

A phylogenetic study of the  
South African representatives  
of the tribe Andropogoneae  
(Poaceae) .

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## LIST OF ABBREVIATIONS

*Bgl1* — *Bacillus grobigii* I

BLFU — Geo Potts Herbarium, Department of Botany & Genetics, University of the Orange Free State, Bloemfontein.

bp — base pair

BT —bootstrap

CI — Consistency index

CO<sub>2</sub> — Carbon Dioxide

CTAB — Hexadecyltrimethyl amonium bromide

d H<sub>2</sub>O — distilled water

DAF — DNA amplification fingerprinting

DNA — Deoxyribonucleic acid

dNTP — Deoxynucleotide triphosphate

EDTA — Ethylene diaminetetra acetic acid

ethanol — Ethylalcohol

HCl — Hydrochloric acid

HCOH — formaldehyde

HI — homoplacy index

*Hind III* — *Haemophilus influenzae* Rd III

*Hinf I* — *Haemophilus influenzae* RF 1

HNO<sub>3</sub> — Nitric acid

i.e. — *it est* (that is)

*ITS* — Internal transcribed spacer

K<sub>2</sub>Cr<sub>2</sub>O<sub>3</sub> — potasiumdichromate

km — kilometers

M — molar

m/v — mass per volume

MgCl<sub>2</sub> — Magnesium chloride

ml — milliliter

mM — millimolar

NaCl — sodium chloride

Na<sub>2</sub>CO<sub>3</sub> — sodium carbonate

Na<sub>2</sub>S<sub>2</sub>O<sub>3</sub> — sodium thiosulfate

PAUP — phylogenetic analysis using parsimony

PCR — polymerase chain reaction

pmol — picomoles

RAPD — Random amplified polymorphic DNA  
RC — rescaled consistency index  
RFLP — restriction fragment length polymorphisms  
RI — retention index  
RNA — Ribonucleic acid  
*rpoC<sub>2</sub>* — RNA polymerase II gene  
SNL — signal to noise  
subsp. — subspecies  
TAE — Tris-acetic acid EDTA  
*Taq* — *Thermus aquaticus*  
Tris — 2-amino-2-(hydroxymethyl)-1,3-propanediol  
UOFS — University of the Orange Free State  
UV — ultraviolet light  
V — Volt  
v/v — volume per volume  
% — percentage  
°C — degrees Celsius  
μg — micrograms  
μl — microlitre  
μM — micromoles



# **Chapter 1**

## **Introduction.**

*That grasses are interesting and important plants are a fact recognised by botanists all over the world, yet it would appear that people in general have hardly appreciated either their interest or their importance. Apart from their almost universal distribution, and quite apart from the fascinating interest attaching to those extraordinary tropical giants, the Bamboos, West Indian Sugar-cane, the huge Reed-grasses of Africa, the Pampas-grasses of South America; and from the utilitarian value of the cereals — Maize, Rice and Wheat — everyone must be struck by the significance of the enormous tracts of land covered by the grasses in all parts of the world (Ward 1901).*

### **1.1 Poaceae**

The grasses form the family Poaceae, a well defined natural group of plants. This family is by no means the largest in terms of species (9 700) and genera (770), coming after Asteraceae and Fabaceae, but its importance is beyond doubt for it provides the grasslands which occupy a third of the lands surface (Schantz 1954), and the cereal crops upon which much of the worlds population depends for its food (Clayton and Renvoize 1986). In southern Africa 194 genera and 967 species, with 847 indigenous species and 329 endemic species (Gibbs Russell *et al.* 1990) represent this family.

Between 1958 and 1966 new systems for the classification of Poaceae were recognised. Most of these systems recognised more than the two initial subfamilies (Panicoideae and Festucoideae). During the last 30 years the number of subfamilies has stabilized at five (Renvoize 1981, Campbell 1985, Watson *et al.* 1985), namely Pooideae Macfarlane and Watson, Bambusoideae Asch and Graeb, Chloridoideae Rouy, Arundinoideae Tateoka and Panicoideae A. Br. (Soderstrom *et al.* 1986). Another subfamily the Centothecoideae Soderstrom is also recognised (Clayton and Renvoize 1986). Recently the GPWG (2001) reclassified the Poaceae into 13 subfamilies namely Anomochlooideae Pilg. ex Potztal, Pharoideae (Stapf) L.G. Clark and Juds, Peulioideae

L.G. Clark. M. Kobay. S. Mathews, Spangler and E.A. Kellogg, Bambusoideae Luerss. Grundz, Ehrhartoideae Link, Pooideae Benth., Fl. Hongk, Aristidoideae Caro, Arundinoideae Burmeist., Danthonioideae Barker and H.P. Linder, Centothecoideae Soderstr., Panicoideae Link, Chloridoideae Kunth ex Beilschm. and *Incertae Sedis*. Of interest to us is the subfamily Panicoideae.

## **1.2 Panicoideae**

This is the largest subfamily including about 3 270 species. Most of the genera fall into two large tribes, the Andropogoneae and Paniceae (Campbell 1985, Kellogg and Campbell 1987).

Earlier systematists (Bentham 1878, Hackel 1890, Hooker 1897) included Zoysieae, Melinideae and Oryzeae as well as the main tribes Paniceae and Andropogoneae and the small tribe Maydeae among the Panicoideae. Stapf and Hitchcock (Bews 1929) later removed the Zoysieae and Oryzeae and placed them among the Pooideae. Stapf increased the number of tribes beyond those recognised by Hackel. From a phylogenetic standpoint this may or may not be advisable, but in the case of two new tribes formed by Stapf, the Arundinelleae and Phoreae, the distinction made is helpful (Bews 1929). Clayton and Renvoize (1986) also included the Steyermarkochloae and the Eriachneae in the Panicoideae, whereas Watson and Dallwitz (1992) placed them in the Arundinoideae. Each of these systems allots the same genera to each tribe, two in each case, suggesting that the tribes themselves internally cohere (Chapman 1996). Most systematists, however, only recognised two main tribes for the Panicoideae, for they see the Maydeae as no more than a subtribe of the Andropogoneae, which are the more advanced, the other and relative primitive tribe being the Paniceae. In 2001 the GPWG divided the subfamily into six tribes, Andropogoneae Dumort, Arundinelleae Sapf, Hubbardieae C.E. Hubb, Isachneae Benth, Paniceae R. Br. and Steyermarkochloae Davidse and R.P. Ellis.

On morphological level it is a fascinating group whose genera lend themselves to arrangement in an orderly sequence of increasing morphological complexity (Clayton and Renvoize 1986). Spikelet characters are diagnostic for the subfamily Panicoideae and distinguish it from the other subfamilies (Ellis 1986). In each spikelet there is one

perfect terminal floret, with a male floret or empty lemma below it (Brown 1810, 1814, Bews 1929, Chapman 1996, Clayton and Renvoize 1986, Watson and Dallwitz 1986, Kellogg and Campbell 1987). In this respect, therefore, the Panicoideae are uniformly advanced. The large tribe of the Andropogoneae shows a distinct advance in having the lower glume always larger than the whole floret, and firmer in texture than the lemmas, usually much hardened and very efficient from a protective standpoint (Bews 1929). In the Panicoideae the reduced one seeded spikelets, falls entire and the axis disarticulate below the glumes. The fruit is thus better protected, especially in the Andropogoneae, where the larger lower glume closely embraces the whole spikelet, and is usually thick and hard. The axes of the racemes or spikelets in the Andropogoneae also break up at maturity.

Not only do the surrounding structures, lemmas and palea serve to protect the fruit, they also are effective in assisting dispersal. They increase the surface exposed to the wind, decrease the specific gravity and in many cases act like the wings in a winged fruit or seed (Bews 1929). Hairs and awns also sometimes assist in wind-dispersal. After dispersal, hygroscopic awns may assist the fruits to become buried in the soil by coiling and untwisting and causing the fruit to rotate. A pointed prickly callus helps in the same direction. The fruits of some xerophytic, advanced types are extraordinary efficient in this way, e.g. *Heteropogon contortus* (L.) Roem. and Schult.

Among the Panicoideae the relative primitive Paniceae are hygrophilous or mesophytic, often forest-margin forms, with only a few xerophytic types, whereas the very highly evolved Andropogoneae are the dominant grasses of enormous areas of tropical and subtropical savannah with, however, a fairly large number of tall coarse, rather hygrophilous types as well and even one or two aquatics (Bews 1929).

Anatomically it is a rather diverse assemblage, with no unique, diagnostic features. Most panicoid grasses have elongated, fingerlike microhairs and horizontally elongated, cross- to dumbbell-shaped silica bodies. Keels are often present and the vasculate is always simple (Ellis 1986). The physiological anatomy of this subfamily is exceptionally variable with all known types being present (Ellis 1977). The supertribe Andropogoneae is homogenous for the C<sub>4</sub> photosynthetic pathway (Sedulsky 1986, Chapman 1996).

Their leaf anatomy are mixed and, therefore, unhelpful in establishing outgroup relationships, though their cuneate lodicules link them to the other tropical subfamilies.

The Panicoideae appear to have had a distinct origin from the Pooideae, but may connect somewhat remotely with ancestral forms, which resemble the Bamboos. It is obvious that the whole of the Panicoideae are rather highly developed in so far as reduction of the number of flowers in the spikelets is concerned. But, this reduction occurs in primitive members of the tribes of the Pooideae as well. Though a very important evolutionary trend, it is after all only one among many. It has taken place within all the separate circles of affinity. Clayton (1981) considered the subfamily Arundinoideae to have provided the ancestral stock for the Panicoideae. Of the other subfamilies it resembles the Chloridoideae in C<sub>4</sub> photosynthesis, chromosome base number and broadly tropical distribution. These two subfamilies differ in numerous ways, however: spikelets, embryos, microhair distal-cell shape and silica bodies. In photosynthetic pathway, 20% of the genera of Panicoideae are non-kranz, of the kranz genera, 89% are of the mestome-sheath (MS) subtype of kranz anatomy (Brown 1977). Chloridoids, on the other hand uniformly have the parenchyma-sheath (PS) subtype. The caryopses of the Panicoideae contain lower levels of glutamine and methionine and higher levels of proline, alanine and leucine than the Chloridoideae (Yeoh and Watson 1981). The levels of proline and glycine in the Panicoideae are intermediate between levels in subfamily Chloridoideae and the tribe Andropogoneae.

Andropogoneae also appears to be related to Arundinelleae and there is sufficient similarity between primitive members of that tribe and of Arundineae to envisage a common ancestry for the group. The origin of Paniceae is more obscure because there are no obvious precursors. Clayton (1981) stated that the subfamily Panicoideae represents a problem and that there is no direct link between the main tribes Paniceae and Andropogoneae. This statement is supported by systematic and cluster analysis (Hilu and Wright 1982, Hilu 1985).

Of interest in this study is the subfamily Andropogoneae.

### **1.2.1 Andropogoneae**

This tribe makes up about half of the grass subfamily Panicoideae with approximately 85 genera and 960 species (Hartley 1958, Clayton and Renvoize 1986, Le

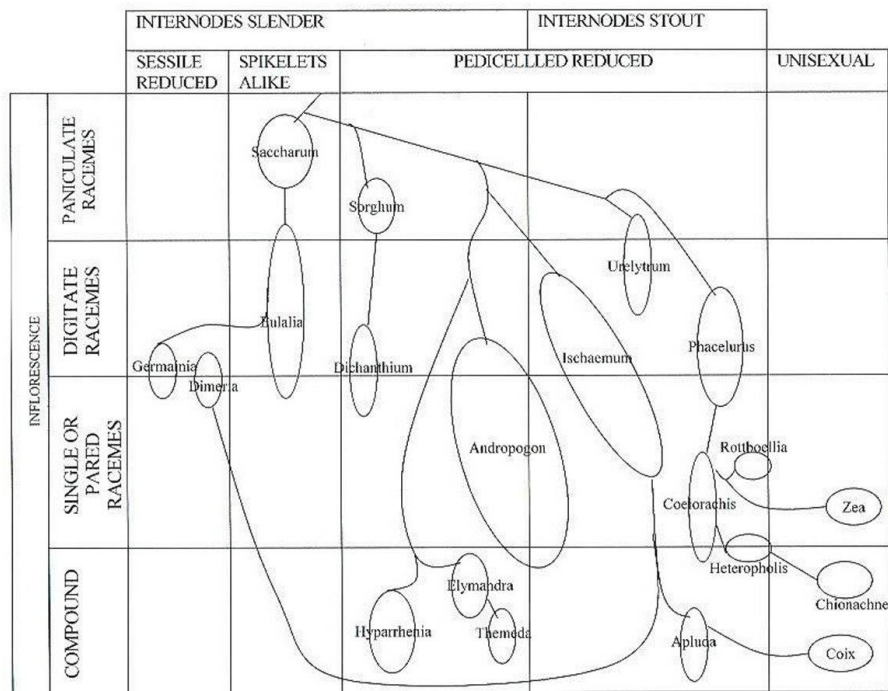


Roux and Kellogg 1999). The genera of the Andropogoneae are typically tropical with only a few species extending beyond the tropics into warm temperate regions.

The anatomy is of the “Kranz MS” type (Brown 1977). These features characters such as a single bundle sheath, chlorenchyma cells arranged radially around the bundles and slender finger-like hairs. This type of anatomy is associated with C<sub>4</sub> photosynthesis (Renvoize 1981).

The most characteristic feature of the tribe is, however, the possession of fragile racemes bearing paired spikelets (Clayton 1972, Connor 1986, Clayton 1986, Clayton and Renvoize 1986, Le Roux and Kellogg 1999). The one spikelet is usually sessile and hermaphrodite, whereas the other is pedicelled and male, sterile or extremely reduced (Hilu and Wright 1982, Clayton 1986, Clayton and Renvoize 1986, Davidse 1986, Le Roux and Kellogg 1999). This makes the entire plant andromonoecious (Le Roux and Kellogg 1999). This unique spikelet structure was hypothesised to be evidence of the monophyly of the subfamily (Kellogg and Campbell 1987) and has been supported by all studies to date (Kellogg and Watson 1993, Kellogg 1998, Masson and Gamer *et al.* 1998, Spangler *et al.* 1999, Mathews *et al.* 2002). Furthermore the great plasticity of this morphological unit gives the Andropogoneae their distinctive flavour (Renvoize 1981). With the only unusual flavonoid reported so far in grasses, arthraxin, a derivative of luteolin, which is reported in *Arthraxon hispidus* (Thunb.) Makino and *Miscanthus tinctorius* (Steud) Hack.A. (Kaneta and Sugiyama 1969). Flavonoid sulphates are usually present in the tropical and subtropical subfamilies for example the Panicoids (18% of species), Chloridoideae (15%) and Arundinoideae *sensu lato* (40%). This chemical difference is linked in part with the distribution of the C<sub>4</sub> photosynthetic pathway (Harbourne and Williams 1986).

Large terminal panicles occur in some of the more primitive genera, but there is a strong tendency, throughout the tribe, to the reduction of the inflorescence, ultimately to a single short raceme. The racemes may be exerted terminally from the spatheole by elongation of the common peduncle or, if paired, they may burst forth laterally by deflection of the stalks supporting the individual racemes. By exploiting the morphological variation in the raceme-segment and panicle, the tribe has evolved some of the most complex structures found in Poaceae and, therefore, a pattern of the advancement along raceme-segment and inflorescence axis can be drawn (Figure 1.1) (Clayton and Renvoize 1986). Another distinctive development in Andropogoneae is the



**Figure 1.1:** Evolution of Andropogoneae. Schematic diagram showing selected genera on two axes of increasing morphological complexity (Clayton and Renvoize 1986).

reduction of the inflorescence and proliferation of axillary branching until, in extreme cases, the branch system come to imitate a panicle (Clayton 1986).

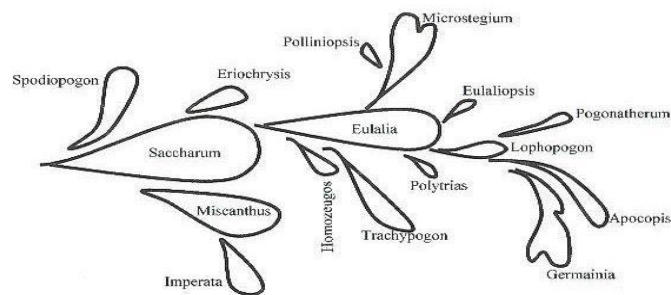
The upper lemmas are membranous and often greatly reduced (Clayton 1986). The tribe divides naturally into two parts: species with an upper lemma and those with the upper lemma entire and awnless.

The original classification of Andropogoneae consists of Dimerieae, Sacchareae, Ischaemeae, Euandropogoneae and Rottboellae (Hackel 1889). Clayton (1972, 1973) recognised ten subtribes with respect to the awns (Table 1). Clayton and Renvoize (1986), on the other hand only recognise five of these subtribes: Saccharinae, Sorginae, Ischaeminae, Andropogoninae and Rottboeliinae (Figure 1.2-1.6) and Watson and Dallwitz (1992) only three: Andropogoninae, Rottboeliinae and Maydeae.

**Table 1:** The ten subtribes recognised by Clayton (1972, 1973) with respect to the presence or absence of awns.

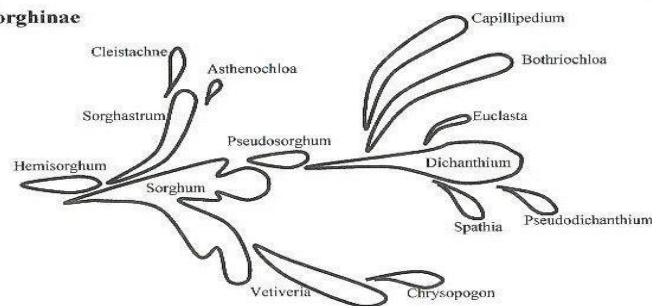
Awned genera	Awnless genera
Andropogoninae	Rottboelliinae
Anthistirrinae	Coicinae
Dimeriinae	Tripsacinae
Saccharinae	
Sorghinae	
Germaniinae	
Ischaemastrae	

**Saccharinae, Germaniinae**



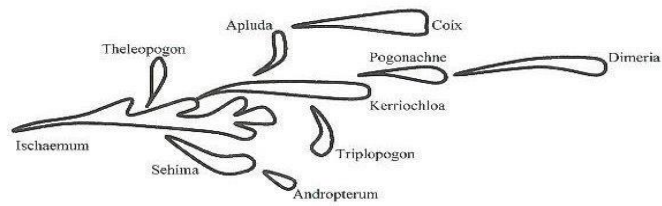
**Figure 1.2:** Subtribe Saccharinae as recognised by Clayton and Renvoize (1986).

**Sorghinae**



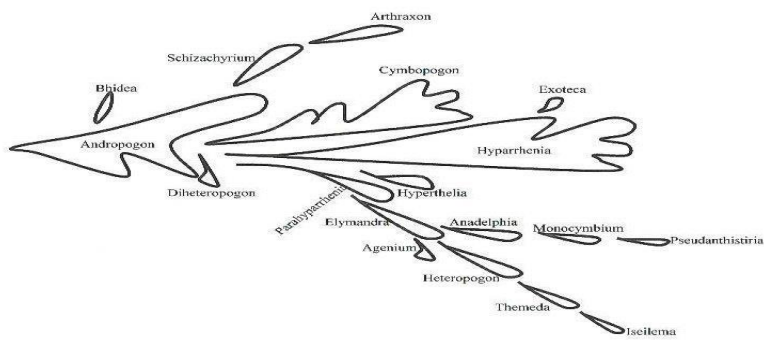
**Figure 1.3** Subtribe Sorghinae as recognised by Clayton and Renvoize (1986).

**Ischaeminae**



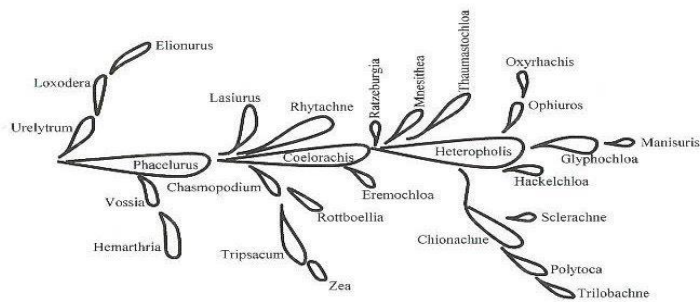
**Figure 1.4:** Subtribe Ischaeminae recognised by Clayton and Renvoize (1986).

**Andropogonae, Anthistirinae**



**Figure 1.5:** Subtribe Andropogonae as recognised by Clayton and Renvoize (1986).

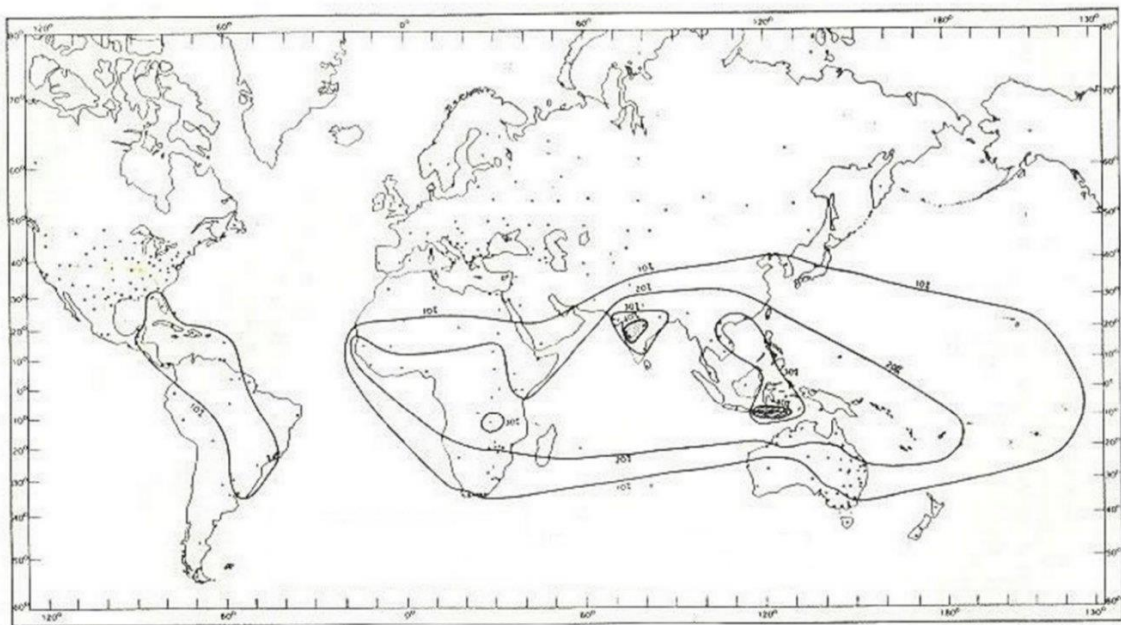
**Rottboelliinae, Tripsacinae, Chionachninae**



**Figure 1.6:** Subtribe Rottboelliinae as recognised by Clayton and Renvoize (1986).



The only important modification to the circumscription of the tribe concerns the genera placed in Maydeae, a tribe whose heterogeneous nature has long been a cause of disputes (Clayton 1986). Celarier (1956) favours the inclusion of the Maydeae in the Andropogoneae and considers the Maydeae a natural extension, through specialization of the Andropogoneae. The inclusion of the Maydeae in the Andropogoneae would also not change the pattern of geographical distribution due to the relatively few species involved (Hartley 1958). In 1958, Hartley composed a geographical distribution map of the Andropogoneae by using 300 floras and floristic lists, each of which is represented on the map by a point located at the approximate geographical centre of the region covered by the flora (Figure 1.7).



**Figure 1.7:** Map of world distribution of the Andropogoneae (Hartley 1958).

The most prominent features of this map are (1) the concentration of the tribe in the tropical and subtropical parts of the world, (2) the majority of species in Indo-Malaysia and (3) the relative lower concentration of the tribe in the western hemisphere in comparison to the eastern hemisphere.

In 1958, Hartley indicated that the distribution of the Andropogoneae is related to the climatic factors, such as winter temperature and rainfall. This is reflected in the predominantly distribution of the tribe in the tropical and subtropical parts of the world.

The main centre of specific differentiation is noted to be in southern Asia. Both taxonomists and cytogenetisists have regarded this taxa rich region to contain the most primitive members of the subtribe. Saccharineae and especially the genus *Miscanthus* Anders. are included here. *Miscanthus* not only has a generalised type of inflorescence from which the more specialised forms are derived, but also shows relationships to other tribes of the grasses (Avdulov 1931, Keng 1939, Celarier 1956). Saccharinae is also a typically tall, hygrophilous grass, which are considered to be ecologically primitive (Bews 1929).

Hartley (1958) confirmed the tribe's adaptation to hot moist conditions, but care must be taken in drawing conclusions about the centres of origin and evolution from these data. In Africa for instance the percentage of species of Andropogoneae does not approach the high levels reached in some parts of Asia, but this does not mean that the tribe has spread into Africa from a centre of origin in Asia. It may in fact well be due to the absence of comparable levels of summer rainfall in Africa.

Evidence also suggests that if spreading has occurred at all, it must have been during the very early history of the tribe. The presence of *Miscanthidium* in Africa, which is closely related to, or perhaps congeneric with *Miscanthus*, indicates that the primitive forms reached Africa at a very early stage, if it did not indeed, originate there. It is also possible that parallel evolution and spread occurred in Africa and Asia, with the tribe retaining its tendency to develop differentiation in regions of high winter temperature and summer moisture (Hartley 1958). The continuing tradition of regional Flora production for Africa and India has provided a reasonable account of the unrevised genera applicable to most of Africa and much of Asia, but otherwise knowledge of their species is uncoordinated and scattered throughout literature (Clayton 1986), making the identification difficult. The problem stems from continuity of variation. Some genera have been studied in detail (De Wet and Harlan 1968, Clayton 1969 a, b, De Wet and Harlan 1970, Clayton 1975, Tothill and Hacker 1976, Soenarko 1977, De Wet 1978), but adequate accounts for many unrevised genera must still be provided (Clayton 1986).

From the information available, it is clear that the African representatives form an integral part of the tribe. For this study, however, we will only concentrate on the South African representatives of the tribe.

### **1.2.1.1 South African representatives**

The South African representatives of this tribe consist of approximately 40 genera and 110 species. These species are represented from the Highveld to the Free State to Lesotho and Kwazulu Natal.

#### **1.2.1.1.1 *Andropogon* L.**

*Andropogon* consists of ~100 species, of which 15 species are indigenous to South Africa, i.e. *A. amethystinus* Nutt, *A. appendiculatus* Nees, *A. brazzae* Franch. W. BE., *A. chinensis* (Nees) Merr., *A. distachyos* L., *A. eucomus* Nees, *A. fastigiatus* SW., *A. festuciformis*, *A. gayanus* Kunth, *A. huillensis* Rendle., *A. lacunosus* J.G. Anders., *A. laxatus* Stapf, *A. mannii* Hook, *A. ravus* J.G. Anders., *A. schirensis* A. Rich. Stapf (1899) described 43 species for the “Flora of Tropical Africa”, of which the most important are: *A. distachyos* L., found in tropical Africa and in the Mediterranean region, from the Canaries to Syria; *A. eucomus* Nees, widespread over the southern half of the continent; *A. amplectens* Nees, from French Guinea and Abyssina through Angola and southward through Zimbabwe, Gauteng and KwaZulu Natal; *A. schirensis* Hochst, all over Africa from Sierra Leone to KwaZulu Natal and *A. gayanus* Kunth, a polymorphic species up to ten feet tall, very common in the Cape Verde Island across the continent to Sudan and Zimbabwe. The species are mostly perennial and together they form an important part of the open savannas of Africa. *Andropogon gayanus*, an important pasture species, as well as some others occur in the high grass savannas of the moist tropical belt (Bews 1929, Clayton and Renvoize 1986).

The genus is divided into four sections according to the shape of the sessile, i.e. *Andropogon*, *Leptopogon*, *Piestium* and *Notosolen*.

The most studied specimen of the genus *Andropogon* seems to be *A. gayanus*, probably due to its pasture importance. From one of these studies, Humphreys (1981) concluded that it is a tussock forming grass with erect (apogeotropic) culms having long

internodes. Clayton and Renvoize (1986) also concluded that *A. gayanus* have an external ligule and false petioles and leaf nectaries. Fisher *et al.* (1994) also discovered the globally important fixation of carbon in the New World savannah *A. gayanus*.

The importance of this genus is mostly for prevention of erosion and pasture purposes (van Oudtshoorn 1999).

#### **1.2.1.1.2 Arthraxon P. Beauv.**

This genus consists of seven species with one indigenous species, i.e. *A. lanceolatus* (Roxb.) Hochst., all delicate grasses, which are perennial or annual. Their habitats include rocky slopes, shady glades and old farmlands.

It is a homogeneous genus with several unusual features, which makes it difficult to place. The only possible comparison being with *Schizachyrium*, which awn is also sometimes basal though from a bifid lemma (Clayton and Renvoize 1986).

#### **1.2.1.1.3 Bothriochloa Kuntze**

This aromatic grass is found throughout the tropics in open grassy places (Clayton and Renvoize 1986). Of the 35 species only three are indigenous to South Africa, i.e. *B. bladhii* (Retz.) S.T. Blake, *B. insculpta* (A. Rich.) A. Camus, *B. radicans* (Lehm.) A. Camus. These are perennial grasses (Watson and Dallwitz 1992).

Due to the complex pattern of hybridisation created by rapacious introgression from *B. bladhii* (De Wet and Harlan 1970), the boundaries between *Bothriochloa*, *Capillipedium* and *Dichanthium* are somewhat blurred. Williams and Gillard (1971) also found that the occurrence of *B. bladhii* decreased with increasing tree density in nature. Because of a capacity for genetic mixing in *B. bladhii*, De Wet and Harlan (1966) called it a compilospecies. However, apart from this one miscellaneous species, the genus is morphologically and genetically distinct.

De Wet and Scott (1965) studied the essential oils and taxonomic implications of *Bothriochloa* and Heslop-Harrison (1961) came to the conclusion that the glume pits play a role in the cleistogamous flowering by obstructing the emergence of the anthers.



Most of the times the occurrence of these grasses in a veldt is an indication of overgrazing and plays an important part in the prevention of soil erosion (van Oudshoorn 1999).

#### **1.2.1.1.4 *Chrysopogon* Trin.**

This genus is commonly found in the tropical and warm temperate regions of the Old World, mainly in Asia and Australia, with one indigenous species in South Africa, i.e. *C. serrulatus* Trin. (Clayton and Renvoize 1986). Their habitat range from open disturbed places, subdesert to rain forest. Some species are used as thatching grasses in some parts of Africa and *C. aciculatus* is often used for lawns in the humid tropics.

The genus *Chrysopogon* are tufted perennial grasses (Clayton and Renvoize 1986, Watson and Dallwitz 1992).

*Chrysopogon* does seem to intergrade with *Vetiveria* via *Chrysopogon sylvaticus*, and the separation of these genera is arbitrary, particularly in Australia (Clayton and Renvoize 1986). It is, however, somewhat justified by the treatment of species with triads as a single entity.

#### **1.2.1.1.5 *Cleistachne* Benth.**

This is a coarse annual grass consisting of one species, i.e. *C. sorghoides* Benth.. Their habitats are mostly old farmland in tropical Africa and India and due to the extreme reduction of the racemes this genus can be baffling, but its spikelets do ally it to *Sorghum* (Clayton and Renvoize 1986).

#### **1.2.1.1.6 *Coelorachis* Brongn.**

This tall perennial grasses are distributed throughout the tropics and consists of ~20 species, with one indigenous to South Africa, i.e. *C. capensis* (Trin.) Roberty. Its habitat includes grassland and open woodland, often favouring damp soils (Clayton and Renvoize 1986, Bews 1929).

*Coelorachis* is closely related to *Rhytachne*, with no character being wholly reliable and a few species are difficult to assign (Bews 1929, Rosengurt 1984, Clayton and Renvoize 1986).

#### **1.2.1.1.7 Coix L.**

The genus *Coix* consists of five species that are annual or perennial. They are helophytic to mesophytic, shade species or species of open habitats. Habitats further include forest margins and swamps (Clayton and Renvoize 1986, Gibbs Russell *et al.* 1990, Watson and Dallwitz 1992).

*Coix lacryma-jobi* L. (Job's tears) has been introduced throughout the tropics (Clayton and Renvoize 1986). This species is widely cultivated for the sake of its false fruits, which are used as beads and in the making of rosaries. It is also employed as a fodder crop and the utricles are sometimes ground up for flour (Bews 1929, Clayton and Renvoize 1986).

#### **1.2.1.1.8 Cymbopogon Spreng.**

This genus consists of ~40 species (6 indigenous to South Africa, i.e. *C. dieterlenii* Stapf ex Phill., *C. excavatus* (Hochst.) Stapf ex Burt Davy, *C. marginatus* (Steud.) Stapf ex Burt Davy, *C. plurinoidis* (Stapf) Stapf ex. Burt Davy, *C. prolixus* (Stapf) Phill., *C. validus* Stapf Burt Davy) distributed in the tropics and subtropics and introduced to tropical America. This is a tall, mostly perennial grass with aromatic leaves.

It is a homogeneous genus of narrowly circumscribed species, which are sometimes difficult to distinguish. The variation in this genus is indicated by five series according to the racemes, spikelets and glumes, namely: Series *Proceri*, Series *Refracti*, Series *Citrati*, Series *Cymbopogon* and Series *Rusae* (Clayton and Renvoize 1986). Series *Proceri* is considered to be primitive due to their straight bases.

*Cymbopogon* shows an affinity to *Andropogon* according to the concave two-keeled sessile spikelet, but there are distinguishing features between the two genera. The aromatic flavour of the leaves is such an example. This aromatic oil in *Cymbopogon* is

extensively used in perfumery (Lavania 1988, Chapman 1996) and *C. citratus* are used as a culinary herb (Watson and Dallwitz 1992, van Oudtshoorn 1999).

#### **1.2.1.1.9 *Dichanthium* Willem.**

This genus consists of ~20 Old World tropical species, with one indigenous species and one naturalized species in South Africa, i.e. *D. annulatum* (Forsk.) Stapf, *D. aristatum* (Poir.) C.E. Hubbard (Clayton and Renvoize 1986). Their habitats are open places, from subdesert to marshland, particularly when subjected to disturbance. It is mostly annual grasses (Chapman 1996) with occasional aromatic leaves and is used for grazing (van Oudtshoorn 1999).

Although *Dichanthium* is recognised by its homogamous pairs and obtuse sessile spikelets, some species lack these features and resembles *Bothriochloa*, from which they are separated by the solid pedicels (Clayton and Renvoize 1986).

#### **1.2.1.1.10 *Diheteropogon* Stapf**

This is an annual or perennial grass distributed in tropical and South African savanna (Clayton and Renvoize 1986), i.e. *D. amplexens* (Nees) Clayton, *D. filifolius* (Nees) WD Clayton, and used for grazing and control of soil erosion (van Oudtshoorn 1999).

*Diheteropogon* is a small homogeneous genus, which links *Andropogon* section *Piestium* to *Pararhynchia*, with a barely justified separation from *Andropogon*.

#### **1.2.1.1.11 *Elionurus* Kunth ex Willd**

The genus *Elionurus* consists of 15 species, throughout Africa, America and Australia, with two species indigenous to South Africa, i.e. *E. muticus* (Spreng.) Kunth, *E. tripsacoides* Humb. & Bonpl. It is a climax or subclimax grass in savanna or often on dry soils. They are generally tufted perennials and not well grazed because of its pungent odour and aromatic taste, which is not commercially exploited. *Elionurus* is said to be a homogenous genus, with an alliance to *Loxodera*, because of the resemblance in the

short callus and lobbed internode tip (Bews 1929, Roberts 1973, Clayton and Renvoize 1986, Roberts and Fourie 1989).

#### **1.2.1.1.12 *Elymandra* Stapf**

This genus consists of 6 species, with one indigenous, i.e. *E. grallata* (Stapf) Clayton. These species are annual or perennial grasses with tall culms found in the shade or open habitats. Although there is some variance between the species, all have the olive-green homogamous spikelets and dark sessile spikelet. There seems to be a correlation between the number of homogamous pairs and fertile spikelets, which decrease as the number of pairs increase (Clayton and Renvoize 1986). Connor (1986) also suggested that there are 4-5 times as many male spikelets as perfect spikelets in *Elymandra*. The pedicelled spikelet callus of *Elymandra* also suggests a relationship to *Parahyparrhenia* (Clayton and Renvoize 1986).

#### **1.2.1.1.13 *Eriochrysis* P. Beauv.**

This is a tufted perennial grass with seven species of which two are indigenous to our region, i.e. *E. brachypogon* (Stapf) Stapf., *E. pallida* Munro.. Their habitats include swamps and moist places (Watson and Dallwitz 1992). It is a homogeneous genus, which is allied to *Saccharum*, but is distinguished by the compact rufous racemes and slightly dimorphic spikelets (Clayton and Renvoize 1986).

#### **1.2.1.1.14 *Eulalia* Kunth**

This genus consists of ~30 species from the Old World tropics, with two indigenous to our region, i.e. *E. aurea* Gravier, *E. villosa* (Thunb.) Nees. They are tufted perennial grasses found in grasslands and sometimes in moist places (Watson and Dallwitz 1992).

*Eulalia villosa* is a significant weed species and *E. aurea* an important native pasture species.



*Eulalia* is related to *Saccharum*, but differ in the digitate racemes and their tendency for wind-dispersed spikelets to rely upon hairs from glumes and internodes rather than from the callus. It does, however, seem that the genus stands at the junction of several divergent lines, with links to *Polytrias*, *Lophopogon* and *Homozeugos* (Clayton and Renvoize 1986).

#### **1.2.1.1.15 Hackelochloa Kuntze**

This coarse annual species (*H. granularis* L. Kuntze) is distributed throughout the tropics in weedy places. Apart from a unique spikelet shape the genus differs little from *Heteropholis* (Bews 1929, Clayton and Renvoize 1986, Gibbs Russell *et al.* 1990).

#### **1.2.1.1.16 Hemarthria R.Br.**

The genus *Hemarthria* consists of 12 species, distributed through the old world tropics, from India, Australia, America through to Africa (*H. altissima* (Poir.) Stapf & C. E. Hubb.) and the Mediterranean region and are used for grazing in some parts of Africa (Bews 1929, Clayton and Renvoize 1986, van Oudtshoorn 1999). This grass is hygrophilous, usually occurring in or near water. Some species have culms of 4.57-6.10m. long, often floating. *Hemarthria* is a rambling perennial grass, with evidence of a fused pedicel, linking it to *Heteropholis*, but evidence suggests a more close relationship to *Phacelurus* (Bews 1929, Clayton and Renvoize 1986).

#### **1.2.1.1.17 Heteropogon Pers.**

This genus is distributed in the tropics and subtropics of both hemispheres. Its habitat includes dry open places on poor soils, and in North America it is found in rocky places from Texas to Arizona (Bews 1929, Clayton and Renvoize 1986). This wide range of geographical distribution is associated with considerable morphological variation and phenological behavior (Tothill 1966, 1968, Tothill and Hacker 1976). *Heteropogon contortus* (L.) Roem. & Schult. is a significant weed species and native pasture species

(Watson and Dallwitz 1992). These grasses can be either annual or perennial (Clayton and Renvoize 1986), i.e. *H. contortus*, *H. melanocarpus*.

Some species of *Heteropogon* also adapted a needle-sharp, sessile spikelet callus, which penetrates clothing, as an efficient dispersal mechanism (Clayton and Renvoize 1986). The homogamous spikelets in *Heteropogon* as well as the developed pedicelled spikelet callus, suggest a loose relationship with *Elymantra*.

#### **1.2.1.1.18 Hyparrhenia Fourn.**

This species are annual or perennial, usually caespitose with tall culms. The genus consists of 55 species and is distributed mostly in Africa (20 indigenous, i.e. *H. anamesa* Clayton, *H. collina* (Pilg.) Stapf, *H. cymbaria* (L) Stapf, *H. dichroa* (Steud.) Stapf, *H. dregeana* (Nees) Stent., *H. filipendula* (Hochst.) Stapf, *H. fili-var pilosa* (Hochst.) Stapf, *H. finitima* (Hochst.) Stapf, *H. gazensis* (Rendle) Stapf, *H. hirta* (L.) Stapf, *H. newtoni var macra*, *H. newtonii* (Hack.) Stapf, *H. nyassae* (Rendle) Stapf. syn, *H. pilgeriana* C.E. Hubb., *H. peocilotricha* (Hack.) Stapf, *H. quarrei* Robyns, *H. rudis* Stapf, *H. rufa var rufa* (Nees) Stapf, *H. schimperii* (A. Rich.) Stapf, *H. tamba* (Steud.) Stapf, *H. umbrosa*, *H. variabilis* Stapf), extending to tropical America, Asia, Australia and to the Mediterranean countries (Bews 1929, Humphreys 1981, Clayton and Renvoize 1986). It is one of the most important genera in high grass savannah, as well as in drier types of grassland (Bews 1929). It is widely used for thatching in Africa and well grazed by animals when immature. It is also a successful coloniser of bare soil, particularly in gravely or stony situations and may be used to revegetate eroded shale slopes and certain poor soils (Roberts 1973, Clayton and Renvoize 1986).

*Hyparrhenia* is linked to *Cymbopogon* through superficial resemblance with *H. glabriuscula*, which is rated as one of the most primitive species (Clayton and Renvoize 1986).

#### **1.2.1.1.19 Hyperthelia Clayton**

This genus consists of six species with one indigenous to South Africa, i.e. *H. dissoluta* (Nees ex Steud.) Clayton syn. It is distributed throughout Africa and introduced

to tropical America, the remaining species localized in southern Sudan and Central African Republic. They are tall annual or perennial grasses, which is used for thatching and sometimes harvested for grain (Clayton and Renvoize 1986, van Oudtshoorn 1999).

The genus is related to *Hyparrhenia*, but it has more features in common with *Parahyparrhenia*.

#### **1.2.1.1.20 Imperata Cirillo**

This rhizomatous perennial grass consists of eight species with one indigenous to our region, i.e. *I. cylindrica* (L.) Raeuschel. The genus is distributed throughout the tropics extending to warm temperate areas in both hemispheres (Bews 1929, Clayton and Renvoize 1986). Varieties occur all over Africa and Asia, and are among the dominant grasses in high grass savannas. Some species are distinctly hygrophylous, but others are psammophilous, or occur as weeds in cultivated land (especially in lands where rice, cotton, coffee and tea are cultivated). Economically these species are also important for they are used for fuel, paper, thatching and ornamental purposes (van Oudtshoorn 1999).

The homogeneous genus *Imperata* is allied to *Miscanthus*, but the contracted panicles and deficient floral parts distinguish the two genera (Clayton and Renvoize 1986). Phenotypically these two genera are also clearly distinguishable.

#### **1.2.1.1.21 Ischaemum L.**

This genus consists of ~60 species, mostly in southern Asia and Australia with two in South Africa, i.e. *I. schaeum afrum* (J.F. Gmel.) Dandy, *I. fasciculatum* Brongn.. Their habitats are mostly damp or shady places and they are scattered through grasslands but not as a rule dominant (Bews 1929, Clayton and Renvoize 1986). They are tufted perennial grasses, with *I. muticum* and *I. rugosum* used as native pasture species (Watson and Dallwitz 1992). There is no adequate treatment of the genus available, but the variation of the sessile spikelet and pedicelled spikelets lead to the recognition of five sections (Clayton and Renvoize 1986), namely: *Fasciculata*; *Ischaemum*; *Coelischaeum*; *Aristata* and *Aurea*.

The sections evolved in this order, with Sect. *Fasciculata* closely related to *Eulalia*. The genus is sometimes difficult to distinguish from *Andropogon*, but the shape of the rachis is usually characteristic (Clayton and Renvoize 1986).

#### **1.2.1.1.22 *Microstegium* Nees**

These are creeping or climbing grasses, which can be either annual or perennial with 15 species commonly adventive (Watson and Dallwitz 1992).

The genus is distributed through tropical Asia and Africa and consists of 15 species with one indigenous to South Africa, i.e. *M. nudum* (Trin.) A. Camus. *Microstegium* resembles *Eulalia* due to the presence of the cordate lemmas (Clayton and Renvoize 1986).

#### **1.2.1.1.23 *Miscanthus* Anderss.**

This is a tufted or rhizomatous perennial tall grass, distributed mainly in Asia, but extending to Africa. There are ~20 species in this genus of which 2 are indigenous to our region, i.e. *M. capensis* (Nees) Anderss., *M. junoeus*. Their habitats are open places, such as hillsides and marshes and several of the larger species are grown as ornamentals (Bews 1929, Clayton and Renvoize 1986, Chapman 1996). Some species are also used in the purifying of water ecosystems (van Oudtshoorn 1999).

An interesting oddity is the suppression of the leaf lamina, resulting in a quill-like blade, formed from the midrib. *Miscanthus* is closely allied to *Saccharum*, with which it hybridises (Clayton and Renvoize 1986).

#### **1.2.1.1.24 *Monocymbium* Stapf**

This perennial grass is distributed in Africa and in South Africa, through the Gauteng and KwaZulu Natal (Bews 1929, Clayton and Renvoize 1986, *M. ceressiforme* (Nees) Stapf). They are usually indicators of sour soil and is often subdominant in *Themeda* grass-veldt (Bews 1929, van Oudtshoorn 1999).

The conspicuous spatheoles of *Monocymbium*, gives it an entire different look from *Anadelphia* even though they are rather similar genera. The fact that there are no intermediate species does, however, makes this separation justifiable (Clayton and Renvoize 1986).

#### **1.2.1.1.25 *Oxyrhachis* Pilg.**

*Oxyrhachis* consists of only one species, distributed in Africa and Madagascar, i.e. *O. gravillima*. It is a perennial with linear, folded, or rolled leaf blades found at upland streamsides and marshy places. *Oxyrhachis* is an isolated genus, which shows resemblance to *Ophiuros* (Clayton and Renvoize 1986, Gibbs Russell *et al.* 1990).

#### **1.2.1.1.26 *Phacelurus* Griseb.**

This genus consists of ~9 species with one indigenous to South Africa, i.e. *P. franksae* Griseb. The distribution of this genus includes Indo-China and Japan.

The genus consists of perennial grasses, with linear, flat or folded leaf blades and 200-600mm high culms and grow in shady places or open habitats (Clayton and Renvoize 1986, Gibbs Russell *et al.* 1990).

This is a variable genus, whose characters are poorly correlated. Attempts at subdivision have had to rely on the subjective weighting of single characters, therefore it is best to treat the species as a single diffuse cluster. Of the other genera *Phacelurus* is the closest to *Ischaemum*.

#### **1.2.1.1.27 *Rhytachne* Desv.**

This genus consists of ~ 12 species, with three indigenous to South Africa, i.e. *R. latifolia*, *R. robusta*, *R. rottboellioides*. The genus consists of plants that are annual or perennial growing in pans and riversides or grassland (Gibbs Russell *et al.* 1990). The distribution of this genus includes Madagascar and tropical South America, with a habitat of moist or seasonally flooded grassland. This genus was derived from *Phacelurus* via *Phacelurus gabonensis*, according to Clayton and Renvoize (1986).

#### **1.2.1.1.28 *Rottboelia* L.f.**

*Rottboelia* is a small genus with four closely allied species (Clayton and Renvoize 1986) of which one is indigenous to South Africa (*R. cochinchinensis* (Lour.) W.D. Clayton). Clayton and Renvoize (1986) describe this species as a serious tropical weed. The habitat of this species is variable and includes swamps, disturbed places or dry soils in woodlands.

*Rottboelia* species are coarse annuals, growing up to 12 feet high. Their leaf blades are broad and flat. According to Chapman (1996), *Rottboelia* is one example among several known to switch from cleistogamy to chasmogamy.

#### **1.2.1.1.29 *Saccharum* L.**

*Saccharum* consists of ~33 species, adapted to temperate to subtropical areas. Important genus in moist to wet habitats. *Saccharum* is commonly split into several genera, but the characters relied on are more appropriate to infrageneric categories (Clayton and Renvoize 1986). The most commonly division into awned (*Erianthus*) and awnless species, seems artificial and Narenga, with its coriaceous glumes, seems no more than the expression of a trend found elsewhere in the genus. Stapf retains *Erianthus ravennae* Beauv. in this genus as *S. ravennae* L.

In South Africa and US the species of this genus behaves as perennials under cultivation (Bews 1929). It grows up to 6 or 8 feet tall, with solid yellowish-green canes and thick rich green leaves.

This genus is a native of India, but is now widely distributed in cultivation. There are many varieties, some of which are used as forage grasses for all classes of stock (Bews 1929). Probably the most important feature of this genus is the use of *S. officinarum* sugar cane, which is harvested as thick culms and pressed through heavy rollers to extract the juice. Three products are produced: sugar; molasses and bagasse. Since all three are the result of photosynthesis, they represent energy capture and could therefore be used for fuel. The sugar and molasses being fermented to ethanol and the bagasse burned (Chapman 1996).

### **1.2.1.1.30 *Schizachyrium* Nees**

This genus consists of ~60 species with six indigenous to South Africa, i.e. *S. brevifolium* (Sw.) Nees, *S. exile* (Hochst.) Pilger., *S. jeffreysii* (Hack.) Stapf, *S. rupestre* (K.Schum.) Stapf, *S. sangiuneum*, *S. ursulus* Stapf. It occurs in the tropics of both hemispheres and forms an important part of the wild prairie hay. Their habitat includes sandy beaches and savannah (Bews 1929, Clayton and Renvoize 1986). It is a tall, sometimes delicate, annual or perennial grass with sexual and asexual reproduction (Connor 1986).

This genus is closely allied to *Andropogon* Sec. *Leptopogon* but is distinguished by its single racemes.

### **1.2.1.1.31 *Sehima* Forssk.**

This genus consists of 6 species ranging from South Africa (2 indigenous, i.e. *S. galpinii* Stent., *S. ischaemoides*) to India and Australia. It is a tufted perennial or annual grass in dry bushland and grows on lava rocks and on seasonally waterlogged black clay's (Bews 1929, Clayton and Renvoize 1986, Watson and Dallwitz 1992).

This genus is remarkably uniform even though they are variable in some key characters. The shape and nervation of the pedicelled spikelet being its most particular characteristic. Some do regard this genus as a segregate of *Ischaemum* (Clayton and Renvoize 1986). An important feature of this genus is that it only grows in clayey soil.

### **1.2.1.1.32 *Sorghastrum* Nash**

This close relative of *Sorghum* consist of ~20 species and range from Africa to tropical America. Two of these species is indigenous to South Africa, i.e. *S. friesii* (Pilg.) Pilger, *S. stipoides* (Kunth.) Nash and is found in savanna and woodland margins. It is either an annual or perennial grass (Bews 1929, Clayton and Renvoize 1986).

*Sorghastrum avenaceum* survives repeated burnings. Their flowering stems can grow up to seven feet tall and the leaves can reach two feet in length and are tapered.



*Sorghastrum nutans* have a tall stalked seedhead with rust-gold, soft seeds. The sheath is usually hairy but can have sparse or no hairs. This species is similar to *Stipa sporea* and *Andropogon gerardi*. It is a fair grazing grass for wildlife and makes good grazing for livestock.

The generic circumscription of this genus is stretched by some species, whose pedicels bear fertile awned spikelets, which also indicates a strong alliance with *Saccharum* (Clayton and Renvoize 1986).

### **1.2.1.1.33 Sorghum Moench**

The genus *Sorghum* Moench consists of ~2 species and is widely distributed throughout the tropics and subtropics of the Old World. In South Africa two species are indigenous. This genus consists of annual or perennial, tufted or sometimes rhizomatous, mostly robust grasses (Clayton and Renvoize 1986, Gibbs Russell *et al.* 1990). Their leaf blades are usually flat with the inflorescence a panicle, either open or contracted. The spikelets are overtly heteromorphic with the pedicellate much narrower and awnless (Gibbs Russell *et al.* 1990).

*Lasiorrhachis*, with pedicelled spikelets that are fertile and rudimentary within the same panicle, lies between *Saccharum* and *Hemisorghum*, but it are not distinct enough to stand on its own. Other species well known in this genus includes: 1) *S. guineense* "Guinea corn", with three varieties and numerous races; 2) *S. dura*, the most common cultivated sorghum in Egypt; 3) *S. caffrorum*, the "Kafir corn" or "Amabela" of South Africa.

This genus is vary variable, and is usually subdivided into sections, *Chaetosorghum*, *Heterosorghum*, *Parasorghum*, *Sorghum* and *Stiposorghum* (De Wet *et al.* 1970, De Wet *et al.* 1972, Harlan and de Wet 1972, De Wet 1978). Section *Parasorghum* and *Stiposorghum* appears to be some what distant from the rest of the genus. Duvall and Doebley (1990) and Sun *et al.* (1994 b) found *Parasorghum* to be paraphyletic in molecular phylogenetic studies.

Section *Sorghum*, which includes cultivated sorghum, a complex of closely related annual taxa from Africa (*S. bicolor*), and a complex of perennial taxa from southern Europe and Asia (*S. halepense*). This section is also related to *Saccharum*, with

which it will hybridize. Numerous classifications of this section are available (Snowden 1936, Murty *et al.* 1967, Jakusyevsky 1969, Harlan and De Wet 1971, 1972).

*Sorghum bicolor* is one of the most economically important crops in the world (Dogget 1976, 1988, FAO Yearbook Production Report 1993, Arriola and Ellstrand 1996) and has received much attention, but the rest of the genus has not received that much attention (Garber 1950, Harlan and De Wet 1972, Lazarides *et al.* 1991).

*Sorghum bicolor* is an annual grass that is wind pollinated and outcross at a rate of 10-15% (Ellstrand and Foster 1983, Eastin and Lee 1985, Dogget 1988, Arriola and Ellstrand 1996). The morphological wild varieties of *S. bicolor* differ from each other primarily in respect to inflorescence structure and distribution (de wet and Harlan 1971).

It has been suggested (Vinall and Getty 1921, Hadley 1953, Baker 1972, Dogget 1988, Ariolla and Ellstran 1996) that *S. bicolor* hybridizes with *S. halepense* (Johnsongrass) under field conditions. *Sorghum halepense* is a perennial grass that is wind pollinated and reproduces vegetatively as well as through sexual means (Arriola and Ellstrand 1996).

*Sorghum* is the staple food of millions of people, mainly in Africa and India. The genus is an excellent silage and is also directly used for grazing. It does, however, contain hydrocyanic acid and can be poisonous if it is welted. The *Sorghum* culm is also used to make syrup and fuel.

#### **1.2.1.1.34 *Thelepogon* Roth.**

This is a coarse annual grass, which is mainly found in tropical Africa, extending to Indonesia. Its habitat include seasonally soils and disturbed ground.

Of the other genera it resembles *Ischaemum*, distinguished only by the barren pedicel (Clayton and Renvoize 1986).

#### **1.2.1.1.35 *Themeda* Forssk.**

*Themeda* is in many ways one of the most advanced genera in the tribe with 18 species and one indigenous to South Africa, i.e. *T. triandra* Forssk. They cover immense areas of the subtropical grassland and are very important grazing species in South Africa.

It is a dominant grass, which will increase when fires occur frequently, providing overgrazing does not take place (Bews 1929, Clayton and Renvoize 1986 and van Oudtshoorn 1999). It is the “Rooigras” or “Red grass” of South Africa or “Insinde” (Zulu) and is a valuable forage grass. This is a relatively mesophytic surface rooting perennial (Bews 1929 and Clayton and Renvoize 1986).

#### **1.2.1.1.36 *Trachypogon* Nees**

This genus consists of ~13 species and is distributed through Africa (1 indigenous to South Africa, i.e. *T. spicatus* (L.f) Kuntze), *Madagascar* and the warmer regions of America. It forms a dense covering in the high rainfall areas and plays an important role in the protection of soil erosion. In some parts of America, it is an important constituent of the grazing areas (Bews 1929, Clayton and Renvoize 1986, Gibbs Russell *et al.* 1990, van Oudtshoorn 1999). It is mostly tufted perennial (Bews 1929, Clayton and Renvoize 1986).

The genus *Trachypogon* has a similar raceme structure to *Germania*, but it was achieved by a different route, for it resembles *Homozeugos* rather than *Lophopogon* (Bews 1929, Humphreys 1981, Clayton and Renvoize 1986, , Gibbs Russell *et al.* 1990).

In 1929 Stapf described the nomenclature of the species of *Trachypogon* in the "Flora of Tropical Africa", to be controversial. He extends it to cover *T. montufari* of Farnier (and of Hitchcock), but the other species of *Trachypogon* are more local (Bews 1929).

#### **1.2.1.1.37 *Urelytrum* Hack.**

This genus consists of seven species, with one indigenous to South Africa, i.e. *U. agropyroides* (Hack.) Hack. The genus consists of annual or perennial grasses with 600-2500 mm high culms. The leaves are auriculate with linear, flat or rolled (convolute) leaf blades.

According to Clayton and Renvoize (1986) *Urelytrum* have the thickened internodes and awnless upper lemma, that is characteristic of their subtribe *Rottboelinneae*, but some features, in particular the callus, are reminiscent of the subtribe

*Ischaemum* and *Andropogonineae*. There is also an alliance between some *Urelytrum* and *Schizachyrium* species.

#### **1.2.1.1.38 *Vetiveria* Bory**

The genus *Vetiveria* consists of ~10 species, with one indigenous to South Africa, i.e. *V. nigritana* (Benth.) Stapf. *Vetiveria* grows about 2m tall and is a densely tufted perennial grass, which is almost seedless. Its habitat is flood plains and stream banks. It is a deep rooted grass that is able to withstand both draught and flooding, with tough foliage that makes it unpalatable to livestock. This grass is mostly cultivated for its aromatic roots. It is the "Khas - Khas" or "Khus Khus", the source of "Vetiver" oil (Bews 1929, Clayton and Renvoize 1986, Gibbs Russell *et al.* 1986, Chapman 1996).

According to Chapman (1996), Vetiver grasses represents the student of grasses. They are related to *Sorghum* subgen. *Parasorghum* and *Vetiveria pauciflora*, with its 2-3 spikelet pairs per raceme also links the genus to *Chrysopogon*.

#### **1.2.1.1.39 *Vossia* Wall. and Griff.**

This genus consists of one species, *V. cuspidata* Griff. and is distributed throughout tropical Africa and India. It is an aquatic perennial, which can be found close to water, or often floating. The culms are 1000-2000mm high, with floating culms up to 7m long. The leaf blades are broad and flat (Bews 1929, Clayton and Renvoize 1986, Gibbs Russell *et al.* 1990).

Gibbs Russell *et al.* (1990) describe this genus as hydrophytic or helophytic and as a segregate from *Phacelurus*.

#### **1.2.1.1.40 *Zea* L.**

Maize's agricultural importance has lead to the use of *Zea* as a model for genetics, molecular biology and systematics (Doebley 1990, Kellogg and Blichler 1993 and Buckler and Holtsford 1996a). The genus consists of ~4 species. They originated in America, but were cultivated from prehistoric times by the races of American aborigines

from Peru to the middle of North America. Introduced into the Old World, it spread rapidly and is grown extensively everywhere in the warmer regions of the world. They are now the chief crop in Africa, being cultivated by natives and Europeans.

In 1980 (Doebly and Iltis) used multivariate techniques to show that maize and the teosintes fell into two groups, one which consisted of maize plus two annual teosintes. These they treated in an emended *Z. mays* as subsp. *mays*, subsp. *mexicana* (Schrader) Iltis, and subsp. *parviglumis* Iltis and Doebley, respectively, *Z. mays* was then the sole species in *Zea* sect. *Zea*. The other phenetic group is comprised of three species, *Z. luxurians* (Durieu and Ascherson) Bird, *Z. diploperenis* Iltis, Doebley and Guzman, and *Z. perenis* (Hitchcock) Reeves and Mangelsdorf, all included in *Zea* sect. *Luxuriantes*.

The genus consists of annual or rarely perennial species with broad leaves. *Z. mays* is morphologically so different from the other species in the genus, that it is sometimes placed in a separate genus. Although the differences are extreme to the eye, they were found to be under relatively simple genetic control and are such as would be expected between species subjected to disruptive selection.

*Zea* L. “Maize”, “Indian Corn”, “Mealie” is widely cultivated, being one of the most important of all the cereal grasses, giving comparatively high yield under suitable conditions. It is an important staple cereal of tropical regions, and also extensively grown as a forage crop and is an important source of oil, syrup and alcohol (Clayton and Renvoize 1986).

The phenetic systematics of *Zea* are confirmed by studies of variation in isozymes (Doebley 1984, Doebley *et al.* 1984, 1987a), in chloroplast DNA restriction sites (Timothy *et al.* 1979, Doebley *et al.* 1987a, 1987b, Doebley 1990), in DNA-DNA hybridization (Hake and Walblot 1980) and in ribosomal and 5S RNA genes (Zimmer *et al.* 1988, Buckler and Holtsford 1996a, 1996b).

From phylogenetic studies (Buckler and Holtsford 1996a, 1996b) it is clear that *Zea* L. is a sister species of *Tripsacum*.

## **1.3 Cytogenetics**

During the past century the part of biological science which deals with evolution and classification has entered a new era, dominated by cytology and genetics and their use for solving evolutionary and taxonomic problems (Stebbins 1956).

No group of plants has been more radically affected by this new approach than the grass family. Because of the many uses of grasses, classification or taxonomy does more than satisfy the curiosity about the diversity of living things and the way in which they evolved. Cereal and sugar cane breeders have learned that many species of wild grasses are closely related to the cultivated crop species and hybrids between them can be obtained. Therefore, the most useful system of classification is one which reflects as nearly as possible the true genetic and evolutionary relationships of the species.

The use of cytogenetics in plant taxonomy is thus important for the determination of phylogenetic relationships (Celarier 1956). It is, therefore, crucial to know what the basic chromosome numbers are, how they originated and their significance in establishing phylogenetic trends (Celarier 1956). The tribe Andropogoneae has been studied extensively over the last millennium, but there is still an uncertainty about the true base chromosome number (Celarier 1956).

Celarier (1956) suggested that the base chromosome number is five, because of the occurrence of many taxa with the haploid chromosome numbers as multiples of five. Other chromosome numbers suggested for the tribe includes nine (Avdulov 1931), seven (Moriya and Kondo 1950) and six (Janaki-Amal 1940), but they are rare and there are no diploid species known with six or seven as the base number. The most common numbers recorded for the tribe are multiples of ten or five (Darlington and Janaki-Ammal 1945, Delay 1951).

The basic number of five was first recorded by Karper (1929) and later confirmed by several others (Longley 1932, Karper and Chisholm 1936, Hushkins and Smith 1932, 1934, Ayyenger and Ponnaiya 1941, Garber 1947, 1950, Garber and Snyder 1951, Celarier and Harlan 1957, De Wet 1958, Olorode 1975, Dujardin 1978, Spies and Du Plessis 1986 a, b, 1987 a, b, Sinha *et al.* 1990, Spies *et al.* 1994, Strydom *et al.* 2000).

Earlier studies of the tribe Andropogoneae indicated basic chromosome numbers of both five and ten for *Andropogon*, *Chrysopogon*, *Cymbopogon*, *Eulalia*, *Elionurus*, *Imperata*, *Monocymbium*, *Themeda*, *Trachypogon* and *Vetiveria* (Appendix 1).

In *Bothriochloa*, *Dichanthium*, *Diheteropogon*, *Hyparrhenia*, *Ischaemum*, *Microstegium*, *Phacelurus*, *Sorghastrum*, *Urelytrum* and *Zea* only the base number of ten has been observed, which was confirmed by the observation of a high multivalent frequency in *Hyparrhenia tamba* with  $2n = 20$  by Spies and Du Plessis (1987 a, b), whereas in *Coix* and *Sorghum* the base number of five has been confirmed (Appendix A).

*Cleistachne* and *Coelorachis* revealed nine as their basic chromosome number and in 1986, Spies and Du Plessis also observed nine as a basic chromosome number in *Andropogon lacunosus* J.G. Anders.

Both basic numbers of nine and ten have been observed in *Arthraxon*, *Hemarthria* and *Rottboelia* (Appendix 1), whereas nine and 15 have been reported in *Miscanthus* (Avdulov 1931, Bremer 1934, Celarier 1956), and ten and 15 in *Hyparrhenia*. Later studies by Spies and Du Plessis (1987 a, b) also indicated five as the basic number for *Hyparrhenia*. Other numbers include 11 in *Heteropogon*, with aneuploidy suspected (Spies and Du Plessis 1986 a, b, 1987 a, b), 17 and 20 in *Sehima* and seven in *Hackelchloa*.

Heterochromatized knobs are found on the chromosomes of *Zea*, *Tripsacum* and *Coix lacryma-jobi* but not on *Coelorachis cylindrica*, *C. ramosa* or *Elionurus argenteus* (Chandravadana and Galinat 1976). Dennis and Peacock (1984) found close similarities between the knob sequences of *Tripsacum* and *Zea*.

Like many other organisms maize (*Zea*) have supernumary or B chromosomes (Langley 1927, McClintock *et al.* 1981, Jones and Rees 1982, Kellogg and Briclher 1993, Rosato *et al.* 1996 and Chiavarino *et al.* 1998). It is reported that B chromosomes are maintained due to various mechanisms of “drive” consisting of:

1. The suppression of meiotic loss when they are in single dose,
2. Nondisjunction at the second pollen grain mitosis (Roman 1947, Carlson 1978, Carlson and Tau-San Chou 1981, Carlson and Roseman 1992),
3. Preferential fertilization by the sperm nucleus carrying the B chromosomes produced during the nondisjunction process (Roman 1948, Carlson 1969) and
4. A higher competitive ability of B-carrying pollen grains (Beckett 1982).



B chromosomes are not generally present in all individuals of a species or even in a population and their number varies with little or no detectable effect on the phenotype. Cytogenetically all B chromosomes are similar and they are highly heterochromatic. Because of their sporadic occurrence, their origin has been a puzzle. B chromosomes have been observed in various genera of the tribe Andropogoneae (Appendix A).

A phenomena that is commonly observed in Andropogoneae is polyploidy. Polyploids have three or more complete sets of chromosomes in their nuclei instead of the two found in diploids. The level of ploidy can be extremely high. The highest level thus far found in Andropogoneae is 12-ploid (Appendix A).

A particular distinctive feature of the evolution and speciation of flowering plants is allopolyploidy – the combination of different diploid nuclear genomes from two or more different ancestral species or genera. Understanding polyploidy assumes even greater significance when it is considered that most important crops are polyploids (eg. Coffee, cotton, maize, potato, sugarcane and wheat) (Leitch *et al.* 1997).

Novel uses of molecular techniques have opened up powerful ways of investigating the genomic or DNA sequence constitution of polyploids.

In this study the cytogenetic data will be used in conjunction with phylogeny to determine the relationships within the tribe Andropogoneae and to establish any correlation between the polyploid level and phylogeny.

## **1.4 Phylogeny**

Phylogenies are providing major insights into our understanding of evolution (Eggleton and Vane-Wright 1994, Harvey *et al.* 1996, Walsh *et al.* 1999). If the evolutionary history of an organism is traced back it connects through shared ancestors to lineages of other organisms (Lipscomb 1998), therefore all is connected through a phylogenetic cladogram.

The first step of a phylogenetic reconstruction is the identification of plesiomorphic (primitive characters) and apomorphic (derived characters) character states. For phylogenetic reconstructions only synapomorphic characters (shared derived characters) can be used. When these synapomorphic characters are found, the phylogeny is reconstructed step by step. This is the search for a sister group. The concept of sister

groups is independent of extinction. It is thus possible to start with any species or monophyletic group, and search for the sister group by surveying the synapomorphies (Bremer and Wanntorp 1978).

There are two ways of grouping taxa together based on synapomorphies. The Hennig Argumentation (Hennig 1966) and the Wagner Method (Kluge and Farris 1969, Farris 1970). In the Wagner method character states are measured on an interval scale and no “a priori” restrictions are imposed on character state changes. Since this procedure minimizes the number of changes, it also minimizes the number of extra change or homoplasy. Thus, the most parsimonious cladogram is obtained (Swofford and Maddison 1987). More homoplasy in the most parsimonious cladogram do not imply that the data provide less information for choosing among cladograms. The consistency index (CI) is an appropriate measure of homoplasy (=deviation from hierarchy). The retention index (RI) seems more appropriate for fit between different cladograms for the same data set (Goloboff 1991). To search for the most parsimonious cladogram, several methods are used, e.g. exhaustive search, branch and bound and branch swapping.

A consensus cladogram is used to show the information about taxa relationships that all the equally parsimonious cladograms have in common. A consensus cladogram is not a character distribution scheme, but merely indicates common, as well as unshared components. Consensus cladograms are consequently neither parsimonious nor unparsimonious. The cladistic parsimony criterion should only be used in relation to the length of cladograms and not in relation to consensus cladograms, which are without length (Anderberg and Tehler 1990). There is a variety of consensus techniques, including:

- Strict (Sokal and Rohlf 1981, Page 1989), a cladogram that is derived by combining only those components from a set of cladograms that appear in all of the original cladograms,
- the Adams consensus cladogram which places taxa that conflict at the node all the conflicting positions have in common (Adams 1972),
- the Combinable components or Semistrict consensus cladogram that is similar to a Strict consensus cladogram, except it will combine those clades that are not contradicted by all of the cladograms (Bremer 1990) and

- the Majority Rules consensus cladogram, which considers the most frequent placement of a taxon in all the cladograms as its placement in the consensus, cladogram (Swofford 1991).

Of these, the most commonly used is Strict consensus, mainly on the grounds that it is the most conservative (Barret *et al.* 1991). Other criteria than those followed by the Strict consensus may be used to choose among the many equally parsimonious cladograms. Such an example is successive weighting. Homoplastic characters being less reliable are down weighted, hence successive weighting employs evidence that would otherwise be discarded, namely, that the characters themselves indicate that some are less reliable than others (Farris 1969, Carpenter 1994). The weighted characters are then used to create a new cladogram. New unit character consistencies are calculated and the characters re-weighted, until the cladograms on successive iterations are identical. The technique thus selects the most consistent characters (Carpenter 1988).

Heritable characters of taxa are the products of their history and are of value in its elucidations. Thus, different data sets, when subjected to reliable methods of phylogenetic analysis are expected to converge onto the true species phylogeny for their group (Miyamoto and Carcraft 1991, Sheldon and Bledsoe 1993, Miyamoto *et al.* 1994 and Miyamoto and Fitch 1995).

These data sets used must be obtained from the appropriate molecular method. Studies of nucleic acids (DNA, RNA) are the most powerful approach to chemosystematics (the use of chemical constituents of plants to assess inter- and intraspecific relationships and to infer phylogeny) of plants (Woodland 1991).

There are two basic methods used to obtain primary sequence information (Clegg and Darbin 1990, Woodland 1991). Restriction endonucleases can be used for analysis of molecular differences among DNA samples. The other choice is the complete sequencing of genes from representative taxa (Clegg and Darbin 1990).

DNA sequencing is more expensive and requires more sophisticated protocols than does restriction analysis, but it represents a more straightforward approach to comparative systematics and is most valuable for higher-level phylogenetic comparisons (Woodland 1991).

### **1.4.1 Chloroplast DNA in phylogeny**

The variation in chloroplast DNA has proven to be valuable in the reconstruction of plant phylogenies. The chloroplast genome of higher plants consists of circular DNA of about 120-217 kilobase pairs (kb), arranged into two single copy regions divided by inverted repeats. Earlier studies of chloroplast genome evaluation indicated conservation in size, structure, gene content and linear order of genes among major lineages of land plants (Palmer 1985, 1991, Palmer and Stein 1986). This suggested that a change in structure arrangement or content of the chloroplast genome may have significant phylogenetic implications.

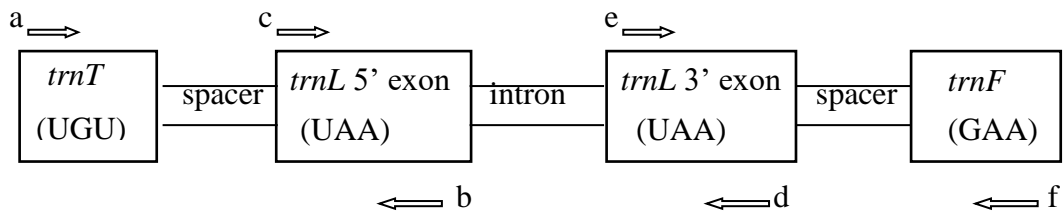
The changes in the genome have been attributed to inversions and additions/deletions that may be found in noncoding DNA, which are used for determining relationships among more closely related species (Palmer *et al.* 1988, Clegg *et al.* 1991, Morton and Clegg 1993). Examples of this strategy used for chloroplast DNA evolution analysis, include studies of:

- the *rbcL* region of the Arundinoideae (Barker *et al.* 1995),
- between *Brassica* species (Lannér 1998), *Korthalsella* species (Molvray *et al.* 1999), Acanthaceae (McDade and Moody 1999), *Pelargonium* species (Bakker *et al.* 1999), *Astragalus* species (Wojciechowshi 1999) and *Doronicum* species (Fernandez *et al.* 2001), based on *trnL-trnF* sequence data,
- the plastid DNA sequences of *Pleurothallidinae* (Priggeon *et al.* 2001), *Cranichideae* with emphasis on *Spiranthinae* (Salazar *et al.* 2003) and from *Cyperaceae* subfamily *Mapanioideae* (Simpson *et al.* 2003),
- The chloroplast genes of the cotton genus *Gossypium: Malvaceae* (Cronn *et al.* 2002),
- chloroplast sequence data of *Capparaceae* and *Brassicaceae* (Hall *et al.* 2002),
- the *rbcL*, *trnL-F* and *ndhF* sequences of *Urticalean rosids* (Sytsma *et al.* 2002),
- the *ndhF* region of *Panicum* (Aliscion *et al.* 2003),

- *rbcL* and *trnL-F* plastid DNA sequences of *Ophioglossaceae* (Hauk *et al.* 2003),
- the DNA sequence variation from three cpDNA fragments: *rbcL*, *atpB* and *trnL-F* of the Hawaiian fern genus *Adenophorus* (Ranker *et al.* 2003).

The *trnL-trnF* region is suitable for evolutionary studies because of the succession of conserved *trn* genes and several base pairs of non-coding regions and the higher rate of molecular evolution of the single-copy regions (Wolfe *et al.* 1987, Taberlett *et al.* 1991).

In 1991 Taberlett *et al.* developed the primers for the amplification of three non-coding regions (Figure 1.8) i) *trnT* (coding for threonine) ii) *trnL* (coding for leucine) and iii) *trnF* (coding for phenylalanine). The first primer pair amplifies the intergenic spacer between *trnT* and *trnL* 5' exon, the second the *trnL* intron and the third pair the intergenic spacer between the *trnL* 3' exon and *trnF*.



**Figure 1.8:** A schematic representation of the *trn* region in the Chloroplast genome with the primers a-f developed for the amplification of this region (Taberlett *et al.* 1991).

Several scientists have observed a faster evolution in the spacer portion of the *trnL-trnF* region, than in the intron (Gielly and Taberlett 1994, 1996, Kita *et al.* 1995, Gielly *et al.* 1996). This suggests a stronger selection for the intron than for the spacer (McDade and Moody 1999).

This combination of regions evolving at different rates increase the phylogenetic range over which these sequences are useful, with the more slowly evolving regions providing support for the older divergences and the more rapidly evolving regions providing support for the closer relatives (McDade and Moody 1999).

In comparison to other chloroplast loci the parsimony-informative variation of *trnL-F* is comparable to *ndhf* in terms of rate of substitutions. These two regions combined, provides twice the rate of informative substitutions as does *rbcL*. The *trnL-F*

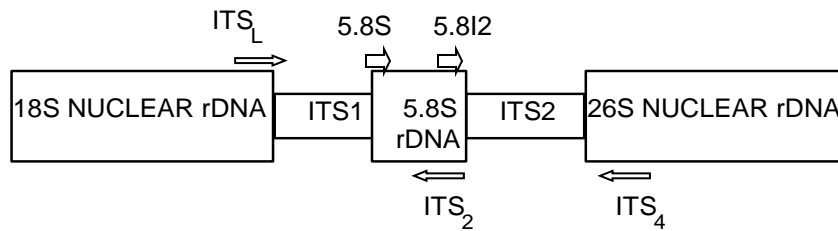
region does, however, exceed the number of informative length mutations within *ndhf* and *rbcL*. This region offer on a whole, more phylogenetically informative variation per nucleotide sequenced than does *ndhf* or *rbcL*, it is also relatively free from homoplasy compared to the other two.

A phylogenetic study on the Andropogoneae, based on *ndhf* sequences has recently been conducted by Spangler *et al.* (1999). For this study, however, the *trnL-trnF* region will be sequenced to determine the relationships within and between the genera of the tribe Andropogoneae and to determine the mutation rate within and between the genera.

### **1.4.2 Nuclear DNA in phylogeny**

Even though plant nuclear DNA is analyzed less easily than cpDNA, it does possess two attributes that make it useful phylogenetically. First, certain nuclear sequences appear to evolve more rapidly than any cpDNA sequences, allowing a finer level of discrimination at the population level than for cpDNA. Second, the nuclear genome is inherited biparently, whereas the chloroplast genome is inherited maternally. Thus, a hybrid plant will possess the nuclear complement of both parents but only the cpDNA of the maternal plant. The nuclear DNA used in such a study must be i) evolutionary conservative, ii) phylogenetically interpretable, iii) easily examined in the laboratory and iv) sufficiently large for phylogenetic reconstruction (Hamby and Zimmer 1992).

An example of this is the internal transcribed spacers (*ITS*) region of 18-26S nuclear ribosomal DNA. This region includes three components (Figure 1.9): the 5.8 S subunit, an evolutionary highly conserved sequence and, importantly from the perspective of this study, two spacer regions designated *ITS1* and *ITS2*. The *ITS* regions are part of the transcriptional unit of nuclear ribosomal DNA (nrDNA), but the spacer segments of the transcript are not incorporated into mature ribosomes. Instead *ITS1* and *ITS2* regions of the nrDNA transcript may function in the maturation of nrRNA's.



**Figure 1.9:** Repeat unit of 18-26S nuclear ribosomal DNA, minus the intergenic spacer and the larger part of the 26S subunit (White *et al.* 1990, Sun *et al.* 1994).

Several aspects of the *ITS* region promote *ITS* use for phylogenetic analysis. Firstly, along with the other components of the nrDNA multigene family, the *ITS* region is one of the most highly repeated sequences in the plant nuclear genome. The entire nrDNA repeat unit, including the subunits, *ITS1*, *ITS2*, the intergenic spacer (*IGS*) and the external transcribed spacer (*ETS*), is present in up to many thousands of copies, arranged in tandem repeats of a chromosomal locus or at multiple loci (Rodgers and Bendich 1987, Hamby and Zimmer 1992). This high copy number promotes detection, amplification, cloning and sequencing of nrDNA. Secondly, and most importantly from the phylogenetic standpoint, this gene family undergoes rapid concerted evolution (Arnheim *et al.* 1981, Hills *et al.* 1991), via unequal crossing over and gene conversion, a property that promotes intragenomic uniformity of repeat unit and accurate reconstruction of species relationships from these sequences (Hamby and Zimmer 1992, Sanderson and Doyle 1992). Thirdly, the small size of the *ITS* region and the presence of highly conserved sequences flanking each of the two spacers makes this region easy to amplify.

The successful sequencing of the *ITS* region is commonly used in phylogenetics (Brochman *et al.* 1998, Emshwiller and Doyle 1998, Downie *et al.* 1998, Fenton *et al.* 1998, Freudenstein 1999, Gernandt and Liston 1999, Hsiao *et al.* 1998, Kelly 1998, Kim *et al.* 1999, Kornkven *et al.* 1998, Alice and Campbell 1999, Douzery *et al.* 1999, Lee and Downie 1999, Li *et al.* 1999, Noyes and Rieseberg 1999, Panero *et al.* 1999, Plunkett and Downie 1999, Steane *et al.* 1999 a, Steane *et al.* 1999b, Vargas *et al.* 1999, Pridgeon *et al.* 2001, Coleman 2002, Nimri *et al.* 2002, Watson *et al.* 2002, Muellner *et al.* 2003, Schnabel *et al.* 2003, Urbatsch *et al.* 2003, van den Heede 2003).

Spies and Kellogg (unpublished data) sequenced the *ITS* region of some representatives of the tribe Andropogoneae and in 1994 Sun *et al.* sequenced this region to infer phylogenetic relationships in *Sorghum* and Buckler and Holstford (1996a,b) also determined the phylogenetic relationships of some species of *Zea*.

In this study we will sequence the *ITS* region of the South African representatives of the tribe Andropogoneae to determine the phylogenetic relationships and to determine the mutation rate within and between the genera of Andropogon.

## **1.5 Aim.**

The morphological variation among members of the tribe Andropogoneae in South Africa provides an interesting context to examine evolutionary relationships within this group. Within the grass family this tribe is considered to be advanced and widespread polyploid. This study is the beginning of research on South African representatives, with the main aim to determine the usefulness of the different genes in providing answers on the South African situation where a lot of variation among the members of the tribe exist.

This study will determine the phylogenetic relationships from chloroplast and nuclear DNA sequences, as well as from the basic chromosome numbers in the tribe. Thus sequencing of the chloroplast region, *trnL-F*, and the nuclear ribosomal DNA, *ITS* region, will be compared with cytogenetic data. The sequences will then be used to determine the divergence within some genera of the tribe Andropogoneae.



# **Chapter 2**

## **Materials and Methods**

### **2.1 Materials.**

The plants used in this study were collected in the veldt. Voucher herbarium specimens (Table 2.1) are housed in the Geo Potts Herbarium (BLFU).

In this study Taq polymerase (Advanced Biotechnology's, Surrey, UK) and DNA Molecular marker VI (pBR328 DNA cleaved with a mixture of *Bgl* I and *Hinf* I) (Boehringer Mannheim Cat. no. 1062590) were used. Different primers were used for *ITS* sequencing (DNAgency Cat. No. GK 071101 and GK 071102) and *trnL-F* sequencing (Whitehead Scientific, 930030, 930031, 930032, 930033). The Thermo Sequence™ dye terminator cycle sequencing pre-mix kit (Amersham Life Science, US 79765 and US 79865), and the BigDye terminator Cycle sequencing ready reaction (PE Applied Biosystems, 4303152) were used to sequence the amplified nrDNA and chloroplast DNA. All the other chemicals used were of analytical grade.

### **2.2 Methods.**

#### **2.2.1 Cytogenetics.**

Young inflorescence were collected and fixed in Carnoy's (1886) fixative. The fixative was replaced with 70% (v/v) ethylalcohol 24-48 hours after fixation. The anthers were squashed in 2% (m/v) aceto-carmine (Darlington & La Cour 1976) on a microscope slide. Improved staining of the chromosomes was achieved by adding iron acetate (Thomas 1940) and heating the slide over a spirit flame. Liquid CO<sub>2</sub> (Bowen 1956) was used to separate the cover slip from the microscope slide for mounting the slides permanently. Removal of the cover slip was followed by dehydration in ethylalcohol and mounting in Euparal. At least twenty cells per specimen were studied for each meiotic stage, except where otherwise indicated.

## **2.2.2 Photography.**

A Nikon microphot-FXa photomicroscope and Ilford Pan-F films, as well as a Nikon Coolpix 990 digital camera were used for the photomicrographs. A selection of the photos is used in this study.

**Table 2.1:** List of specimens studied, their voucher numbers and localities according to the degree reference system (Edwards & Leistner 1971).

---

*Andropogon amethystinus* Steud.

FREE STATE.—2828 (Bethlehem): Witsieshoek vacation resort (-DB), *Spies* 6677, 6679, 6685.

*Andropogon appendiculatus* Nees

MPUMALANGA.—2729 (Volksrust): Hartebeesfontein (-CC), *Spies* 6543.

FREE STATE.—2729 (Volksrust): Giddys Home (-DC), *Spies* 6566. 2827 (Senekal): 33km from Senekal to Ficksburg (-DB), *Spies* 6638. EASTERN CAPE PROVINCE.—3027 (Lady Gray): 35km from Aliwal North to Lady Gray (-CA), *Spies* 6946, 6949.

*Andropogon distachyos* L.

FREE STATE.—2828 (Bethlehem): Witsieshoek vacation resort (-DB), *Spies* 6684; 24km from Fouriesburg to Clarens (-CA), *Spies* 5869.

*Andropogon eucomus* Nees

MPUMALANGA.—2530 (Lydenburg): 32km from Carolina to Belfast (-CC), *Spies* 5107.

FREE STATE.—2829 (Harrismith): Sterkfontein (-CA), *Spies*, 6587.

*Andropogon schirensis* A. Rich

FREE STATE.—2827 (Senekal): 5km from Nebo to Fouriesburg via Generaalsnek (-DB), *Spies* 4821. 2829 (Harrismith): Sterkfontein (-CA), *Spies* 5084, 6581.

KWAZULU/NATAL.—2729 (Volksrust): 99km from Harrismith to Normandien (-DC), *Spies* 5084.

*Aristida adscensionis* L. (Trin. & Rupr.) Henrard

FREE STATE.—2926 (Bloemfontein): 2km from Reddersburg to Smithfield (-CA), *Spies* 6713.

*Arthraxon lanceolatus* (Roxb.) Hochst. var. *lanceolatus*

MPUMALANGA.—2430 (Pilgrim's Rest): On top of Blyderiviers Poort (-DB), *Spies 1424*.

*Bothriochloa bladhi* (Retz.) S.T. Blake

MPUMALANGA.—2530 (Lydenburg): 27km from Lydenburg to Machadodorp (-AD), *Spies 5143*.

FREE STATE.—2926 (Bloemfontein): UOFS campus (-AA), *Spies 5173*.

*Bothriochloa insculpta* (A. Rich.) A. Camus

GAUTENG.—2528 (Pretoria): Soutpan experimental farm (-AC), *Spies 3321*.

NORTHERN CAPE.—2820 (Kakamas): Augrabies waterfall (-CB), *Spies 5681*. 2824 (Kimberley): 21km from Kimberley to Griekwastad (-DA), *Spies 7489*.

KWAZULU/NATAL.—2832 (St. Lucia): 22km from Cape Vidal to St. Lucia (-AD), *Spies 2407*.

*Bothriochloa radicans* (Lehm.) A. Camus

FREE STATE.—2827 (Senekal): 6km from Nebo to Fouriesburg (-DB), *Spies 4831*. 2927 (Maseru): 26km from De Wetsdorp to Hobhouse (-DB), *Spies 4784*.

*Brachiaria serrata* (Thunb.) Stapf syn.

FREE STATE.—2828(Bethlehem): 10km from Fouriesburg to Clarens (-CA), *Spies 6654*.

*Cymbopogon dieterlenii* Stapf ex Phill.

FREE STATE.—2827 (Senekal): 41km from Rosendal to Ficksburg via Nebo (-DB), *Spies 6642*.

*Cymbopogon excavatus* (Hochst.) Stapf ex Burt Davy

FREE STATE.—2828 (Bethlehem): 34km from Fouriesburg to Clarens (-CA), *Spies 5877*. 2829 (Harrismith): Sterkfontein (-CA), *Spies 6577*.

*Cymbopogon marginatus* (Steud.) Stapf ex Burt Davy

WESTERN CAPE.—3219 (Wupertal): On top of Uitkyk Pass (-AC), *Spies 4397*. 3420 (Bredasdorp): 1km from De Hoop (-AB), *Spies 5051*.

*Cymbopogon plurinodis* (Stapf) Stapf ex. Burt Davy

GAUTENG.—2528 (Pretoria): Soutpan experimental farm (-AC), *Spies 3296, 3322*.

FREE STATE.—2827 (Senekal): 6km from Nebo to Fouriesburg (-DB), *Spies 4842*. 2926 (Bloemfontein): UOFS campus (-AA), *Spies 6404, 6406*. 2927 (Maseru): 19km from De Wetsdorp to Hobhouse (-DB), *Spies 4772*.

EASTERN CAPE PROVINCE.—3324 (Steytlerville): 53km from Patensie to Willowmore in Baviaanspoort (-CB), *Spies 5206*.

*Cymbopogon prolixus* (Stapf) Phill.

MPUMALANGA.—2629 (Bethal): 46km from Ermelo to Breyton (-DB), *Spies 5101*.

FREE STATE.—2828 (Bethlehem): Witsieshoek vacation resort (-DB), *Spies 6682*.

*Cymbopogon* sp.

WESTERN CAPE.—3119 (Calvinia): 5km from Algiers to Citrusdal in Nieuwoudt Pass (-CA), *Spies 5800*.

*Digitaria tricholaenoides* Stapf

FREE STATE.—2826 (Brandfort): Glen camp (-CD), *Spies 6691*. 2827 (Senekal): Nebo (-DB), *Spies 6636*.

*Diheteropogon amplexans* (Nees) Clayton

FREE STATE.—2828 (Bethlehem): 19km from Fouriesburg to Clarens (-CA), *Spies 4821*.

*Diheteropogon filifolius* (Nees) WD Clayton

FREE STATE.—2828 (Bethlehem): 34km from Fouriesburg to Clarens (-CA), *Spies 5872, 5874*.

*Elionurus muticus* (Spreng.) Kunth

SWAZILAND.—2631 (Mbabane): Sebebe Hill, 10 km North-east of Mbabane (-AC), *Spies 2537*.

MPUMALANGA.—2629 (Bethal): 46km from Ermelo to Breyten (-DB), *Spies 5104*. 2729 (Volksrust): Hartebeesfontein (-CC), *Spies 6539*.

FREE STATE.—2729 (Volksrust): Giddys Home (-DC), *Spies 6559*. 2828 (Bethlehem): 10km from Clarens to Golden Gate (-DA), *Spies 6664*. 2927 (Maseru): 42km from De Wetsdorp to Hobhouse (-CA), *Spies 4792*.

KWAZULU/NATAL.—2729 (Volksrust): 93km from Harrismith to Normandien (-DC), *Spies 5074*.

EASTERN CAPE PROVINCE.—3026 (Aliwal North): 35km from Lady Gray to Aliwal North (-AA), *Spies 6945*. 3027 (Lady Gray): 28km from Barkley East to Rhodes (-DC), *Spies 6054*.

*Eulalia villosa* (Thunb.) Nees

KWAZULU/NATAL.—2729 (Volksrust): 99km from Harrismith to Normandien (-DC), *Spies 5085*.

*Helictotrichon longifolium* (Nees) Schweick.

FREE STATE.—2729 (Volksrust): Giddys Home (-DC), *Spies 6560*.

*Hemarthria altissima* (Poir.) Stapf & C. E. Hubb.

SWAZILAND.—2631 (Mbabane): Motshane 16km from Mbabane to Oshoek (-AC), *Spies 5698*.

*Heteropogon contortus* (L.) Roem. & Schult.

MPUMALANGA.—2729 (Volksrust): Hartebeesfontein (-CC), *Spies* 6538.

FREE STATE.—2729 (Volksrust): Giddys Home (-DC), *Spies* 6552. 2827 (Senekal): 33km from Senekal to Rosendal (-BC), *Spies* 6628; 6km from Nebo to Fouriesburg (-DB), *Spies* 4841.

EASTERN CAPE PROVINCE.—3026 (Aliwal North): 15km from Lady Gray to Aliwal North (-AA), *Spies* 4271, 4741; 10km from Lady Gray to Aliwal North (-AA), *Spies* 6942.

*Hyparrhenia anamesa* Clayton

FREE STATE.—2827 (Senekal): 8km from Clocolan (-DC), *Spies* 4807.

*Hyperthelia dissoluta* (Nees ex Steud.) Clayton syn.

MPUMALANGA.—2530 (Lydenburg): 27km from Lydenburg to Machadodorp (-AD), *Spies* 5145.

*Hyparrhenia filipendula* (Hochst.) Stapf. var. *filipendula*

MPUMALANGA.—2530 (Lydenburg): 27km from Lydenburg to Machadodorp (-AD), *Spies* 5144.

*Hyparrhenia hirta* (L.) Stapf

MPUMALANGA.—2530 (Lydenburg): 5km from Belfast to Dullstroom (-CA), *Spies* 5112. 2629 (Bethal): 9km from Ermelo to Breyten (-DB), *Spies* 5100.

FREE STATE.—2827 (Senekal): 5km from Nebo to Fouriesburg via Generaalsnek (-DB), *Spies* 4822, 15km from Senekal to Rosendal (-BC), *Spies* 6618. 2828 (Bethlehem): 10km from Clarens to Golden Gate (-DA), *Spies* 6656.

KWAZULU/NATAL.—2729 (Volksrust): 97km from Harrismith to Normandien (-DC), *Spies* 5079.

EASTERN CAPE PROVINCE.—3324 (Steytlerville): 7km from Humansdorp to Patensie (-DD), *Spies* 5184.

*Hyparrhenia pilgeriana* C. E. Hubb.

EASTERN CAPE PROVINCE.—3027 (Lady Gray): 15km from Barkley East to Lady Gray (-DC), *Spies* 4738.

WESTERN CAPE PROVINCE.—3319 (Worcester): After tunnel on Du Toitskloof Pass (-CA), *Spies* 4603. 3419 (Caldon): Caledon (-AC), *Spies* 4649. 3420 (Bredasdorp): 4km from Bredasdorp to Swellendam (-CA), *Spies* 4851.

*Hyparrhenia quarrei* Robyns

FREE STATE.—2829 (Harrismith): Sterkfontein (-CA), *Spies* 6586

KWAZULU/NATAL.—2729 (Volksrust): 106km from Harrismith to Normandien (-DC), *Spies* 5088.

*Hyparrhenia tamba* (Steud.) Stapf

MPUMALANGA.—2729 (Volksrust): Hartebeesfontein (-CC), *Spies* 6537.

FREE STATE.—2829 (Harrismith): Sterkfontein (-CA), *Spies* 6585.

*Hyparrhenia rudis* Stapf

FREE STATE.—2829 (Harrismith): Sterkfontein (-CA), *Spies* 6576.

*Hyparrhenia* sp.

WESTERN CAPE.—3420 (Bredasdorp): 11km from Bredasdorp to Cape Agullas (-CA), *Spies* 5033.

*Imperata cylindrica* (L.) Raeuschel

MPUMALANGA.—2530 (Lydenburg): 5km from Belfast to Dullstroom (-CA), *Spies* 5114.

FREE STATE.—2829 (Harrismith): Sterkfontein (-CA), *Spies* 6582. 2827 (Senekal): Nebo (-DB), *Spies* 6632.

KWAZULU/NATAL.—2729 (Volksrust): 10km from Newcastle to Cundycleugh (-DD), *Spies* 5090.

*Ischaemum fasciculatum* Brongn.

SWAZILAND.—2631 (Mbabane): Sebebe Hill (-AC), *Spies* 2563.

KWAZULU/NATAL.—2832 (Mtubatuba): 6km from Hluhluwe to Sordwana (-AA), *Spies* 2358.

*Miscanthus capensis* (Nees) Anderss.

FREE STATE.—2829 (Harrismith): Sterkfontein (-CA), *Spies* 6584. 2828 (Bethlehem): 10km from Clarens to Golden Gate (-DA), *Spies* 6655.

*Monocymbium cerasiiforme* (Nees) Stapf

KWAZULU/NATAL.—2829 (Harrismith): Oliviershoek Pass (-CA), *Spies* 6578.

*Panicum schinzii* Hack.

FREE STATE.—2925 (Jagersfontein): 46km from Petrusburg to Kimberley (-AB), *Spies* 6696.

*Paspalum dilatatum* Poir.

FREE STATE.—2827 (Senekal): 20km from Senekal to Rosendal (-BC), *Spies* 6617.

*Phacelurus franksae* Griseb.

EASTERN CAPE PROVINCE.—3028 (Matatiele): 78km from Rhodes (-CC), *Spies* 4705.

*Setaria* sp.

FREE STATE.—2729 (Volksrust): Giddys Home (-DC), *Spies* 6567.

*Sorghum bicolor* (L) Moench subsp. *arundinaceum* (Desv.) De Wet & Harlan

FREE STATE.—2926 (Bloemfontein): UOFS grounds (-AA), *Spies* 6588, 6589.

*Themeda triandra* Forssk.

MPUMALANGA.—2530 (Lydenburg): Hartebeesfontein (-CC), *Spies* 6525.

FREE STATE. —2729 (Volksrust): Giddys Home (-DC), *Spies* 6555. 2926 (Bloemfontein): UOFS campus (-AA), *Spies* 6382. 2828 (Bethlehem): 10km from Clarens to Fouriesburg (-CB), *Spies* 6652.

EASTERN CAPE PROVINCE.—3027 (Lady Gray): Near Dordrecht on the way to Lady Gray (-DC), *Spies* 2489.

*Trachypogon spicatus* (L. f.) Kuntze

FREE STATE.—2828 (Bethlehem): 19km from Fouriesburg to Clarens (-CA), *Spies* 4846. 2829 (Harrismith): Sterkfontein (-CA), *Spies* 6580.

KWAZULU/NATAL.—2829 (Harrismith): Oliviershoek Pass (-CA), *Spies* 6579.

WESTERN CAPE.—3321 (Ladysmith): 62km from Barrydale to Riversdal over Garcia Pass (-CC), *Spies* 5205, 5480.

*Tristachya leucothrix* Nees

FREE STATE.—2729 (Volksrust): Giddys Home (-DC), *Spies* 6549. 2827 (Senekal): 33km from Senekal to Rosendal (-BC), *Spies* 6627. 2828 (Bethlehem): 38km from Clarens to Phutaditjhaba via Golden Gate (-DA), *Spies* 6671.

*Urelytrum agropyroides* (Hack.) Hack.

GAUTENG.—2528 (Pretoria): 1km from Cullinan to Pretoria (-DA), *Spies* 2081.

### **2.2.3 DNA extraction.**

The leaves of the different specimens (Table 2.1) were collected and stored in a saturated sodium chloride (NaCl) and hexadecyltrimethylammonium bromide (CTAB) solution (Rogstad 1992). The leaves were washed with distilled water and cut into pieces. One gram of the material was ground to a fine powder in liquid nitrogen. The tissue was immediately resuspended in warm (65°C) 1% CTAB (500µl) extraction buffer

[1% (m/v) CTAB, 50mM Tris-HCl (2-amino-2(hydroxymethyl)-1,3-propanediol) (pH 8,0), 10mM ethylene diaminetetra-acetic acid (EDTA), 0.7M NaCl (Hills & Moritz 1990), to which 1% (v/v)  $\beta$ -mercapto-ethylalcohol had been added prior to use]. This mixture was incubated for half an hour at 65°C. After incubation the same volume of chloroform:iso-amylalcohol (24:1) was added to the mixture and thoroughly mixed. This solution was centrifuged at 3 000g for five minutes. The supernatant (which contained the DNA) was transferred to a clean eppendorf tube. Ethylalcohol (two times the volume of the supernatant), containing 3M sodium acetate (25:1), was added to the supernatant to precipitate the DNA at -20°C for half an hour. This eppendorf tube was centrifuged at 10 000g for 10 minutes. The supernatant was discarded and the pellet washed with 70% (v/v) ethylalcohol, containing 10mM ammonium acetate. This pellet was air dried and dissolved in sterile water (Rogstad 1992). The DNA was visualised by intercalating it with Ethidium Bromide and studying it on an ultraviolet (UV) light illuminator.

For the extraction of DNA from herbarium specimens the same method was used but a 4% CTAB solution was used instead of the 1% CTAB as previously mentioned, with 0.5% PVP (Polyvinyl-Pyrrolidone) added to the CTAB extraction buffer.

### **2.2.4 Gel electrophoresis.**

An agarose gel [0.8% for genomic DNA] was prepared by melting the desired amount of electrophoresis grade agarose in 1x TAE buffer [Tris (48.44g), acetic acid (11.42ml) and EDTA (2.92g) (pH 8.0)]. This mixture was cooled down to 55°C and Ethidium Bromide was added to a final concentration of 0.05  $\mu$ g/ml. Electrophoresis was done in 1x TAE running buffer.

The DNA samples (6  $\mu$ l) were mixed with 2  $\mu$ l 6x loading buffer [0.25% (m/v) bromophenol blue, 0.25% (m/v) xylene cyanol and 40% (m/v) glycerol in water] and loaded into the slots of the submerged gel. The DNA fragments were separated at 80V for 30 minutes and examined under ultraviolet (UV) light.



## **2.2.5 Taguchi optimisation.**

The optimisations of the PCR based sequencing analysis were done according to the Taguchi method (Cobb & Clarkson 1994). According to this method, four reaction component optima can be assessed by using nine reactions. Each reaction component is varied in an orthogonal array by three different concentrations of each variable (Table 2.2). The primer, dNTP, MgCl<sub>2</sub> and DNA concentrations were the variable components and DNA polymerase and buffer concentrations were kept constant.

**Table 2.2:** The orthogonal arrangement of the components to be optimised by the Taguchi method (Cobb & Clarkson 1994). Three different volumes of the four components, which are to be optimised, are varied in an orthogonal array.

Reactions	[Primer] 3.5pmol/ µl	[dNTP] 2mM/ µl	[MgCl <sub>2</sub> ] 25mM/ µl	[DNA] 1:40/ µl
1	1	2	3	2
2	2	3	3	2.5
3	3	4	3	3
4	1	3	4	3
5	2	4	4	2
6	3	2	4	2.5
7	1	4	5	2.5
8	2	2	5	3
9	3	3	5	2

## **2.2.6 PCR Sequencing.**

The *trnL-F*, as well as the *ITS1* and *ITS2* regions, were amplified by using the polymerase chain reaction (PCR). For *ITS* sequencing 40 cycles of 94°C for 30 seconds, 50°C for 30 seconds and 72°C for 90 seconds were applied. Amplification of *ITS1*

(Figure 1.1) was obtained by using the primers ITS<sub>L</sub> and ITS<sub>2</sub> {ITS<sub>L</sub>—5'-TCG TAA CAA GGT TTC CGT AGG TG-3' and ITS<sub>2</sub>—5'-GCT GCG TTC TTC ATC GAT GC-3'} (White *et al.* 1990). The ITS2 region was amplified using the 5.8S or the 5.8I2 in combination with ITS<sub>4</sub> primers {5.8S—5'-ACGACTCTCGGCAAC-3', 5.8I2—5'-GCCTGGGCGTCACGC-3' and ITS<sub>4</sub>—5'-TCC TCC GCT TAT TGC TAT GC-3'} (White *et al.* 1990, Sun *et al.* 1994).

For *trnL-F*, primers c and d {c—5'-CGAAATCGGTAGACGCTACG-3' and d—5'-GGGGATAGAGGGACTTGAAC-3'} (Taberlet *et al.* 1991) were used to amplify the *trnL* (UAA) intron and primers e and f {e—5'-GGTTCAAGTCCCTCTATCCC-3' and f—5'-ATTTGAACTGGTGACACGCG-3'} (Taberlet *et al.* 1991) for the intergenic spacer between the *trnL* (UAA) 3' exon and *trnF* (GAA).

The amplified DNA template was diluted with sterile water (1:60; ~5ng). A total volume of 11µl of this DNA template was mixed with 1µl of one of the primers and 8µl of the sequence reagent pre-mix. This total reaction volume of 20µl was amplified in a Perkin Elmer thermal cycler. This reaction consists of 94°C for 60 seconds and then 25 cycles of 94°C for 30 seconds, 45°C for 15 seconds and 60°C for 4 minutes. Single stranded DNA was subsequently produced using the double stranded DNA templates and only one primer. Single stranded DNA fragments were purified according to the following protocol of the Thermo Sequence™ dye terminator cycle sequencing pre-mix kit: a volume of 68µl of concentrated ethylalcohol (-20°C) was added to each reaction and mixed using a vortex and placed on ice for 15-20 minutes to precipitate the DNA. This solution of DNA and ethylalcohol was centrifuged for 15 minutes at 10 000 g and the supernatant discarded. The pellet was washed with 250-500µl of 70% (v/v) ethylalcohol (-20°C) and briefly centrifuged. The supernatant was discarded and the pellets were vacuum dried for 2-5 minutes. The pellets were resuspended in 4µl of formamide loading dye. Each sample was heated to 70°C for 2-5 minutes and then placed on ice. The entire sample was loaded onto the sequencing gel and run on the ABI 377 automated sequencer for four to six hours.

## **2.3 Statistical analysis.**

### **2.3.1 Analysis of Taguchi products.**

Using the Taguchi method, the product yield for each reaction was used to estimate the effects of individual components on amplification. This is done by using the quadratic loss function, which Taguchi refers to as signal-to-noise ratios:

$$\text{SNL} = -10 \log [1/n \sum 1/y_i^2],$$

where SNL is the signal-to-noise ratio, n is the number of levels and y is the yield. The yield was calculated according to the equation  $P = (r \times s) + 1$ , where P is the product yield, r is the number of products (in this case  $r = 1$ ) and s is the size range. For each component the optimal conditions are those that give the largest SNL. The reactions are further refined by using the polynomial regression from SNL values for each component to obtain curves, the maximum of which represent the reaction optima.

### **2.3.2 Computer analysis.**

For phylogenetic analyses the computer program Phylogenetic Analysis using Parsimony (PAUP) version 4.0 \* beta version (Swofford 1998), which connect taxa, one at a time, until all the taxa have been added, were used. The heuristic search option with branch swapping was used. Each heuristic search comprised 500 replicates of random taxon addition with tree bisection-reconnection (TBR) branch swapping and MULPARS selected. A bootstrap was done from 1000 replications, using the heuristic option (Felsenstein, 1985). Relative levels of support for clades present in the most parsimonious trees were assessed by calculating decay values (Bremer 1988). A search for multiple islands of parsimony was also conducted in cases where the cladogram yielded a RI value of less than 0.67 (Maddison 1991).

Most sequence analyses were carried out using programs designed for, or adapted

to, the Apple Macintosh. Basic analysis was carried out using the Sequence Navigator DNA and Protein Sequence Comparison Software (Applied Biosystems 1994). The sequence of *ITS1*, *ITS2* and *trn L-F* regions were aligned using the computer program Clustal W version 1.6 (Thompson *et al.* 1994). Final alignment was visually examined and manually optimised for phylogenetic analysis, using Clustal X (Thompson *et al.* 1997).

### **2.3.3 Sequence divergence.**

The rate of sequence divergence of all genes and spacers were measured using the formula by O'Donnell (1992).

$$\text{Percentage Divergence} = \{(\text{TS} + \text{TV} + \text{I}\backslash\text{D}) / \text{sequence length}\} \times 100$$

Where:

TS = Transitions

TV = Transversions

I\backslashD = insertion\deletion; multi-base differences are scored as 1.

This formula permits a comparison of the phylogenetic informations on the relevant sequences (Gielly and Taberlet 1994).

# **Chapter 5**

## **General Conclusions**

The tribe Andropogoneae with its morphological variation was an interesting context to examine. Previously the evolutionary pattern in the tribe has been derived from phenetic (Clayton 1972, 1973) and cytotoxic (Celarier 1956-1959) studies.

The most characteristic feature of this tribe is the possession of fragile racemes bearing paired spikelets (Le Roux and Kellogg 1999), and inflorescence structure (Clayton 1972, 1973). The original classification of the tribe was not surprisingly done in accordance to these morphological features. The development of molecular systematics brought new insights into the classification of Andropogoneae. In 2001 the GPWG used molecular data to reclassify the grass family (Poaceae) and the tribe Andropogoneae. This tribe is however complex and everyday new data becomes available that can help answer taxonomic questions.

In 1999 Spangler *et al.* studied the *ndhF* region of Andropogoneae and in 2002 Mathews *et al.* combined the *phytochrome B*, *GBSSI* and *ndhF* data. This data supported the monophyly of the Andropogoneae and a sister relationship with *Arundinella* and suggest that the tribe Arundinelleae should be abandoned as a taxonomic group. This tribe is still recognized in the GPWG study in 2001. The *ITS* and *trnL-F* data in this study further support the sister relationship between Andropogoneae and Arundinelleae and the suggestion that *Arundinella* should be placed within the Andropogoneae.

The previous data sets do, however, support the division of the tribe into awned and awnless genera (Clayton 1972, 1973), whereas in this study, with the South African representatives, this feature is polyphyletic and the groupings are not supported.

The small number of samples per specimen in the previous studies could be responsible for this difference as sampling within a monophyletic group may improve the phylogenetic accuracy (Rosenberg and Kumar 2001), however sampling outside the group “pushes the common ancestor of the new data set” back in time and could decrease accuracy. Random sampling of additional taxa is also thought to decrease phylogenetic accuracy.

Analyzing a single large data set, consisting of a combination of different data sets versus analyzing the individual data sets and then examining the trees for congruence is a topic of contemporary debate in systematics. Mathews *et al.* (2002) combined the *GBSSI*, *PHYB* and *ndhF* data sets. This phylogeny resulted in strong support for clades, which rejected the subtribal classification of Clayton and Renvoize (1986). The combined *ITS* and *trnL-F* data of the South African representatives also rejected the subtribal classifications. Furthermore this data suggest that most genera in the tribe are polyphyletic. In 2002, Hodkinson *et al.* studied the *ITS* and *trnL-F* regions in *Mischanthus* and *Saccharum* and also found these genera to be polyphyletic. This polyphyletic nature of the tribe has been hypothesized by Mathews *et al.* (2002) and is now confirmed.

These variations within a genus can be explained by hybridization events. Since allopolyploidy was indicated by cytological studies (Chapter3), these hybridization events are further supported, suggesting the tribe is much more complex than originally thought. Allopolyploidy has generally been agreed to be widespread in plants. Because rDNA is biparentally inherited, additively of parental rDNA patterns is expected in hybrids and allopolyploids, as was observed in the *ITS* analysis, e.g. *Hyparrhenia* and *Bothriochloa bladhii*. A lack of additively was however previously observed in *Zea* (Zimmer *et al.* 1988), this is however a disadvantage for the molecular analysis of the origins of polyploid species. In this study we were able to demonstrate reticulate evolution in the tribe Andropogoneae in South Africa, because gene conversion apparently did not eliminate the divergent alleles in the allopolyploids.

The hybridization events also give an answer to the basic chromosome number of the tribe Andropogoneae. The event of  $n = 5 + 5$  is a simpler explanation for  $n = 10$  than an aneuploid event leading from  $n = 9$  to  $n = 10$ .

The phylogeny could not prove the basic chromosome number of  $n = 5$  for the tribe Andropogoneae, however the lower polyploid levels were concentrated at the base of the tree, supporting the maturity of the polyploid complex. Therefore, systematics rejects the hypothesis of a basic number of  $n = 10$  with  $n = 5$  derived.

The “core Andropogoneae” group as observed by Spangler *et al.* (1999) and Mathews *et al.* (2002) is not supported by *ITS* and *trnL-F* data sets. Strong evidence of a sister lineage between *Hyparrhenia* and *Andropogon* exists. This relationship was also

noted in previous studies. The genes, ITS and trnL-F proved to be useful in delimiting various relationships in the tribe with the South African representatives.

This study only included specimens from South Africa, increased taxon sampling from especially southern Asia, which is regarded as the region containing the most primitive members of the tribe, will help elucidate the relationships in the tribe. Future studies need to concentrate on the gathering of as much data as possible from all sources available, including morphological characters. Most of what we know about phylogeny is based on morphology and this has been confirmed time and again by molecular studies (Donnoghue and Sanderson 1992). It is thus a mistake to set morphological data aside and base phylogenetic relationships only on molecular evidence.

The study of Andropogoneae is of importance to all mankind. They affect the environment, acting to prevent soil erosion. They are used in pharmaceuticals and most important as a food source, yet uncertainty remains about the number of genes present and the evolutionary patterns. To meet the demands and challenges of the world, we need to increase our knowledge of plant genes in order to manage the environment and food supply.

## **Chapter 6**

### **Summary**

The tribe Andropogoneae makes up half of the grass subfamily Panicoideae, with approximately 85 genera and 960 species (Hartley 1958, Clayton and Renvoize 1986). The genera of the tribe are typically tropical with only a few species extending beyond the tropics into warm temperate regions. From information available it is clear that the African representatives form an integral part of the tribe. In this study we only concentrated on the South African representatives of the tribe.

The tribe Andropogoneae has been studied extensively over the last millennium, but there is still an uncertainty about the true basic chromosome number. Previous molecular studies include sequencing of the *ndhF*, *GBSSI* and *phytochrome B*.

The morphological variation in the tribe provides an interesting context to examine. This study focused on the sequencing of plastid chloroplast gene *trnL-F* and nuclear ribosomal DNA the Internal Transcribed Spacer (*ITS*) to determine the phylogenetic relationships within the tribe.

In this study the chromosome numbers of 58 specimens were determined. The genetic chromosome numbers varied with  $n = 5, 9, 10, 10, 11, 20, 30$ . For the first time the basic diploid number of  $n = 5$  was observed in the genus *Andropogon*. *Urelythrum aggrophyroides* was studied for the first time with a somatic chromosome number of  $2n = 20$ .

The absence of multivalent and prevalence of bivalents in this study indicate that the genomes of the specimens studied are homologous. This lead us to the conclusion that the tribe consists of allopolyploid species, derived from interspecific hybridization and chromosome doubling.

Both the *trnL* intron and *trnL-F* intergenic spacer was sequenced. Analysis of the *trnL-F* gene included 56 accessions and 61 accessions for the *ITS* gene. Combined analysis of both *ITS* and *trnL-F* included 59 accessions. Sequencing data indicated *Cymbopogon* to be a diploid progenitor of *Hyparrhenia* and *Andropogon* of *Bothriochloa bladhii*. Reticulate evolution was demonstrated in nr*ITS* alleles. *ITS* and *trnL-F* data supports the inclusion of *Arundinella* in Andropogoneae and rejects the subtribal



classification of Clayton and Renvoize (1986). This data also does not support the division of the tribe into awned and awnless genera and does not support the “core Andropogoneae” lineage as previously been observed. Most genera in the tribe are polyphyletic, suggesting a much more complex nature for the South African representatives.

Molecular data could not prove the basic chromosome number but supported the maturity of the polyploid complex for the South African specimens.

**Keywords:** Poaceae, Panicoideae, Andropogoneae, cytogenetics, polyploidy, allopolyploidy, DNA sequencing, trnL-F, Internal transcribed spacer (ITS), reticulate evolution.

# **Chapter 7**

## **Opsomming**

Die tribus Andropogoneae vorm die helfte van die gras subfamilie Panicoidae, met ongeveer 85 genera en 960 spesies (Hartley 1958, Clayton and Renvoize 1986). Die genera in die tribus is gewoonlik tropies, met slegs 'n paar wat buite die tropiese gebiede na gebiede met 'n gematigde klimaat versprei het. Die Afrika verteenwoordigers vorm 'n integrale deel van die tribus. Hierdie studie konsentreer hoofsaaklik op die Suid-Afrikaanse verteenwoordigers van die tribus.

Die tribus Andropogoneae is dikwels bestudeer gedurende die laaste millenium, maar daar bestaan nog steeds onsekerheid oor die ware basiese chromosoomgetal van die tribus. Vorige molekulêre studies het die nukleotiedvolgordebepaling van *ndhF*, *GBSSI*, en *phytochrome B* ingesluit.

Die morfologiese variasie in die tribus verskaf 'n interessante konteks om te bestudeer. Hierdie studie het hoofsaaklik gekonsentreer op die nukleotiedvolgordebepaling van die chloroplaste gebied *trnL-F* en die kern ribosomale DNA se intern getranskribeerde streke (*ITS*), om die filogenetiese verwantskappe binne die tribus vas te stel.

Tydens hierdie studie is die chromosoomgetalle van 58 eksimplare bepaal. Die gametiese chromosoomgetalle het gewissel tussen  $n = 5, 9, 10, 11, 20, 30$ . Die basiese diploïede getal van  $n = 5$  is vir die eerste keer in die genus *Andropogon* waargeneem. *Urelythrum aggrophyroides* is vir die eerste keer betudeer, met 'n somatiese chromosoom getal van  $2n = 20$ .

Die afwesigheid van multivalente en die teenwoordigheid van bivalente in hierdie studie is 'n aanduiding dat die genome van die spesies in die studie homeologies is. Dit dui daarop dat die tribus bestaan uit allopoliploïede spesies, wat afgelei is vanaf interspesifieke verbastering en chromosoomverdubbeling.

Beide die *trnL* intron en die *trnL-F* intergeniese streek se nukleotiedvolgorde is bepaal. Analises van die *trnL-F* geen het 56 eksimplare ingesluit terwyl 61 eksimplare vir *ITS* volgorde bepaling gebruik is. 'n Gekombineerde analise van beide *ITS* en *trnL-F* het 59 eksimplare ingesluit. Nukleotiedvolgordebepaling dui aan dat die genus

*Cymbopogon* 'n diploïede voorganger van *Hyparrhenia* is en *Andropogon* van *Bothriochloa bladhii*.

Retikulerende evolusie kom voor in die nrITS allele. ITS en *trnL-F* data ondersteun die insluiting van *Arundinella* in die tribus Andropogoneae en verwerp die subtribus klassifikasie van Clayton en Renvoize (1986). Hierdie data ondersteun ook nie die verdeling van die tribus in geangelde en angellose genera en verder ondersteun dit ook nie die suiwer basiese Andropogoneae groepering soos in vorige studies waargeneem nie. Die meeste genera in die tribus is polifeleties, wat 'n baie meer komplekse samestelling vir die Suid-Afrikaanse verteenwoordigers van die tribus voorstel.

Molekulêre data het nie bewyse gelewer vir die ware basiese chromosoomgetal nie maar het wel die volwasse aard van die poliploïede kompleks vir die Suid-Afrikaanse spesies ondersteun.

**Sleutel woorde:** Poaceae, Panicoideae, Andropogoneae, sitogenetika, poliploïede, allopoliploïede, DNA nukleotiedvolgordebepaling, *trnL-F*, Intergeniese getranskribeerde streke (ITS), retikulerende evolusie.

# **Chapter8**

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**Appendix A:** List of the polyploid levels in Andropogoneae from all existing chromosome data.

Species	2n	Reference
Andropogon		
africanus Franch	40	Dujardin 1979
amethystinus Steudel	c.30	Hoshino & Davidse 1988
	40	Spies et al. 1994
angustatus (Presl) Steud	40	Pohl & Davidse 1971
appendiculus Nees	20, 40, 60	Spies et al 1994
arctatus Chapman	20	Campbell 1982
arenarius Hackel	60	Campbell & Windisch 1986
ascinodis C.B. Clarke	40	Mehra 1982
bicornis L.	30	Davidse & Pohl 1972
	60	Pohl & Davidse 1971, Norrmann 1985, Campbell & Windisch 1986
brachystachyus Chapman	20	Campbell 1983
brazzae Franch	20	Dujardin 1979
canaliculatus Schumach.	20	Tateoka 1965c
Chinensis (Nees) Merr.	20, 40	Spies et al. 1994
distachyos L.	36	Faruqi et al. 1987
eucomus Nees	40	Spies & Du Plessis 1986, 1987
	20	Spies & Du Plessis 1987, Spies et al. 1991, Spies et al. 1994
exaratus	60, 60 + 2B	Norrmann 1985
floridanius Scribner	20	Campbell 1983
gayanus Kunth	20	Tateoka 1965c
	40	Okoli & Olorode 1983
gerardii Vitman	40	Davidse & Pohl 1972
	60	Gould 1968, Riley & Vogel 1982, Keeler et al. 1987
	70, 80	Keeler et al. 1987
	gam 30II, 5I+12II+8III +5IV+1V+2	Norrmann et al. 1997
glaziovii Hackel	VI	Campbell & Windisch 1986
glomeratus (Walt.) BSP	60	Gould & Soderstrom 1970
	20	Davidse & Pohl 1972
glomeratus (Walter) BSP var. glomeratus Mohr	10	Campbell 1983
“ var. hirsutior (Hackel)	20	Campbell 1983
“ var. pumilus Vasey	20	Campbell 1983
gyrans Ashe var. gyrans	20	Campbell 1983
“ var. stenophyllus (Hackel) Campbell	20	Campbell 1983
hallii Hack	20	Riley & Vogel 1982
hirtifolius Presl	60	Gould 1965
huillensis Rendle	60	Spies & Du Plessis 1987
	c. 50	Spies et al. 1994
hypogynus Hack	80	Norrmann 1985, Campbell & Windisch 1986
ischaemum L. var. ischaemum	60	1986
kilimandscharicus Pilger	40	Gould 1965
lacunosus J.G. Anders	20	Tateoka 1965c
lateralis	18	Spies & Du Plessis 1986
	60	Campbell & Windisch 1986, Norrman 1985, Norrmann et al. 1994
leucostachyus HBK	60 +1B	Norrman 1985
liebmannii Hackel var. pungensis (Ashe) Campbell	20	Pohl & Davidse 1971, Norman 1985
lindmanii Hackel	20	Campbell 1983
lividus Thwaites	60	Campbell & Windisch 1986
	20	Saxena & Gupta 1970, Christopher &

longiberbis Hackel		Samraj 1985
longipes Hack.	20	Campbell 1983
macrothrix Trin.	20	Christopher & Samraj 1985
	20	Norrmann 1985
patris Robyns	60	Campbell & Windisch 1986
polypticus Steud. var. decanensis Bor	20	Dujardin 1979
pseudapricus Stapf	20	Christopher & Samraj 1985
reevesii Gould	40	Dujardin 1979
selloanus (Hackel) Hackel	120	Gould 1965
	20	Norrmann 1985, Honfi et al. 1990
schirensis Hochst ex A. Rich	20	Norrmann & Quarin 1987
ternatus (Spr.) Nees	20	Spies et al. 1994
tectorum Schum. & Thann.	30	Norrmann 1985, Norrmann & Quarin
	18	1987
	20	Okoli 1982
tracyi Nash	30	Okoli 1983, Okoli & Olorode 1983
urbanianus Hitchc.	20	Okoli 1982, 1983, Okoli & Olorode 1983
virgatus Desv.	40	Campbell 1983
	20	Davidse & Pohl 1972
virginicus L. var. glaucus Hackel	60	Pohl & Davidse 1971
“ var. virginicus	20	Campbell & Windisch 1986
	20	Campbell 1983
		Campbell 1983

Arthraxon breviaristatus Hack.	18	Mehra & Kalia 1975, Kalia 1978, Mehra 1982, Sinha et al. 1990
	36	Kalia 1978, Mehra 1982
castratus (Griffith) Nayayan ex Bar	18	Sinha et al. 1990
	36	Probatova & Sokolovskaya 1984a
centrasiaticus (Griseb.) Gamajun.	40	Christopher 1978
depressus Stapf	18	Christopher & Samraj 1985
echinatus (Nees) Hochst.	36	Cristopher & Samraj 1985
hispidus (Thunb.) Makino	36	Mehra & Sharnma 1975, Kalia 1978
inermis Hock.f.	18	Mehra 1965
lanceolatus (Rosb) Hochst.	18	Mehra & Sharma 1972
	30	Mehra & Sharma 1975, Sharma & Sharma 1979
	36	Mehra & Kalia 1975
lanceolatus (Roxb.) Hochst. var. lanceolatus		Spies & Du Plessis 1986
lanciofolius (Trin.)Hochst..	18	Mehra & Sharma 1975, Kalia 1978,
	36	Mehra 1982, Sahnı & Bir 1985, Ahsan et
langsdorffii (Trin.) Hochst. ex Rosch.	36	al. 1994
micans (Nees) Hochst.	36	Mehra & Kalia 1975, Kalia 1978, Mehra
microphyllus (Trin.) Hochst.	16	1982
nudans (Steud.) Hochst.	18	Probatova & Sokolovskay 1981, 1984a
prionodes (Steud) Dandy	16	Mehra & Sharma 1972 & 1975
	20	Kalia 1978, Mehra 1982
	36	Mehra & Kalia 1975, Kalia 1978, Mehra
	36	1982
	36	Kalia 1978, Mehra 1982
quirtinianus (A.Rich) Nash	10	Farugi et al. 1979, Ahsan et al. 1994
	36	Mehra 1965, Farugi et al. 1979, Ahsan et
	38	al. 1994
	36	Tateoka 1965c
	36	Sinde et al. 1975
		Christopher 1978, Merha & Sharma
sikkimensis Bar	16	1975, Dujardin 1979a
	18	Gill et al. 1980
		Gould 1966, Gould & Soderstrom 1970,
		Pohl & Davidse 1971, Mortor 1993
		Merha & Sharma 1975
		Sharma et al. 1978
Bothriochloa		
barbinodis (Lag.) Herter	180	Gould 1966
caucasia (Trin.) C.E. Hubb	40	Sinha et al. 1990
ensiformis (Hook.f.) Henr.	20	Christopher & Samraj 1985
exaristata (Nash) Henrard	60	Allred & Gould 1982
glabra (Roxb.) A. Camus	34-44	Srivastava 1970 & 1980
	40	Tateoka 1965c, Gupta & Srivastava
		1974, Srivastava 1979 & 1980, Sinha et
grahamii (Haines) Bor	40	al. 1990
		Borgaonkar 1964, Gupta & Srivastava
grahamii (Haines) Bor x B. intermedia (R.Br.) A. Camus	40	1974, Farugi et al 1979
	50	Srivastava & Purnima 1990
grahamii (Haines) Bor x D. annulatum (Forsskal)	120	Srivastava & Purnima 1990
Stapf	30	Gould 1968
hybrida (Gould) Gould	40	Srivastava 1980

intermedia (R. Br.) A. Camus var intermedia	50	Celarier et al 1961, Kalia 1978, Srivastava 1979 & 1979a, Farugi et al. 1979, Mehra 1982, Sinha et al. 1990, Srivastava & Purnima 1990
	40	Gupta & Srivastava 1974, Srivastava 1979 & 1980
intermedia (R.Br) A.Camus var. punctata (Roxb) Keng	20	Mehra & Sharma 1975, Sinha et al. 1990
	50, 60	Srivastava & Purnima 1990
intermedia (R. Br.) A. Camus x D. annulatum (Fors.) Stapf	c.60	Spies & Du Plessis 1986a, Strydom et al. 2000
insculpta (A. Rich.) A. Camus	20	Spies & Du Plessis 1987
	40	Spies & Du Plessis 1987a
		Farugi et al. 1979, Koul & Gohil 1987, Ahsan et al. 1994
ischaemum (L.) Keng	30	Hindakova 1974, Petrova in Prokudin et al. 1977, Kliphuis & Wieffering 1979, Skalinska et al. 1968, Love & Love 1982b
	50	
	20	
	40	
	60	Farugi et al. 1979, Ahsan et al. 1994
ischaemum var. songaricus (Rupr.) Celarier & Harlan	60	Gould 1968, Singh 1965
	120	Gill et al. 1980
kuntzeana (Hack.) Henr	n = 20+0-2B	Borgaonkar 1964
.	20	Allred & Gould 1983
laguroides (DC) Herter ssp. laguroides	40	Allred & Gould 1983
laguroides Herter ssp. torreyana (Steud) Allred & Gould		Allred & Gould 1983
longipaniculata (Gould) Allred & Gould		Kalia 1978, Mehra 1982
odorata (Lisboa) A. Camus		Mitra & Datta 1967, Sahni & Bir 1985
pertusa (L) A. Camus		Tateoka 1965c, Mitra & Datta 1967, Gupta 1969, Gould & Soderstrom 1970, Pohl & Davidse 1971, Davidse & Pohl 1972, Gupta 1971, Gupta & Srivastava 1974, Mehra & Sharma 1975, Christopher 1978, Sharma & Sharma 1979, Gill et al. 1980, Roy 1981, Sinha et al. 1990
	36	Sharma et al. 1978
	60	Malik & Tripathi 1970
	40	Farugi et al. 1979
	120	Allred & Gould 1983
	60	Ward & Spellenberg 1986
	120	Gould & Soderstrom 1970
pseudoischaemum (Nees) Henr		
saccharoides ssp. reevesii (Gould) Allred & Gould		
saccharoides var. torreyana (Steud) Gould		
schlumbergi (Fourn. Ex Hemsl.) Henrard		
Chrysopogon aciculatus (Retz.) Trin	20	Gupta 1969, Saxena & Gupta 1970, Gupta 1971, Christopher 1978, Dujardin & Beyne 1975, Mehra & Sharma 1975, Kalia 1978, Mehra 1982
asper (Heyna) Blatt. & McCann	20	Muniyamma et al. 1976, Nagabhushana Rao Sindhe 1980, Sinha et al. 1990
aucheri (Boiss.) Stapf	20	Baquer & Saeed 1969, Farugi et al. 1979, Ahsan et al. 1994
echinulatus (Nees) Watson	20	Mehra 1965, Mehra & Sharma 1972
fulvus (Sprengel) Chiov.	20	& 1975, Farugi et al. 1979
		Mehra & Sharma 1975, Christopher 1978, Sharma & Sharma 1979, Sinha et al.

<i>gryllus</i> (L) Trin	40 n = 10+0-2B 20	1990, Bir & Chauhan 1990a Sinha & Bir 1985 Kalia 1978, Mehra 1982 Mehra & Sharma 1975, Sharma & Sharma 1979, Sinha et al. 1990 Christopher 1978, Petrova in Prokudin et al. 1977
<i>gryllus</i> subsp. <i>Echinulatus</i> (Nees) Cope <i>hackelii</i> (Hook.f.) Fisch	20 20	Ahsan et al. 1994 Sinde et al. 1975, Shanthamma 1976, Christopher & Samraj 1985
<i>lancearius</i> (Hook.f.) Haines	20	Sinha et al 1990
<i>montanus</i> (Forssk.) Stapf	20 80	Sinha et al. 1990 Sinha et al. 1990
<i>orientalis</i> (Desv.) A. Camus <i>serrulatus</i> Trin	20 20	Malik & Tripathi 1970 Christopher & Samraj 1985
<i>verticillatus</i> (Roxb.) Trin <i>zeylanicum</i> (Nees) Thw.	20 20	Saxena & Gupta 1969, Saxena & Gupta 1970, Farugi et al. 1979, Ahsan et al. 1994 Christopher 1978 Christopher 1978
<i>Coelorachis</i> <i>aurita</i> (Steud.) A. Camus <i>balansae</i> (Hack.) A. Camus <i>capensis</i> Stapf <i>khasiana</i> (Hack.) Stapf <i>parodiana</i> Henrard <i>selloana</i> (Hack) A. Camus	18 36 18 18 18 18	Quarin 1979 Quarin 1979 Spies & Du Plessis 1986a Kalia 1978, Mehra 1982 Quarin 1979 Quarin 1979, Honfi et al. 1990
<i>Coix</i> <i>aquatica</i> Roxb.	10  20 40	Koul 1965, Venkateswarlu & Chaganti 1966, Venkateswarlu et al. 1976, Sapre & Barve 1983a, Sapre et al. 1985, Rao & Nirmala 1990, Rao & Nirmala 1994 Venkateswarlu et al. 1976 Venkateswarlu et al. 1976, Christopher & Singh 1986, Christopher et al. 1989
<i>aquatica</i> Roxb. x <i>C. gigantea</i> Koen ex Roxb <i>gigantea</i> Koeng ex Roxb	10-21 12	Sapre & Deshpande 1987 Christopher & Jacob 1991, Jacob & Christopher 1992, Christopher et al. 1995a, Christopher et al. 1996
	n = 9II	Sapre & Barve 1983, Sapre & Barve 1987, Sapre & Naik 1990
	n = 10II 5-8, 9,11,19,21, 22,23 12	Sapre & Barve 1984 Sapre & Barve 1984, Christopher & Jacob 1991, Jacob & Christopher 1992, Christopher et al. 1995a, Christopher et al 1996 & 1997
	18	Sapre & Barve 1983 & 1984, Barve & Sapre 1985, Sapre et al. 1985, Rao & Nirmala 1994
	20	Mehra 1982, Sapre & Barve 1984, Barve & Sapre 1986, Sapre et al. 1985, Sapre & Naik 1990, Rao & Nirmala 1993 & 1994
<i>gygantea</i> x <i>C. aquatica</i>	24	

lacryma – jobi L. var. lacrymajobi	18, 24 10 20	Sapre & Barve 1983 & 1984 Sapre & Deshpande 1987 Venkateswarlu et al. 1976 Koul 1965, Malik & Mary 1970, Saxena & Gupta 1970, Davidse & Pohl 1971 & 1972, Olorode 1975, Venkateswarlu et al. 1976, Sindhe 1977, Christopher 1978, Dujardin 1979a, Mehra 1982, Wang et al. 1984b, Lin et al. 1985, Sarma & Sharma 1986, Sinha et al. 1990, Rao & Nirmala 1993, Christopher et al. 1997 Christopher & Nirmala 1994 Mehra 1982 Christopher & Jacobs 1989 & 1990 Sinde et al. 1975 Christopher et al. 1997
lacryma-jobi L. var. ma-yuen (Rom du Gaill) Stapf	30 10	
lacryma-jobi L. x C. gigantea J.Ko	30 10 16	
Cymbopogon afronardus Stapf bombycinus (R.Br.) Domin caecius (Nees) Stapf	20 20 n = 10II+0- 6B 10II 20	Tateoka 1965c Soenarko 1977 Sudharshan & Jagadishchandra 1984 Christopher & Raj 1986 Gupta 1965 & 1970, Christopher 1978, Lavania et al. 1988, Jagadishchandra 1992
citratu (DC) Stapf	40 40+0-6B 40 60	Quraish & Faruqi 1979, Faruqi et al. 1979, Faruqi & Quraish 1985 Jagadishchandra 1992 Gupta 1965 Soenarko 1977, Christopher & Raj 1986, Lavania et al. 1988, Sinha et al. 1990
coloratus (Nees) Stapf confertiflorus (Steud.) Stapf distans (Nees) Wats	n = 20II, 40 20 20	Christopher & Raj 1986 Gupta 1970 Gupta 1969 & 1970, Mehra & Sharma 1975, Quraish & Faruqi 1979, Faruqi et al. 1979
distans (Nees) Wats. Var. mundensis Gupta excavatus (Hochst.) Stapf ex Burt Davy	40 60 20 20	Gupta 1969 & 1970, Mehra & Sharma 1975 Saxena & Gupta 1969, Saxena & Gupta 1970
exsertus (Hack) A.Camus flexuosus (Nees ex. Steud.) Wats	40 20 n = 20II 8-20 20	Gupta 1970 Spies & Du Plessis 1987a, Hoshino & Davidse 1988, Spies et al. 1994 Spies & Du Plessis 1987a Kalia 1978, Mehra 1982 Christopher & Raj 1986 Lavania 1987
flexuosus (Nees ex. Steud.) Wats var cormbatorensis Gupta	40 20-40	Gupta 1965 & 1970, Christopher 1978, Verma & Sobiti 1982, Verma & Sobti 1985a & 1985b, Christopher & Raj 1986, Banerji et al. 1987, Lavania et al. 1988, Sinha et al. 1990
flexuosus (Nee.) W. Watson var. microstachys (Hook.) Bor	30 n = 10II+0- 2B	Gupta 1970, Christopher 1978, Verma & Sobiti 1982, Christopher & Raj 1986
gidarba (Ham. Ex Hook.) Haines	n = 20II 40	Gupta 1970 Malik & Tripathi 1970



iwarancusa (Sones) Schult jwarancusa Schult	20 n = 10II 20+2B, 20+6B 20+0-8B 20	Christopher & Raj 1986 Christopher & Raj 1986 Christopher & Raj 1986 Mitra & Datta 1967 Christopher & Raj 1986 Verma & Sobti 1985b Sobti et al. 1979 Gupta 1965 & 1970, Baquar & Saeed 1969, Kalia 1978, Mehra & Sharma 1975, Quraish & Faruqi 1979, Faruqi et al. 1979, Verma & Sobiti 1982, Mehra 1982, Verma & Sobti 1985a & b, Christopher & Raj 1986, Sreenath & Chandra 1987, Lavanina et al. 1988 Baquar & Saeed 1969, Christopher & Raj 1986, Sinha et al. 1990 Gupta 1970 Kalia 1978, Mehra 1982 Sobti et al. 1979 Gupta 1965 & 1970, Verma & Sobiti 1982, 1985a Gupta 1970 Gupta 1970c Gupta 1970c Ahsan et al. 1994 Spies et al. 1994 Du Plessis & Spies 1988 Gupta 1965, Mitra & Datta 1967, Gupta 1970, Mehra & Sharma 1975, Bir & Chauhan 1990a Gupta 1965 & 1970, Mehra & Sharma 1975, Ahsan et al. 1994 Gupta 1965 Verma & Sobiti 1982 & 1985a & b, Lavanina et al. 1988 Christopher & Raj 1986 Sobti et al. 1979, Verma & Sobiti 1985b Christopher & Raj 1986 Verma & Sobiti 1985b Gupta 1970, Verma & Sobiti 1982 & 1985a, Christopher & Raj 1986, Lavanina et al. 1988 Mehra & Sharma 1972 & 1975 Gupta 1970d Gupta 1965 & 1970, Christopher & Raj 1986, Sinha et al. 1990 Gupta 1965 & 1970, Lavanina et al. 1988 Gupta 1965, Kalia 1978, Verma & Sobiti 1982, Mehra 1982, Verma & Sobiti 1985b, Christopher & Raj 1986, Lavanina et al. 1988 Christopher & Raj 1986 Gupta 1965, Mitra & Datta 1967, Mehra & Sharma 1975, Christopher & Raj 1986 Gupta 1969 & 1970, Quraish & Faruqi 1979, Faruqi et al. 1979
jwarancusa (Jones) Schult var. assamensis Gupta khasianus (Hack.) Stapf ex Bor.	40 20 20 60+2B 60	
kravancorensis Bor ladakhensis Gupta ladakensis Gupta lanceifolium L. Liou marginatus (Steudel) Stapf ex Burtt Davy	20 20 20 40 20	
martinii Wats	40 20	
martinii (Roxb.) Wats var. motia	40 20+B 20	
martinii (Roxb.) var. sofia Gupta	n = 10II+0- 2B 20+0-2B 20+6B n = 10II, 20II	
microtheca (Hook.f) A. Camus motia Gupta nardus (Linn.) Rendle	40 20 20 + 2B 20	
nardus var. confertiflorus	40 20	
olivieri (Boiss.) Bor	40 60	
parkeri Stapf	20	
parkeri var. jammuensis Gupta parkeri Stapf x C. caesius (Nees) Stapf	n = 10II 20 30 40	

pendulus Wats	40 60 20 30 40 60	Chrisopher & Raj 1986 Gupta 1970, Sahni & Bir 1985 Gupta 1970 Gupta 1970, Quraish & Faruqi 1979, Sharma & Kumar 1980, Christopher & Raj 1986
plurinodus (Stapf) Stapf ex Burt Davy	60	Gupta 1970
polyneuros (Steud) Stapf	20 40 80	Faruqi et al. 1979 Kalia 1978, Mehra 1982 Gupta 1969
procerus (R.Br.) Domin	n = 10II+0- 1B	Lavania et al. 1988 Gupta 1965 & 1970, Verma & Sobiti
prolixus (Stapf) Phill	n = 20II	1982, Verma & Sobiti 1985a
ramnagarensis Gupta	20	Spies & Du Plessis 1987, Strydom et al.
schoenanthus (L.) Sprengel	20 40 20	2000 Du Plessis & Spies 1988 Spies et al. 1994
stracheyi (Hook.f.) Raiz. & Jain	n = 20II+2B 20+1B	Christopher & Raj 1986 Christopher & Raj 1986
travencorensis Bor	20 40 20	Gupta 1965 & 1970 Soenarko 1977 Spies & Du Plessis 1987, 1988
validus (Stapf) Stapf ex Burt Davy	40 20	Gupta 1970e Christopher & Raj 1986 Christopher & Raj 1986
winterianus Jowitt.	40 20 30 60 20	Pandy & Pal 1980a, Faruqi et al. 1987 Gill et al. 1980 Mehra & Sharma 1972 Gupta 1969 & 1970 Gupta 1965 & 1970, Christopher & Raj 1986, Lavania et al. 1988 Christopher 1978 Spies & Du Plessis 1987a, Hoshino & Davidse 1988, Spies et al. 1994 Spies & Du Plessis 1986a Spies et al. 1994 Gupta 1965, Verma 1985, Lavania et al. 1988
Dichanthium annulatum (Forsskal) Stapf	20 40 32 50 60	Roy et al. 1965, Singh 1966, Gupta 1971, Sarkar et al. 1975a Sinha et al. 1980, Borgaonkar & Singh 1962 Singh & Godword 1960, Borgaonkar & Singh 1962, Gupta 1969, Reddy & D'Cruz 1969 & 1969b, Baquar & Saeed 1969, Malik & Tripathi 1970, Gupta & Srivastava 1974, Kalia 1978, Faruqi et al. 1979, Sinha et al. 1980, Ray 1981, Mehra 1982, Bir & Singh 1983, Spies & Du Plessis 1986a, Bir et al. 1987, Koul & Gohil 1988, Bir & Chauhan 1990, Ahsan et al. 1994 Bir & Chauhan 1990 Gupta & Srivastava 1974, Srivastava & Purnima 1990

annulatum (Forssk.) Stapf var. annulatum annulatum (Forssk.) Stapf var. fecundum annulatum (Forssk.) Stapf var. papillosum annulatum (Forssk.) Stapf x B. intermedia (R.Br.) A Camus annulatum (Forssk.) Stapf x B. grahamii (Haines) Bor aristatum (Poir.) C.E. Hubb.	20, 40 40 60 50 50 20 40  80 40 20 40 40 20 60	Faruqi et al. 1979, Spies & Du Plessi 1986a, Bir & Chauhan 1990 Singh & M 1965 Singh & M 1965 Singh & M 1965 Srivastava & Purnima 1990 Srivastava & Purnima 1990 Reddy & D'Cruz 1969 Knox & Heslop-Harrison 1966, D'Cruz & Reddy 1969, Sinha et al. 1990, Strydom et al. 2000 Gould 1968 Rao & Mwasumbi 1981a Reddy & D'Cruz 1969 & 1969b D'Cruz & Reddy 1969, Sinha et al. 1990 Ahsan et al. 1994 Christopher & Samraj 1985 Sing 1966 Borgoankar & Sharma 1962
bladhi (Reitz.) Clayton caricosum (L.) A. Camus  foveolatum (Delile) Roberly pallidum (Hook.f.) Stapf panchganiense Blatt. & McCann papillosum	20 20 10 20 30 20 20 10	Gupta 1969 Dujardin 1978 Dujardin 1978 Normann et al. 1994, Spies et al. 1994 Strydom et al. 2000 Ahsan et al. 1994 Gould 1965, Gould & Soderstrom 1970 Strydom et al. 2000
Elionurus adustus (Trin.) Ekm. brazzae Franch hesii K. Schum muticus (Spreng.) Kuntze  royleanus Nees ex A. Rich tripsacoides Humb. & Bonpl. var. ciliaris (HBK) Hack vilosa (Thunb.) Nees	20 20 10 20 30 20 20 10	Gupta 1969 Dujardin 1978 Dujardin 1978 Normann et al. 1994, Spies et al. 1994 Strydom et al. 2000 Ahsan et al. 1994 Gould 1965, Gould & Soderstrom 1970 Strydom et al. 2000
Eriochrysis brachypogon (Stapf) Stapf cayennensis Beauv.	20 20	Dujardin 1978 & 1979 Pohl & Davidse 1971
Eulalia fastigiata (Nees) Haines hirtifolia (Hack.) A. Camus  mollis (Griseb.) O. Kuntze  phaeothrix (Hack.) Kuntze quadri-versis (Hackel) Kuntze speciosa (Debeaux) Kuntze trispicata (Schult.) Henr.  villosa (Thunb.) Nees wightii (Hook.f.) Bor.	18 20 40 18  20 n = 7+2B 40 18 20 20 20	Prince 1965, Parkash 1979, Mehra 1982 Mehra & Sharma 1972 Sindhe 1980 Mehra & Sharma 1972, Parkash 1979, Mehra 1982 Christopher & Samraj 1985 Parkash 1979, Mehra 1982 Parkash 1979, Mehra 1982 Parkash 1979, Gill et al. 1980, Mehra 1982 Faruqi et al. 1979 Hoshino & Davise 1988 Christopher & Samraj 1985
Hackelchloa granularis (Sw.) Kuntze granularis (L.) Kuntze	14 14	Gupta 1969 & 1971 Christopher 1978, Dujardin 1978, Kalia 1978, Pohl & Davidse 1971, Reeder 1971, Mehra 1982, Christopher 1986, Bir

porifera (Hack.) Rhind.	30	& Sahni 1986, Ahsan et al. 1994 Kalia 1978, Mehra 1982
Hemarthria altissima (Poir.) Stapf & C.E. Hubb.	20 18 36	Du Plessis & Spies 1988 Davidse & Pohl 1972, Quesenberry et al. 1982
compressa (L.f.) R. Br.	n = 9+0-2B n = 18+0-2B 18	Davidse & Pohl 1972, Quarin 1977, Quesenberry et al. 1982 Kalia 1978, Mehra 1982 Kalia 1978, Mehra 1982 Mehra & Sharma 1975e, Faruqi et al. 1979, Sharma & Sharma 1979
longiflor (Hook. f.) A. Camus	36	Gupta & Yashavir 1971, Mehra & Sharma 1975, Sinha et al. 1990
protensa Steud	54 54	Quesenberry et al. 1982
uncinata R. Br.	18	Kalia 1978, Mehra 1982
Uncinata R. Br. Var. spathacea (Damin) Vickery	20 36 36	Mehra & Kalia 1975, Kalie 1978, Mehra 1982 Sinha et al. 1990 Quesenberry et al. 1982 Quesenberry et al. 1982
Heteropogon contortus (L.) Beauv.	39 40	Tothill & Hacker 1976 Gupta 1969 & 1971, Gupta & Srivastava 1974, Mehra & Sharma 1975, Christopher 1978, Kalia 1978, Mehra 1982
contortus (L.) Beauv. Ex Roem & Schult	50 60	Gupta 1969 & 1971, Gupta & Srivastava 1974, Tothill & Hacker 1976 Gould 1968, Gupta 1969 & 1971, Gupta & Srivastava 1974, Tothill & Hacker 1976, Borgen 1980, Bir et al. 1987
contortus (L.) Roem & Schult.	69, 70, 80, 90 20 36 40	Tothill & Hacker 1976 Christopher 1978, Ahsan et al. 1994 Bir & Chauhan 1990a Tateoka 1965c, Faruqi et al. 1979, Dujardin 1979, Gill et al. 1980, Rao & Mwasumbi 1981, Sinha et al. 1990, Ahsan et al. 1994
melanocarpus (Elliott) Elliott ex Benth triticeus (R. Br.) Stapf ex Craib	60 20 40 46 56 58 60 22	Davidse & Pohl 1972, Davidse & Pohl 1972, Srivastava & Purnima 1990, Sinha et al. 1990 Spies & Du Plessis 1986a Spies & Du Plessis 1986, 1987a Spies & Du Plessis 1986 Spies & Du Plessis 1987a Bir & Sahni 1983, 1985 Srivastava & Purnima 1990 Sinha et al. 1990
Hypparrhenia anamesa Clayton	40 60	Spies & Du Plessis 1988, Du Plessis & Spies 1988, Spies et al. 1994 Spies & Du Plessis 1988, Spies et al.

conspersum Schrad. Ex Schult.	40	1994
cymbaria (L.) Stapf	30	Burson 1978
diplandra (Hack.) Stapf	40	Tateoka 1965b
dissoluta (Nees) C.E. Hubb.	40	Dujardin 1979
dregeana (Nees) Stapf	20	Davidse & Pohl 1972
familiaris (Steud.) Stapf	40	Spies & Du Plessis 1987
filipendula (Hochst) Stapf var. filipendula	40	Dujardin 1978
filipendula var. pilosa (Hochst. ex Hack) Stapf	20	Dujardin 1979, Spies et al. 1994
	40	Spies & Du Plessis 1987a
griffithii Bor.	36	Spies & Du Plessis 1987a, Spies et al.
hirta (L.) Stapf	c.20	1994
	30	Kalia 1978, Mehra 1982
	36	Romerozarco et al. 1988
	40	Talavera 1978, Devesa et al. 1990
		Faruqi & Quraish 1978, Humphries 1978
		Fernandez & Queiros 1969, Gould 1970,
	40+1B, c.48	Faruqi et al. 1979, Miraval 1983,
	45	Hoshino & Davidse 1988, Spies et al.
	54, 56	1994
	60	Gould 1970
		Fernandez & Queiros 1969, Baldini 1991
intermedium Manro ex Morong	20	Faruqi & Quraish 1978
involucrata	20,21,40	Gould 1970, Spies & Du Plessis 1986a,
jurgensii Hackel	20	1988, Devesa et al. 1990a
pilgeriana C.E. Hubb.	20, 80	Burson 1978
pubescens (Vis.) Chiov.	30	Olorode & Morakinyo 1980
quarrei Robyns	60	Burson 1978
rufa (Nees) Stapf	20	Spies et al. 1994
	30	Miraval 1983
	36	Spies & Du Plessis 1986a
	40	Tateoka 1965c, Sinha et al. 1990
smithiana (Hooker f.) Stapf	40	Rao & Mwasumbi 1981a
subplumosa	40	Tateoka 1965b, Nassar 1977
tamba (Steud.) Stapf	20	Dujardin 1979
	40	Morton 1993
variabilis Stapf	20	Olorode & Morakinyo 1980
	40	Spies & Du Plessis 1986a
		Spies & Du Plessis 1987a
		Tateoka 1965c
		Spies et al. 1994
Hyperthelia		
dissoluta (Nees ex Steud.) W.D. Clayton	20	Spies et al. 1994
	40	Dujardin 1978, Rao & Mwasumbi 1981a
Imperata		
cylindrica (L.) P. Beauve.	20	Gupta 1971, Christopher 1978, Faruqi et
		al. 1979, Rao & Mwasumbi 1981, Bir &
		Singh 1983, Sahni & Bir 1985, Bir et al.
		1987, Sinha et al. 1990
	40	Faruqi et al. 1987, Sinha et al. 1990
	52	Faruqi et al. 1979
	60	Fernandez & Queiros 1969
cylindrica (L.) Beauv. var. africana (Anderss.)	60	Tateoka 1965b, Harvey 1966a, Dujardin
Hubb.	20	1979
cylindrica (L.) Beauv. var. cylindrica	20	Kalia 1978, Mehra 1982

cylindrica (L.) Beauv. var. koenigii (Retz.) Dur. & Schinz.	20	Tateoka 1967
cylindrica (L.) Beauv. var. major (Nees) C.E. Hubb.	20	Kalia 1978, Mehra 1982
conferta (Presl) Ohwi	20	Price & Daniels 1968, Reeder & Soderstrom 1968
contracta (HBK) Hitchc.	20	Pohl & Davidse 1971
tenius Hack.		Norrmann et al. 1994
Ischaemum		
afrum (J.F. Gmel.) Dandy	20	Spies & Du Plessis 1987a, Hoshino & Davidse 1988, Spies et al. 1991
aristatum L.	40	Oke 1971
	60	Oke 1971
ciliare Retz.	18	Pohl & Davidse 1971
	20	Oke 1971
commutatum Hack.	20	Christopher & Samraj 1985
diplopogon Hook.	20	Oke 1971
duthiei Stapf	20	Gill et al. 1980
fasciculatum Brongn.	20	Spies & Du Plessis 1987a
flumineum Bor.	40	Oke 1971
goebelii Hackel	38	Parkash 1979, Mehra 1982
hirtum Hook.	20	Parkash 1979, Mehra 1982
hubbardii Bor.	40	Parkash 1979, Mehra 1982
impressum Hack.	20	Oke 1971
indicum (Houtt.) Merr.	20	Davidse & Pohl 1978
	40	Reeder & Soderstrom 1968
indicum (Houtt.) Merr. Var. indicum	18	Christopher 1978
	40	Sindhe & Harayon 1976
indicum (Houtt.) Merrill var. wallichii (Hack.) Bor.	54	Christopher 1978, Parkash 1979, Mehra 1982
kingii Hook.	20	
latifolium (Spreng.) Kunth	36	Oke 1971
mangaluricum (Hack.) Stapf	20	Pohl & Davidse 1971
		Sindhe & Narayon 1976, Christopher & Samraj 1985
molle Hook.	20	
	30	Christopher 1978
muticum L.	34	Oke 1971
nilagiricum Hack.	20	Christopher 1978
	40	Christopher 1978
pilosum Hack.	60	Sindhe & Narayon 1976
Rangacharianum Fish.	40	Oke 1971
ritchiei Stapf	20	Christopher 1978
rugosum Salisb.	18	Oke 1971
	20	Pohl & Davidse 1971, Parkash 1979, Davidse & Pohl 1978
	40	Singh & Godward 1960, Oke 1971, Bir & Sahni 1986
	44	Oke 1971, Bir & Sahni 1983, Bir & Sahni 1985
	60	Sindhe & Narayon 1976
santapau Bor.	20	
	40	Oke 1971
semisagittatum Roxb.	20	Oke 1971
thomsonianum Stapf	20	Oke 1971
	40	Oke 1971
travancorense Stapf	20	Christopher 1978
zeylanicum Bor.	20	Oke 1971, Sindhe & Narayon 1976
		Christopher 1978

		Christopher 1978
<p><i>Microstegium</i> <i>ciliatum</i> (Trin.) A. Camus</p> <p><i>japonicum</i> (Miq.) Koidz <i>nudum</i> (Trin.) A. Camus</p> <p><i>petiolare</i> (Trin.) Bov.</p> <p><i>vagans</i> (Nees) A. Camus</p> <p><i>vimineum</i> (Trin.) A. Camus <i>vimineum</i> (Trin.) var. <i>polystachyum</i> (Fr. &amp; Sav.) Chwi <i>vimineum</i> (Trin.) var. <i>willdenowianum</i> (Nees) Bor</p>	<p>20 40 n = 40+0-2B 60 c60</p> <p>20 20 42</p> <p>20 n = 40+1-2B 20 60 n = 30+2-4B 40 40 40</p>	<p>Kalia 1978, Mehra 1982 Mehra &amp; Sharma 1975 Mehra 1982 Mehra &amp; Kalia 1976, Kalia 1978, Mehra 1982</p> <p>Reeder &amp; Soderstrom 1968 Tateoka 1967 Mehra &amp; Sharma 1975, Kalia 1978, Mehra 1982</p> <p>Mehra &amp; Sharma 1975, Sharma &amp; Kumar 1980 Kalia 1978, Mehra 1982 Kalia 1978, Mehra 1982 Mehra &amp; Sharma 1975 &amp; 1975e, 1977 Mehra &amp; Kalia 1976, Kalia 1978, Mehra 1982 Mehra &amp; Kalia 1975 Mehra &amp; Kalia 1976 Tateoka 1967 Kalia 1978, Mehra 1978</p>
<p><i>Miscanthus</i> <i>capensis</i> (Stapf) Pilg. <i>floridulus</i> (Labill.) Warb. <i>nepalensis</i> (Trin.) Hack. <i>nudipes</i> (Griseb.) Hack.</p> <p><i>purpurascens</i> Anders. <i>sacchariflorus</i> (Maxim.) Hackel</p> <p><i>sacchariflorus</i> var. <i>ganchaiensis</i> <i>sacchariflorus</i> var. <i>sacchhaiensis</i> <i>sinensis</i> Anders.</p>	<p>30 38 40 40 n = 20+4B n = 20+2-4B 38, 40 38 40</p> <p>38 76 40</p>	<p>Strydom et al. 2000 Price 1965, Price &amp; Daniels 1968 Mehra &amp; Sharma 1975, Kalia 1978, Mehra 1982 Mehra &amp; Sharma 1972 &amp; 1975 Mehra 1982 Kalia 1978 Sokolovskaya 1966 Sokolovskaya 1966, Zhang &amp; Xie 1989 Sokolovskaya 1966, Probatova &amp; Sokolovskaya 1983 Xie et al. 1989 Xie et al. 1989 Probatova et al. 1989</p>
<p><i>Monocymbium</i> <i>ceresiiforme</i> (Nees) Stapf</p>	<p>20</p>	<p>Dujardin 1978, 1979, Hoshino &amp; Davidse 1988</p>
<p><i>Phacelurus</i> <i>speciosus</i> (Steud.) C.E. Hubb.</p> <p><i>typhoides</i></p>	<p>20 80 40</p>	<p>Gould &amp; Soderstrom 1970, Mehra &amp; Sharma 1975 &amp; 1977, Sharma &amp; Sharma 1979, Ahsan et al. 1994 Mehra 1965 Ramulu &amp; Rangasamy 1971</p>
<p><i>Rottboelia</i> <i>compressa</i> L. <i>digitata</i> Sibth. &amp; Sm.</p>	<p>18 38 20</p>	<p>Bir et al. 1987 Bir &amp; Sahni 1983 Kozuharov &amp; Nicolova 1975</p>

exaltata L.f.	20	Mitra & Datta 1967, Pohl & Davids 1971, Trouin 1972, Sindhe et al. 1975, Dujardin 1978a, Christopher 1984, 1984a, 1986, Christopher et al. 1989a, Mini 1994
	40	Chatterji 1975, Sindhe et al. 1975, Christopher 1978, Kalia 1978, Mehra 1982, Sahni & Bir 1985, Christopher 1984, 1986
formosa R. Br.	36	Sindhe et al. 1990
goalparenensis Bor	18	Singh & Godward 1960
latifolia Wall.	20	Kalia 1978
	32	Mallia et al. 1977a
Saccharum		
alopecuroideum (L.) Nutt	30	Burner & Webster 1994
arundinaceum Retz.	40	Xiao et al. 1992
	60	Sahni & Bir 1985, Xiao et al. 1992
baldwinii Spreng.	30	Burner & Webster 1994
bengalense Retz.	20	Kalia & Mehra 1989, Bir & Sahni 1983, 1985 & 1987, Sahni 1988, Bir et al. 1990b
	20+1-5B	Kalia & Mehra 1989
	22	Bir & Sahni 1983, 1985 & 1987, Sahni 1988, Bir et al. 1990b
	40	Kalia & Mehra 1989, Bir & Sahni 1983, 1985 & 1987, Bir et al. 1987, Sahni 1988, Bir et al. 1990b
	60	Kalia & Mehra 1989, Bir & Sahni 1983, 1985 & 1987, Sahni 1988, Bir et al. 1990b
brevibarbe var. breviarbe	60	1990b
brevibarbe var. contortum (Baldwin ex Elliot) R.D. Webster	60	Burner & Webster 1994
coarctatum (Fernald) R.D. Webster	60, 80	Burner & Webster 1994
edule Hassk.	70	Burner & Webster 1994
	30, 60, 90	Roach 1972
giganteum (Walter) Pers	20	Price & Daniels 1968, Roach 1972
griffithii Munro ex Boiss	40, 60	Burner & Webster 1994
	40	Ahsan et al. 1994
officinarum L.	80	Faruqi et al. 1979
		Nair 1972, Iwo & Agboire 1992
		Jagatheson & Sisodia 1967, Sisodia 1967, Nair 1972, Nair 1975, Sreenivason & Sreenivason 1984, Iwo & Agboire 1992, Chen 1993, Zhuang et al. 1996
	108	Price & Daniels 1968, Iwo & Agboire 1992
	120	Price & Daniels 1968, Iwo & Agboire 1992
	70-81, 99	Price & Daniels 1968, Iwo & Agboire 1992
	30, 48, 50,	Nair 1972, Iwo & Agboire 1992
	54, 60, 64,	Price & Daniels 1968
	68, 72, 88, 96	
	n = 84, 86, 88	
	c83	
officinarum L. clone Black Fiji	84	Iwo & Agboire 1992
officinarum L. clone 21 N.G. 2	107-114	Fernandez et al. 1978
officinarum L. clone N.C. 30	c92-93	Li & Price 1965
officinarum L. clone N.C. 76	100	Li & Price 1965
officinarum L. clone N.C. 92	84, 96, 102,	Li & Price 1965
officinale	104, 108, 112	Li & Price 1965



procerum Roxb.	40	Li & Price 1965
ravennae	60	
ravennae subsp. Parviflorum (Pilger) Maire	c60	Hinojo et al. 1985
robustum Brandes & Jeswiet ex Grassl.	60	Kalia 1978, Mehra 1982
	80	Bir & Sahni 1987
	88, 92, 100,	Faruqi et al. 1987
	c85	Jagathesan & Rathambal 1969,
robustum Brandes & Jeswiet ex Grassl. Clone 28	n = 44II+7I	Sreenivason & Sreenivason 1984
N.G. 219	c64	Jagathesan & Rathambal 1969
robustum Brandes & Jeswiet ex Grassl. Clone 28	c64	Sreenivason & Sreenivason 1984
N.G. 129A	110	Sreenivason & Sreenivason 1984
sinense Roxb.	81, 82, 83	Li & Price 1965
sinense 'Mungro'	82, 83	Li & Price 1965
sinenese 'Dhaultu'	91, 91+f, 92	Zhuang et al. 1996
sinense 'Saretha'	105-119, 124	Price 1968
sinense 'Nargori'	106-120	Price 1968
sinense 'Pansahi'	104-121	Price 1968
sinense unclassified	36, 38	Price 1968
spontaneum L.	40	Price 1968
	48	Sreenivasan & Sreenivasan 1984
	52	Sreenivasan 1975, Bir & Sahni 1983,
	54	1985, 1986, 1987
	56	Sreenivasan 1975, Bir & Sahni 1986,
	58	1987, Singh et al. 1991
	60	Faruqi et al. 1979, Sreenivasan &
	62	Sreenivasan 1984
	64	Bir & Sahni 1983, 1985, Sreenivasan &
	72	Sreenivasan 1984, Sinha et al. 1990,
	80	1990a, Singh et al. 1991, Ahsan et al.
	90	1994
	116	Moriya 1965, Sreenivasan & Sreenivasan
	120	1984, Bir & Sahni 1985
	112, 124, 126	Bir & Sahni 1983, 1985, Bir et al. 1987
spontaneum L. clone Burma	n = 57II+II	Faruqi et al. 1979, Bir & Sahni 1986,
spontaneum Glagah	c96	1987, Sinha et al. 1990
spontaneum Glagah cvs	112	Singh et al. 1989
	40, 52, 62-66,	Jagathesan & Sreen 1966, Christopher
	124, 128	1978, Bir & Sahni 1986, 1987
spontaneum Coimbatore	n = 27, 28,	Sreenivasan 1975, Bir & Sahni 1986,
	30+1, 31+1,	1987, Zhuang et al. 1996
	32, 40, 56,	Moriya 1965, Sreenivasan 1975,
	62, 64	Sreenivasan & Sreenivasan 1984, Bir &
trinii (Hack.) Renvoize	60	Sahni 1986, 1987
		Sinha et al. 1990
		Sreenivasan & Sreenivasan 1984
		Jagathesan & Sreen 1966
		Sreenivasan 1975
		Sreenivasan & Sreenivasan 1984
		Li & Price 1965
		Nair 1972
		Nair 1972
		Nair 1975

		Norrman et al. 1994
Sehima nervosum (Rottl.) Stapf	40	Tateoka 1965c, Christopher 1978, Sinha et al 1990
nervosum var. robusta Stapf	40	Oke 1971
notatum (Hack.) A. Camus	30	Mehra 1965
sulcatum A. Camus	40	Oke 1971
Schizachyrium brevifolium (Sw.) Nees ex Buse	20	Dujardin & Beyne 1975, Kalia 1978, Dujardin 1979, Mehra 1982
	40	Christopher & Samraj 1985
cirratum (Hack.) Wook. & Stadl	20	Hatch 1980
condensatum (HBK) Nees	20	Pohl & Davidse 1971
delavayi (Hackel) Bor	20	Kalia 1978, Mehra 1982
exile Stapf	20	Christopher 1978
gracile (Spreng.) Nash	40	Davidse & Pohl 1972
hirtiflorum Nees	40	Davidse & Pohl 1972
	60	Pohl & Davidse 1971
	70, 100	Hatch 1980
platyphyllum (Franch.) Stapf	n = 12+1B	Dujardin 1979
pulchellum (Don ex Benth.) Stapf	20	Dujardin 1978
sangiuneum (Retz.) Alts.	40	Spies & Du Plessis 1987a
	50	Dujardin 1978, Norrman et al. 1994
	60, 70	Norrman et al. 1994
	80	Kalia 1978, Mehra 1982
scoparium (Michx.) Nash var. frequens (F.T. Hubb.) Gould	40	Gould 1968
semiberbe Nees	40, 80	Hatch 1980
thollonii (Franch.) Stapf	20	Dujardin 1979
Sorghastrum agrostoides (Speq.) Hitchc.	20	Pereira Flores 1983, 1986, Pereira Flores & Montenegro Valls 1987
albescens (Hackel) Flores	20	Pereira Flores 1983, 1986, Pereira Flores & Montenegro Valls 1987
bipennatum (Hack) Pilg.	20	Olorode 1975
	40	Dujardin 1979a
elliottii (Mohr.) Nash	20	Hatch 1980
incompletum (Presl.) Nash	20	Pohl & Davidse 1971
minarum (Nees) Hitchc.	20	Pereira Flores 1983, 1986, Pereira Flores & Montenegro Valls 1987, Norrman et al. 1994
nutans L.	40	Love & Love 1981a, Riley & Vogel 1982, Vahidy et al. 1987
rigidifolium (Stapf) Chippindall	40	Tateoka 1965c
scaberrimum (Nees) Herter	20	Pereira Flores 1983, 1986, Pereira Flores & Montenegro Valls 1987
setosum (Griseb) Hitchc.	20	Davidse & Pohl 1972, Pereira Flores 1983, 1986, Pereira Flores & Montenegro Valls 1987, Norrman et al. 1994
stipoides (H.B.K.) Nash	20	Pereira Flores 1983, 1986, Pereira Flores & Montenegro Valls 1987

Sorghum		
aethiopicum	20	Magoon et al. 1967c
almum Parodi	40	Reddi 1970
amplum Lazarides	10	Lazarides et al. 1991
angustum S.T. Blak.	10	Lazarides et al. 1991
arundinaceum (Desv.) Stapf	20	Magoon et al. 1967b, Magoon & Tayyab 1968, Gadella 1977c, Dujardin 1979
australiense Garbver & Snyder	20	Spies et al. 1991
bicolor (L.) Moench.	10	Morakinyo & Olorode 1988
	20	Vachova 1978, Gu et al. 1984, Feng et al. 1985, Morokinyo & Olorode 1988, Laurie & Bennett 1989, Wu 1990, Ahsan et al. 1994, Ingaki & Mujeeb-Kazi 1995, He et al. 1997
	40	Laurie & Bennett 1989
bicolor ssp. arundinaceum (Desv.) De Wet & Harlan	40	Spies & Du Plessis 1987
brachypodium Lazarides	10	Lazarides et al. 1991
bulbosum Lazarides	20	Lazarides et al. 1991
caudatum	20	Saini et al. 1982
cernuum	20	Magoon et al. 1967b
cernuum var. orbiculatum	20	Magoon et al. 1967c
conspicuum	20	Magoon & Tayyab 1968
coriaceum	20	Magoon & Tayyab 1968
dura	20	Saini et al. 1982
dura var. mediocre	10	Magoon et al. 1967c
ecarinatum Lazarides	10	Lazarides et al. 1991
extans Lazarides	30, 40	Lazarides et al. 1991
grande Lazarides	18	Lazarides et al. 1991
halepense (L.) Pers.	20	Bir et al. 1980a, Bir et al. 1987
	26	Christopher 1978, Faruqi et al. 1979, Bir & Sahni 1983
	40	Raman et al. 1966
		Gould 1968, Fernandez & Queiros 1969, Reddi 1970, Pohl & Davidse 1971, Gupta & Yashavir 1971, Gupta 1971, Ferakova 1976, Muniyamma et al. 1976, Kalia 1978, Faruqi et al. 1979, Mehra 1982, Bir & Singh 1983, Probatova & Sakolovskaya 1983, Warwick & Black 1983, Gu et al. 1984, Mulligan 1984, Gohil & Koul 1986, Bir et al. 1987, Faruqi et al. 1987, Ahsan et al. 1994, Warwick et al. 1997
halepense (L.) Pers. var. halepense	41-43	
halepense var. muticum (Hackel) Hayek	60	Fernandez & Queiros 1969
interjectum Lazarides	40	Bir & Chauhon 1990
intrans F. Mueller ex Benth.	10	Devesa et al. 1991
intrans x S. plumosum (R.Br.) P. Beauv.	20	Devesa et al. 1991
laxiflorum F.M. Bailey	40	Lazarides et al. 1991
macrocheata Snowden	40	Gu et al. 1984, Lazarides et al. 1991
macrospermum Garber	40	Lazarides et al. 1991
margaritiferrum	20	Lazarides et al. 1991
matarakense Garber & Snyder	10	Reddi 1970
melaleucum	20	Wu 1990
membranaceum	20	Magoon et al. 1967c
	40	Lazarides et al. 1991, Spies et al. 1991

miliaceum (Roxb.) Snowden	20	Magoon & Tayyab 1968
miliforme	20	Magoon et al. 1967b, Magoon & Tayyab 1968
nervosum	20	Reddi 1970
nitens	40	Magoon et al. 1967b, Magoon & Tayyab 1968
	10	Magoon et al. 1967b
nitidum (Vahl) Pers.	10+1B	Sadasiwaiah & Mitra 1966, Magoon et al. 1967b, Magoon & Tayyab 1968a & 1968
	20	Sadasiwaiah & Mitra 1966
	40	Magoon & Tayyab 1968a, Wu 1982, Rao & Rao 1990
panicoides Stapf in Prain	10	Wu & Pi 1975, Wu 1978
plumosum (R.Br.) P. Beauv. var. plumosum	20	Kalia 1978, Mehra 1982
propinquum	10	Gu et al. 1984
purpureo-sericeum Aschers & Schweinf.	10	Reddi 1970
pupureo-sericeum subsp. aimidiatum (Stapf) Garber	20	Lazarides et al. 1991
roxburghii	20	Magoon et al. 1967 a & b
roxburghii var. nanum	20	Wu & Pi 1968, Wu 1971, Gu et al. 1984, Rao et al. 1989
saccharatum (L.) Moench.	20	Roa et al. 1989
splendidum	10	Magoon & Tayyab 1968, Saini et al. 1982
stipoideum (Ewart & White) C.A. Gardner & C.E. Hubb	20	Magoon et al. 1967c
	20	Vachova 1978, Magoon & Tayyab 1968
subglarens	20	Magoon & Tayyab 1968
sudanense	10, 20	Gu et al. 1984, Lazarides et al. 1991, Spies et al. 1991
technicum	10	Magoon et al. 1967b
timorensis (Kunth) Buse ex De Vries	20	Vachova 1978, Magoon et al. 1967b, Magoon & Tayyab 1968, Saini et al. 1982
versicolor	20	Magoon et al. 1967b
verticilliflorum (Steud.) Stapf	20	Lazarides et al. 1991
virgatum	20	Wu 1982, Gu et al. 1984, Laurie & Bennett 1989, Sun et al. 1994
vulgare	20	Magoon et al. 1967b, Magoon & Tayyab 1968, Renard et al. 1983
		Magoon et al. 1967b, Magoon & Tayyab 1968, Saini et al. 1982
		Magoon et al. 1967a & b, Saini et al. 1982
Thelepogon		
elegans Roth ex R. & S.	10	Sisodia 1970, Sinde et al. 1975
Themeda		
anathera (Nees) Hackel	20	Mehra & Sharma 1975, Birari 1980a & 1981
	60	Mehra & Sharma 1975, Kalia 1978, Sharma & Sharma 1979, Mehra 1982
anathera var. glabrescens Hackel	n=10II+1f, 13, 20	Faruqi et al. 1979, Faruqi & Quresh 1987
anathera var. hirsuta Anders.	n=10II+1f, 10, 11, 30	Faruqi et al. 1979, Faruqi & Quresh 1987
anathera var. submarginata	n=10II+1f,	Faruqi et al. 1979, Faruqi & Quresh 1987

arundinacea (Roxb.) Ridley	10, 11, 30 20 20+4B 22 40 60	Faruqi et al. 1979, Faruqi & Quresh 1987 Sinha et al. 1990 Sinha et al. 1990 Kalia 1978, Mehra 1982 Mehra & Sharma 1975, Kalia 1978, Mehra 1982
caudata (Nees) A. Camus	40	Singh & Godward 1960, Kalia 1978,
cymbaria Hack.	20	Birari 1980a & 1981, Mehra 1982
dacrusii Birari	60	Birari 1980a & 1981
hookeri (Griseb.) A. Camus	20	Christopher 1978 Birari 1980a & 1981
intermedia (Hack.) Bor.	40	Mehra & Kalia 1976, Kalia 1978, Birari
laxa (Anderss.) A. Camus	20	1980a & 1981, Mehra 1982
longispatha Raiz. & Jain	50	Kalia 1978, Mehra 1982
mooneyi Bor.	20	Christopher 1978
quadrivalvis (L.) Kuntze	18 40 80	Birari 1980a & 1981 Birari 1980a & 1981 Sahni & Bir 1985
strigosa (Ham.) A. Camus	20	Birari 1980a & 1981
subsericans (Nees) Ridley	40 54	Christopher 1978 Birari 1980a & 1981
tremula (Nees) Hook.	20	Mehra & Kalia 1975, Kalia 1978, Mehra
triandra Forsk.	20 20+4B 20+6B 21 22 24 30 40 40+4B 42 50 59 60 80 90 100 110	1982 Kalia 1978, Mehra 1982 Christopher 1978, Birari 1980a & 1981 Christopher 1978, Spies & Du Plessis 1986a, Liebenberg 1986, Fossey & Liebenberg 1987 Liebenber 1986 Liebenber 1986, Liebenberg 1990 Liebenberg 1986, Fossey & Liebenberg 1987, Liebenberg 1990 Fossey & Liebenberg 1987 Fossey & Liebenberg 1987 Du Plessis & Spies 1988 Mehra & Sharma 1975, Kalia 1978, Gill et al. 1980 & 1980a, Birari 1981, Mehra 1982, Liebenberg 1990, Liebenberg et al. 1993 Liebenberg 1990 Liebenberg 1990 Liebenberg 1990, Liebenberg et al. 1993 Liebenberg 1990 Sharma & Sharma 1979, Liebenberg 1990, Sinha et al. 1990, Fossey &
triandra Forsk. var. hispida (Nees) Stapf	40, 60	Liebenberg 1992, Liebenberg et al. 1993
villosa (Poir) A. Camus	40	Shanthamma 1976, Gill et al. 1980a, Birari 1981 Gill et al. 1980a, Birari 1981 Liebenberg et al. 1993 Gill et al. 1980a, Birari 1981 Tateoka 1965c Kalia 1978, Gill et al. 1980a & 1981, Mehra 1982

Trachypogon ligularis Nees montufari (HBK) Nees spicatus (L.F.) Kuntze	20 60 20 40	Davidse & Pohl 1972 Davidse & Pohl 1972 Olorode 1975, Strydom et al. 2000 Dujardin 1978
Vetiveria lawsonii (Hook.f.) Blatt & McCann nigritiana (Benth) Stapf zizanoides (L.) Nash	20 20 20	Muniyamma et al. 1976 Olorode 1975, Dujardin 1978 Mitra & Datta 1967, Saxena & Gupta 1970, Pohl & Davidse 1971, Sindhe 1977, Sharma et al. 1978, Shanthamma et al. 1978, Christopher 1978, Kalia 1978, Mehra 1982, Bir & Sahni 1983, Bir et al. 1987, Lavania 1985, Sinha et al. 1990, Ahsan et al. 1994
Vossia cuspidata (Roxb.) Griff.	20 40	Mehra & Kalia 1976, Kalia 1978, Mehra 1982 Dujardin 1978
Zea diploperennis  perennis (Hitche) Reeves & Mangelsdorf mays L.  mays L. cultivars  mays subsp. amylacea (Sturlev.) Zhuk. mays subsp. ceratina (Kuleshov) Zhuk. mays subsp. everta (Sturtev.) Zhuk. mays subsp. indentata (Sturtev.) Zhuk. mays subsp. indurata (Sturtev.) Zhuk. mays subsp. sacharata (Sturtev.) Zhuk. mays subsp. semidentata Kuleshov mays subsp. tunicata Sturtev.	20  20+1B 40 20  20+0-6B 21, 22 20 20+1-3B 20 20 20 20 20 20 20 20 20	Molina 1982, Paupulet & Galinat 1982, Molina 1985 Molina 1982 Kato 1984, Molina 1985, Molina 1986 Weber 1969, Das 1970, Vochova 1978, Chokairi & Gorenflot 1979, Sarr & Gorenflot 1979, Palmer 1971, Filion & Walden 1973, Maquire 1974, Sharma & Agarwal 1980, Joshi & Ranjekar 1982, Sapre & Barve 1983a, Maquire 1982, Sharma & Sharma 1984, Khuong & Schubert 1985, Aguiar Perecin 1985, Zhang et al. 1985, Khuong & Sharma 1986, Chen et al. 1987, Heslop-Harrison et al. 1988, Chen 1988, Laurie & Bennett 1989, Ahsan et al. 1994, Inagaki & Mujeeb-Kazi 1995, Carvalho & Saraiva 1997 Mogford 1981a, Aguiar Perecin 1985 Weber 1969 Sharma & Sharma 1986 Sharma & Sharma 1986 Zhang & Li 1990 Zhang & Li 1990 Zhan et al. 1987, Zhang & Li 1990 Zhang & Li 1990 Zhang & Li 1990 Zhang & Li 1990 Zhang & Li 1990 Zhang & Li 1990 Zhang & Li 1990eddc3

**Appendix B:** CLUSTAL W (1.74) multiple sequence alignment data of the Andropogoneae specimens for *tnrL* intron.

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Andropogonappendiculatus6543      -----TCGGTATA--GGTTTTGTGG-----TTTGGGGG-----
Andropogonappendiculatus6677      GAA-TCGGTAG--CGCTACGTTGTATTCCCG--CCTTCCCAGGA-ACC
Andropogonappendiculatus6946      GAA-TGGAAACCTTGCTAGGTTTAACTCCCN---AAATCCAGAGCCCC
Andropogonappendiculatus6949      GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Aristidaadscencianis6713          GAA-TCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Bothriochloa7489                   GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Brachiaria                          GAA-TCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Brachyariaserrata6654               GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Cymbopogonexcavatus5877             GAAATCGGTACA-CGCTACGTTCAAGTCCCT--CTATCCCAGCCGAAA
Cymbopogonexcavatus6577             -----
Cymbopogonmarginatus4397           GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Cymbopogonprolixus6682              GAA-TCGGACC---GCTACGTTAAGTCCCA---ATTCCAGAGAAACCC
Diheteropogonfiliifolius5872       GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Elionurusmuticus5074                GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Elionurusmuticus6954                GAA-TCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Helictotrichonlongifolium6560      GAA-TCGGTAGA-CGCTACGTTAAGT---CCC---TCCTCCCAGCCGAAA
Hemarthriaaltissima5698            GAA-TGGAAACCTTGCTAAGTGTAACTTCCC---AATCCCAGAACCT
Hyparrheniaanamesa4841              GAA-TCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Hyparrheniadissoluta5145            GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Hyparrheniahirta6550                GAA-TCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Hyparrheniapilgeriana4649           GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Hyparrheniapilgeriana4738           GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Imperatacylindrica5114              GAA-TCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Imperatacylindrica6582              GAA-TCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Miscanthuscapensis6655              GTA-TGGGAAAC---TGCTAAGTGTAACTCCCA---ATTTCCAGAACCT
Panicumschinzii6652                 GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Paspalum5                            GAA-TCGGTAGA-CGCTACGTTCAAGTCCCT--CTATCCCAGCCGAAA
Paspalumdilatatum6617               GAA-TCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Phacelurusfranksae4705              GAAATCGGTAGA-CGCTACGTTCAAGTCCCT--CCTTCCCAGCCGAAA
Setarianigrostis6567                GAAATCGGTAGA-CGCTACGTTTAAATCCCC--CATTTCCAGCCGAAA
Sorghumbicolour6589                 GAA-T-GGT-GA-CGCTACGTT--ACTCCCT--CCTTCCCAGCCGAAA
Sorghumhalepense9318                GAA-TCGGAACC-CGCTACGTTAAGT---CCC---ATTC--AGC-GAAA
Themedatriandra6555                 GAAATCGGTAGA-CGCTACGTTCAAGTCCCGACATAAGCTGCCACCT
Trachypogonspicatus4846              GAAATCGGTAGA-CGCTACGTTCAATCCCT--CCTTCCCAGCCGAAA
Trachypogonspicatus6580             GAA-TGG-AAC--TGCTACGTTCAAGTGCC-----TCCTAGGGGAAA
Zea                                   TTTCT-ATCTA--AGTGAACCTCCAATTT-AGAAGTAGT-TAATAACT
Icylindrical                         -----TTCCA--AATTCAGAGAA-ACC
Sacharumofficinatum                  -----TTCCA--AATTCAGAGAA-ACC
Sorghumhalepense                     -----TTCCA--AATTCAGAGAA-ACC
Ttrianadra                            -----TTCCA--AATTCAGAGAA-ACC
Pschinzii                             ATG-GAAACCT--GCTAAGTGGTAACTTCCA--AATTCAGAGAA-ACC

Andropogonappendiculatus6543      ---GGT---GTGTGTTGGGCAA-TGTTGAGATA-TTCCCTTTTTTTTGAA
Andropogonappendiculatus6677      --TGGAATGATA-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Andropogonappendiculatus6946      --TGGAATGAAA-G-TGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Andropogonappendiculatus6949      CTGGAAT-GACG-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Aristidaadscencianis6713          -TGGGT-GGATGCAATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Bothriochloa7489                   -TGGGAAAGATGCAATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Brachiaria                          -TCGGAATGACG-CATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Brachyariaserrata6654               -TGGGAATGACGCAATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Cymbopogonexcavatus5877             -TCGGTA-GATT-CTACGGCAA-TGCCCGCGGGCTCC-CTTTATTTAAA
Cymbopogonexcavatus6577             -----
Cymbopogonmarginatus4397           -TCGGTA-GATT-AATGGGCAA-TCCTGAGCCAACTCC-CTTTATTTGAA
Cymbopogonprolixus6682              --TGGAATGAAA-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Diheteropogonfiliifolius5872       -TCGGAA-GACG-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Elionurusmuticus5074                -TCGGTG-GATG-AATGGGCAA-T-----CCA-CCGTCCTCCCTTG
Elionurusmuticus6954                ATCGGTA-GACG-AATGGGCAA-TCCTGAACCAATCC-CTTTATTTGAAA
Helictotrichonlongifolium6560      -TTGG----ATG-AATGGGCAA-TCCTGAGCCAAATCC-GTGTTTTCTGA
Hemarthriaaltissima5698            --CGGAATGAAA-A-TGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Hyparrheniaanamesa4841              -TCGGTA-GACG-CATGGGCAA-TCCTGA-CCATCTCC-CTTT-TTT--A
Hyparrheniadissoluta5145            -TCGGTA-GAGA-AATGGGCAA-TCCTGA-CCAAATCC-CTTTTTTTGAAA
Hyparrheniahirta6550                -TGGAAT-GACA-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTTGAAA
Hyparrheniapilgeriana4649           -TCGGTA-GACG-CTAGGGCAA-TCCTGAGCCATATCC-CTTGATTTGAA

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*Hyparrheniampilgeriana*4738  
*Imperatacylindrica*5114  
*Imperatacylindrica*6582  
*Miscanthuscapensis*6655  
*Panicumschinzii*6652  
*Paspalum*5  
*Paspalumdilatatum*6617  
*Phacelurusfranksae*4705  
*Setarianigrostis*6567  
*Sorghumbicolour*6589  
*Sorghumhalepense*9318  
*Themedatriandra*6555  
*Trachypogonspicatus*4846  
*Trachypogonspicatus*6580  
*Zea*  
*Icylindrical*  
*Sacharumofficinarium*  
*Sorghumhalepense*  
*Ttriandra*  
*Pschinzii*

-TCGGTA-GATG-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA  
-TCGGTT-GACG-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA  
--TGGAATGATA-AATGGGCAA-TCCTGAGCCAAATCCACTTTTTTGAAA  
--TGGAATGAAA-AATGGGCAA-TCCTGAGCCTTTGGG-GGT----GAAA  
-TCGGA-TGACG-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA  
-TCGGAA-GACG--CTGGGCTCCTGCTCCTCCATATCC-CTTTAATTCAA  
-TGGGA-TGATG-CATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA  
-TCGGAT-GAGG-AATGGGCAA-TCCTGA-CCAAATCC-CTTTTTTGAAA  
-TGGAA-TGATA-CATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA  
--TGGAATGATA-AATGGGCAA-TCCTGAGCCAAATCCACTTTTTTCAA  
-TCGG--TGATG-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA  
TGGCCTTAATGTCTAGCTAACCACGCGCAATAT-ATGATCTG--A  
-TCGGTG-GATT-AAAGGGCAA-TGGTGAGC---CTCA-CTTTTTTTTTT  
--TCGGATGATA-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA  
AAGATTAATAATTAAGATCTGACTTTACAGCCCTATAT-ATAATTTT--T  
-TGGAA-TGAAA-AATGGGCAA-TCCTGAGCCAAATCCACTTTTTTGAAA  
CTGGAA-TGAAA-AATGGGCAA-TCCTGAGCCAAATCCCTTTTTTGAAA  
CTGGAA-TGAAA-AATGGGCAA-TCCTGAGCCAAATCCACTTTTTTCAA  
CTGGAA-TGAAA-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA  
CTGGAA-TGAAA-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA

*Andropogonappendiculatus*6543  
*Andropogonappendiculatus*6677  
*Andropogonappendiculatus*6946  
*Andropogonappendiculatus*6949  
*Aristidaadscencianis*6713  
*Bothriochloa*7489  
*Brachiaria*  
*Brachyariaserrata*6654  
*Cymbopogonexcavatus*5877  
*Cymbopogonexcavatus*6577  
*Cymbopogonmarginatus*4397  
*Cymbopogonprolixus*6682  
*Diheteropogonfilifolius*5872  
*Elionurusmuticus*5074  
*Elionurusmuticus*6954  
*Helictotrichonlongifolium*6560  
*Hemarthriaaltissima*5698  
*Hyparrheniaanamesa*4841  
*Hyparrheniadissoluta*5145  
*Hyparrheniahirta*6550  
*Hyparrheniampilgeriana*4649  
*Hyparrheniampilgeriana*4738  
*Imperatacylindrica*5114  
*Imperatacylindrica*6582  
*Miscanthuscapensis*6655  
*Panicumschinzii*6652  
*Paspalum*5  
*Paspalumdilatatum*6617  
*Phacelurusfranksae*4705  
*Setarianigrostis*6567  
*Sorghumbicolour*6589  
*Sorghumhalepense*9318  
*Themedatriandra*6555  
*Trachypogonspicatus*4846  
*Trachypogonspicatus*6580  
*Zea*  
*Icylindrical*  
*Sacharumofficinarium*  
*Sorghumhalepense*  
*Ttriandra*  
*Pschinzii*

AAAGAAAGCGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACTT-CCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
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AAACTT-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GC-GGTT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACTA-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACTT-GCGG-TCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAACGAGTTCT--CGAAC TAGAATACAAAGGAAAAGGATAGGTGC  
AAACAA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
--ACCA-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GCG-----CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACCA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAAATGC  
AAACTA-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
TGACTA-CATGGTCTACCTAATGATGATCCAAAGGTAATGGATAGGTGC  
AAACAA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACCC-CTGCTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AA-CAA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAA-----GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AA-CAA-GTGGTT-T--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
GGACAAAAGA---TC---CATATTGGTC-----GGTCCCGTAGCG-TCT  
AAACTA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GTGGTTCT--CTAAGTAGAATCCAAAGGAAAAGGATAGGTGC  
TGTTACATT---TC---TGTTATGTAA-----GCCCACTTAG---CTC  
AAACAA-GTG-----CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAA-----GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GCGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC  
AAACAA-GTGGTTCT--CAAAC TAGAACCCAAAGGAAAAGGATAGGTGC

*Andropogonappendiculatus*6543  
*Andropogonappendiculatus*6677  
*Andropogonappendiculatus*6946  
*Andropogonappendiculatus*6949  
*Aristidaadscencianis*6713  
*Bothriochloa*7489  
*Brachiaria*

AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
AGAGACTCATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
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AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----



*Brachyariaserrata*6654  
*Cymbopogonexcavatus*5877  
*Cymbopogonexcavatus*6577  
*Cymbopogonmarginatus*4397  
*Cymbopogonprolixus*6682  
*Diheteropogonfilifolius*5872  
*Elionurusmuticus*5074  
*Elionurusmuticus*6954  
*Helictotrichonlongifolium*6560  
*Hemarthriaaltissima*5698  
*Hyparrheniaanamesa*4841  
*Hyparrheniadissoluta*5145  
*Hyparrheniahirta*6550  
*Hyparrheniapilgeriana*4649  
*Hyparrheniapilgeriana*4738  
*Imperatacylindrica*5114  
*Imperatacylindrica*6582  
*Miscanthuscapensis*6655  
*Panicumschinzii*6652  
*Paspalum*5  
*Paspalumdilatatum*6617  
*Phacelurusfranksae*4705  
*Setarianigrostis*6567  
*Sorghumbicolour*6589  
*Sorghumhalepense*9318  
*Themadatriandra*6555  
*Trachypogonspicatus*4846  
*Trachypogonspicatus*6580  
*Zea*  
*Icylindrical*  
*Sacharumofficinarium*  
*Sorghumhalepense*  
*Ttriandra*  
*Pschinzii*

*Andropogonappendiculatus*6543  
*Andropogonappendiculatus*6677  
*Andropogonappendiculatus*6946  
*Andropogonappendiculatus*6949  
*Aristidaadscencianis*6713  
*Bothriochloa*7489  
*Brachiaria*  
*Brachyariaserrata*6654  
*Cymbopogonexcavatus*5877  
*Cymbopogonexcavatus*6577  
*Cymbopogonmarginatus*4397  
*Cymbopogonprolixus*6682  
*Diheteropogonfilifolius*5872  
*Elionurusmuticus*5074  
*Elionurusmuticus*6954  
*Helictotrichonlongifolium*6560  
*Hemarthriaaltissima*5698  
*Hyparrheniaanamesa*4841  
*Hyparrheniadissoluta*5145  
*Hyparrheniahirta*6550  
*Hyparrheniapilgeriana*4649  
*Hyparrheniapilgeriana*4738  
*Imperatacylindrica*5114  
*Imperatacylindrica*6582  
*Miscanthuscapensis*6655  
*Panicumschinzii*6652  
*Paspalum*5  
*Paspalumdilatatum*6617  
*Phacelurusfranksae*4705  
*Setarianigrostis*6567  
*Sorghumbicolour*6589  
*Sorghumhalepense*9318  
*Themadatriandra*6555  
*Trachypogonspicatus*4846  
*Trachypogonspicatus*6580

*Zea* -----  
*Icylindrical* -----  
*Sacharumofficinorum* -----  
*Sorghumhalepense* -----  
*Ttriandra* -----  
*Pschinzii* TTGGTAGTGGAACTCCCTCGAAATTATAGAAAGAAGGGCTTTATA-----

*Andropogonappendiculatus6543* -----AACGATTAATCACA--  
*Andropogonappendiculatus6677* -----AACGATTAATCACA--  
*Andropogonappendiculatus6946* -----AACGATCAATCACA--  
*Andropogonappendiculatus6949* -----AACGATTAATCACA--  
*Aristidaadscencianis6713* CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--  
*Bothriochloa7489* -----AACGATTAATCACA--  
*Brachiaria* -----  
*Brachyariaserrata6654* CATTTAATACACACGTATAGATACTGACATAGCAAACGATTAATCAG--  
*Cymbopogonexcavatus5877* CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--  
*Cymbopogonexcavatus6577* -----AACGATTAATCACAAA  
*Cymbopogonmarginatus4397* -----AACGATTAATCACA--  
*Cymbopogonprolixus6682* -----AACGATTAATCACA--  
*Diheteropogonfilifolius5872* -----AACGATTAATCACA--  
*Elionurusmuticus5074* -----AACGATTAATCACA--  
*Elionurusmuticus6954* -----AACGATTAATCACA--  
*Helictotrichonlongifolium6560* CACCTAATAAACACGTATAGATACTGGCATAGCAAACGATTAATCAG--  
*Hemarthriaaltissima5698* CATCTTATATACACGCATATATACTGACATAGCAAACGATTAATCAG--  
*Hyparrheniaanamesa4841* -----AACGATTAATCACA--  
*Hyparrheniadissoluta5145* -----AACGATTAATCACA--  
*Hyparrheniahirta6550* -----AACGATTAATCACA--  
*Hyparrheniapilgeriana4649* -----AACGATTAATCACA--  
*Hyparrheniapilgeriana4738* -----AACGATTAATCACA--  
*Imperatacylindrica5114* -----AACGATTAATCACA--  
*Imperatacylindrica6582* -----AACGATTAATCACA--  
*Miscanthuscapensis6655* -----GCAGCTTAATCACA--  
*Panicumschinzii6652* CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCATA--  
*Paspalum5* AATACAC---CACGTATAGATACTGACATACCAAACATATTATTCACA--  
*Paspalumdilatatum6617* CATCTAATATACACGCATATATACTGACATAGCAAACGATTAATCAG--  
*Phacelurusfranksae4705* -----AACGATTAATCACA--  
*Setarianigrostis6567* CATTTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--  
*Sorghumbicolour6589* -----AACGATTAATCACA--  
*Sorghumhalepense9318* CATTTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--  
*Themedatriandra6555* -----ACGAGT-TTCAGG--  
*Trachypogonspicatus4846* -----TACGATTAATCACA--  
*Trachypogonspicatus6580* -----AACGATTAATCACA--  
*Zea* -----CCGATT-TTTTTT--  
*Icylindrical* -----AACGATTAATCACA--  
*Sacharumofficinorum* -----AACGATTAATCACA--  
*Sorghumhalepense* -----AACGATTAATCACA--  
*Ttriandra* -----AACGATTAATCACA--  
*Pschinzii* CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCATA--

*Andropogonappendiculatus6543* GAACCCATATTATAATATAGGTTCTTTATTTGATTT-----TT  
*Andropogonappendiculatus6677* GAACCCATATTATAATATAGGTTCTTTATTTGATTT-----TT  
*Andropogonappendiculatus6946* GAACCCATATTATAATATAGGTTCTTTATTTGATTT-----TT  
*Andropogonappendiculatus6949* GAACCCATATTATAATATAGGTTCTTTATTTGATTT-----TT  
*Aristidaadscencianis6713* GAACCCATATCATAATATAGGTTCTTTATTTTATAT-----TTTTTTT  
*Bothriochloa7489* GAACCCATATTATAATATAGGTTCTTTATTTTATTT-----TT  
*Brachiaria* GAACCCATATCATAATAGGGGTTCTTTATTTTATAT-----TTTTTTCT  
*Brachyariaserrata6654* GAACCCATATCATAATATAGGTTCTTTATTTTATATATATATTTTTTTTT  
*Cymbopogonexcavatus5877* GAACCCATATTATAATATAGGTTCTTTATTTT-----T  
*Cymbopogonexcavatus6577* -----CATNNGGCTACGTGTTGCGCAGTCG-----  
*Cymbopogonmarginatus4397* GAACCCATATTATAATATAGGTTCTTTATTTT-----T  
*Cymbopogonprolixus6682* GAACCCATATTATAATATAGGTTCTTTATTTTATTT-----TT  
*Diheteropogonfiliifolius5872* GAACCTATATTATAATATAGGTTCTTTATTTTATTT-----TT  
*Elionurusmuticus5074* GAACCCATATTATAATATAGGTTCTTTATTTTATTT-----TT  
*Elionurusmuticus6954* GAACCCATATTATAATATAGGTTCTTTATTTTATTT-----TT  
*Helictotrichonlongifolium6560* GAACCCATATTATAATATAGGTTCTTTATTTTATTT-----TAGAATG  
*Hemarthriaaltissima5698* GAACCCATATCATAATATAGGTTCTTTATTTTATTT-----TT  
*Hyparrheniaanamesa4841* GAACCTATATTATAATATAGGTTCTTTATTTTATTT-----TT  
*Hyparrheniadissoluta5145* GAACCCATATTATAATATAGGTTCTTTATTTTATTT-----TT  
*Hyparrheniahirta6550* GAACCCATATTATAATATAGGTTCTTTATTTTATTT-----TT  
*Hyparrheniapilgeriana4649* GAACCCATATTATAATATAGGTTCTTTATTTTATTT-----TT





*Sacharumofficinatum* -GAGTAATCA-AATCCTT-CAATTCATTGTTTTTCG----AGATCTTTT  
*Sorghumhalepense* -GAGTAATCA-AATCCTT-CAATTCATTGTTTTTCG----AGATCTTTT  
*Ttriandra* -GAGTAATCA-AATCCTT-CAATTCATTGTTTTTCG----AGATCTTTT  
*Pschinzii* -GAGTAATCA-AATCCTT-CAATTCATTGTTTTTG-----AGATCTTCA

*Andropogonappendiculatus6543* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Andropogonappendiculatus6677* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Andropogonappendiculatus6946* CCTCAGTGGATTAATC-GGACGAGGATAAAGAGA--GACTAACCA  
*Andropogonappendiculatus6949* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Aristidaadscencianis6713* AAAAAGTGG-ATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Bothriochloa7489* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Brachiaria* AAAAAGTGG-ATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Brachyariaserrata6654* AAAAAGTGGGATTAATC--GACGAGGATAAAGAGA--GAGTCCCA  
*Cymbopogonexcavatus5877* TAAAAGCGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Cymbopogonexcavatus6577* ATAAGATGTGATAGTC-GAAAGTATACATAAAGAATGACGTACC  
*Cymbopogonmarginatus4397* TAAAAGCGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Cymbopogonprolixus6682* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Diheteropogonfilifolius5872* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Elionurusmuticus5074* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Elionurusmuticus6954* TAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Helictotrichonlongifolium6560* AACAGGTGATTAATCCGGACGAGGACAAAGAGA--GAGTCCCA  
*Hemarthriaaltissima5698* AAAAAGTGGATTAAGC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Hyparrheniaanamesa4841* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Hyparrheniadissoluta5145* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Hyparrheniahirta6550* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Hyparrheniapilgeriana4649* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Hyparrheniapilgeriana4738* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Imperatacylindrica5114* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Imperatacylindrica6582* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Miscanthuscaspensis6655* AAAAAGTGGATAAATC-GGCGGAGGATTAAGCGT--GGTTCATA  
*Panicumschinzii6652* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Paspalum5* ANNAGTNGTNTTAAATCCNGAGGGGGGTAAGAAAGNNAAGNNCCNN  
*Paspalumdilatatum6617* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Phacelurusfranksae4705* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Setarianigrostis6567* AAAAAGTGG-ATTAATC--GACGAGGATAAAGAGA--GAGTCCCA  
*Sorghumbicolour6589* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Sorghumhalepense9318* GAAAGTGG-TATACTC--GAATCGGACGAGGATAA--GAGTCCCA  
*Themedatriandra6555* AAAAAGTGGATTAATCCGGACGAGGATAAAGAGA--GATTCCCA  
*Trachypogonspicatus4846* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Trachypogonspicatus6580* AAAAAGTGGATTAATC-GGATGAGGATAAAGAGA--GAGTAGAA  
*Zea* GA----TGGTGAAT--GGTAGAC----ACGCGA--GACTCAA  
*Icylindrical* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Sacharumofficinatum* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Sorghumhalepense* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Ttriandra* AAAAAGTGGATTAATC-GGACGAGGATAAAGAGA--GAGTCCCA  
*Pschinzii* AAAAAGTGGATTAATC-GAACGAGGATAAAGAGA--GAGTCCCA

*Andropogonappendiculatus6543* TTCTACATGTCAATACTGACAACATT--AATTTCTAGTAAAA  
*Andropogonappendiculatus6677* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Andropogonappendiculatus6946* TTTTGAATCCCAACGGAATCGGG--ACACATAACTTTCAAAT  
*Andropogonappendiculatus6949* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Aristidaadscencianis6713* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Bothriochloa7489* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Brachiaria* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Brachyariaserrata6654* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Cymbopogonexcavatus5877* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Cymbopogonexcavatus6577* GTCAACGATTTTCNGG--NNTATAGTTCTTTATATTTTANANA  
*Cymbopogonmarginatus4397* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Cymbopogonprolixus6682* TTCTACATGTCAATACTGACGGC--AGATGGATTTGAATCCC  
*Diheteropogonfilifolius5872* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Elionurusmuticus5074* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Elionurusmuticus6954* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Helictotrichonlongifolium6560* TTCTACATGTCAATACTG--AACAAATGAAATTTCTAGTAAAA  
*Hemarthriaaltissima5698* TGCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Hyparrheniaanamesa4841* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Hyparrheniadissoluta5145* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Hyparrheniahirta6550* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Hyparrheniapilgeriana4649* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA  
*Hyparrheniapilgeriana4738* TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA

<i>Imperatacylindrica</i> 5114	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Imperatacylindrica</i> 6582	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Miscanthuscapensis</i> 6655	TTCGGTATGTCAATGCGATCGTC--TGACTTTACTTTCTAAC
<i>Panicumschinzii</i> 6652	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Paspalum</i> 5	TTNGAGGTGTCAATACNGGANNCAATAGGATTTTNTNGNNA
<i>Paspalumdilatatum</i> 6617	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Phacelurusfranksae</i> 4705	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Setarianigrostis</i> 6567	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Sorghumbicolour</i> 6589	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Sorghumhalepense</i> 9318	TTCTACATGTCAATACTGACAACAATGAA- TTTCTAGTAAAA
<i>Themedatriandra</i> 6555	TCCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Trachypogonspicatus</i> 4846	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Trachypogonspicatus</i> 6580	TTATACATGTCNAT-CTGACAACAA-GAATAGAGGTTATGAT
<i>Zea</i>	ATCT-CGTG-CTAAAGAGCGTGAGGTTTCGAGTCCCT-CTTCAA
<i>Icylindrical</i>	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Sacharumofficinarum</i>	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Sorghumhalepense</i>	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Ttriandra</i>	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Pschinzii</i>	TTCTACATGTCAATACTGACAACAATGAAATTTCTAGTAAAA
<i>Andropogonappendiculatus</i> 6543	GGA
<i>Andropogonappendiculatus</i> 6677	GGG
<i>Andropogonappendiculatus</i> 6946	TCC
<i>Andropogonappendiculatus</i> 6949	GGA
<i>Aristidaadscencianis</i> 6713	GGA
<i>Bothriochloa</i> 7489	GGA
<i>Brachiaria</i>	GGA
<i>Brachyariaserrata</i> 6654	GGA
<i>Cymbopogonexcavatus</i> 5877	GGA
<i>Cymbopogonexcavatus</i> 6577	ATG
<i>Cymbopogonmarginatus</i> 4397	GGA
<i>Cymbopogonprolixus</i> 6682	TCC
<i>Diheteropogonfilifolius</i> 5872	GGA
<i>Elionurusmuticus</i> 5074	GGA
<i>Elionurusmuticus</i> 6954	GGA
<i>Helictotrichonlongifolium</i> 6560	GGA
<i>Hemarthriaaltissima</i> 5698	GGA
<i>Hyparrheniaanamesa</i> 4841	GGA
<i>Hyparrheniadissoluta</i> 5145	GGA
<i>Hyparrheniahirta</i> 6550	GGA
<i>Hyparrheniapilgeriana</i> 4649	GGA
<i>Hyparrheniapilgeriana</i> 4738	GGA
<i>Imperatacylindrica</i> 5114	GGA
<i>Imperatacylindrica</i> 6582	GGG
<i>Miscanthuscapensis</i> 6655	ACC
<i>Panicumschinzii</i> 6652	GGA
<i>Paspalum</i> 5	GAA
<i>Paspalumdilatatum</i> 6617	GGA
<i>Phacelurusfranksae</i> 4705	GGA
<i>Setarianigrostis</i> 6567	GGA
<i>Sorghumbicolour</i> 6589	GG-
<i>Sorghumhalepense</i> 9318	GGA
<i>Themedatriandra</i> 6555	GGG
<i>Trachypogonspicatus</i> 4846	GGA
<i>Trachypogonspicatus</i> 6580	TGG
<i>Zea</i>	GGC
<i>Icylindrical</i>	GGA
<i>Sacharumofficinarum</i>	GGA
<i>Sorghumhalepense</i>	GGA
<i>Ttriandra</i>	GGA
<i>Pschinzii</i>	GGA

**Appendix C:** CLUSTAL W (1.74) multiple sequence alignment data of the Andropogoneae specimens for *tnrL-F* intergenic spacer.

<i>Diheteropogonfillifolius</i> 5872	CCTCC
<i>Hyparrheniadissoluta</i> 5145a	ACCCC
<i>Hyparrheniafillipendula</i> 5144	GC CCC
<i>Sorghumhalepense</i> 9318	GACCC
<i>Hyparrheniapilgeriana</i> 4738	GACCC
<i>Trachypogonspicatus</i> 4846	GACCA
<i>Phacelurusfranksae</i> 4705	GACCA
<i>Hyparrheniaquarrei</i> 6586	GACCA
<i>Andropogonappendiculatus</i> 6692	GATCC
<i>Elionurusmuticus</i> 6559	GACTC
<i>Elionurusmuticus</i> 6539	GACTC
<i>Elionurusmuticus</i> 6954	GACTC
<i>Hyparrheniahirta</i> 6656f	GAACC
<i>Themedatriandra</i> 6383	GACCC
<i>Themedatriandra</i> 6555	GACCC
<i>Themedatriandra</i> 6525	GACCC
<i>Andropogonschirensis</i> 6581	GACCC
<i>Imperatacylindrica</i> 5114a	GACCC
<i>Bothriochloainsculpta</i> 5681	GACCC
<i>Bothriochloa</i> 7489	GACCC
<i>Heteropogoncontortus</i> 6628	GACC-
<i>Hyparrheniapilgeriana</i> 4603	GACCC
<i>Hemarthriaaltissima</i> 5698	GACCA
<i>Andropogonamethystinus</i> 6679	GACCC
<i>Andropogonamethystinus</i> 6685	GACCC
<i>Andropogondistachyos</i> 6684	GACCC
<i>Hyparrheniatamba</i> 6585	GACCC
<i>Hyparrheniarudis</i> 6576	GACCA
<i>Hyparrheniatamba</i> 6537a	GACC-
<i>Cymbopogonexcavatus</i> 6577	GACCC
<i>Cymbopogonsp</i> 5800	GACCC
<i>Andropogonappendiculatus</i> 6677	GACCA
<i>Heteropogoncontortus</i> 6942	GACCA
<i>Andropogoneucomus</i> 6587	GACCC
<i>Cymbopogondieterlenii</i> 6642	GACCC
<i>Cymbopogonprolixus</i> 6682	GACCC
<i>Imperatacylindrica</i> 6632a	GACCA
<i>Trachypogonspicatus</i> 6580	GACCC
<i>Cymbopogonplurinodus</i> 6406	GACCC
<i>Hyparrheniahirta</i> 6618	GACCC
<i>Hemarthria</i> 9319	GACCC
<i>Sorghumbicolour</i> 6588	GACCA
<i>Miscanthuscapensis</i> 6655	GACCC
<i>Miscanthuscapensis</i> 6584	GACCC
<i>Heteropogoncontortus</i> 6538	GACCC
<i>Hyparrheniaanamesa</i> 4807	GACCC
<i>Pdilatum</i> 6617	GACCC
<i>Zea</i>	ATAAT
<i>Icylindrical</i>	AAATC
<i>Sacharumofficinarum</i>	AAATC
<i>Sorghumhalepense</i>	AAATC
<i>Ttriandra</i>	AAATC
<i>Pschinzii</i>	AAATC
<i>Diheteropogonfillifolius</i> 5872	CNTTTT TAGGACCAAGTTCAG---GGAGA---TAGGGGTTTGAA
<i>Hyparrheniadissoluta</i> 5145a	TTATAGTCTACACAGTCC--TGCACA---AAAGGCACTCCTT
<i>Hyparrheniafillipendula</i> 5144	TC TTAGCCCGACCCAAGTTTC--GGCACA---TAGGGCCCTATTT
<i>Sorghumhalepense</i> 9318	TCTTACCGGACCCCATATA---CCGATT---TCCCGGTATTTT
<i>Hyparrheniapilgeriana</i> 4738	TCTTACCGGACCC--AAATTTTC---GGGAGA---TAGGGGTATTTT
<i>Trachypogonspicatus</i> 4846	TCGTTGTCCANAAAAGTTTC---GCANA---TAGGGGTATTTT
<i>Phacelurusfranksae</i> 4705	AAGT TTTGAGACCAAGTGCG--GGGAGA---TAGGGGTCATTGA
<i>Hyparrheniaquarrei</i> 6586	AGTTCAGGAGATAGGGGGGTG--TGTTT---TAAGGTCAGGTA
<i>Andropogonappendiculatus</i> 6692	ATATTTTGGGACCAAATTCG---GGAGA---TAGGGG---ATAT
<i>Elionurusmuticus</i> 6559	CGATTGTGCGACCAAAGTTCA---GGAGA---TAGGGG---AAAA
<i>Elionurusmuticus</i> 6539	CGATTGGCGGCCCATTTTCG---GGGA---CANGGCCNATTA

*Elionurusmuticus*6954  
*Hyparrheniahirta*6656f  
*Themedatriandra*6383  
*Themedatriandra*6555  
*Themedatriandra*6525  
*Andropogonschirensis*6581  
*Imperatacylindrica*5114a  
*Bothriochloainsculpta*5681  
*Bothriochloa*7489  
*Heteropogoncontortus*6628  
*Hyparrheniapilgeriana*4603  
*Hemarthriaaltissima*5698  
*Andropogonamethystinus*6679  
*Andropogonamethystinus*6685  
*Andropogondistachyos*6684  
*Hyparrheniatamba*6585  
*Hyparrheniarudis*6576  
*Hyparrheniatamba*6537a  
*Cymbopogonexcavatus*6577  
*Cymbopogon*5800  
*Andropogonappendiculatus*6677  
*Heteropogoncontortus*6942  
*Andropogoneucomus*6587  
*Cymbopogondieterlenii*6642  
*Cymbopogonprolixus*6682  
*Imperatacylindrica*6632a  
*Trachypogonspicatus*6580  
*Cymbopogonplurinodis*6406  
*Hyparrheniahirta*6618  
*Hemarthria*9319  
*Sorghumbicolour*6588  
*Miscanthuscapensis*6655  
*Miscanthuscapensis*6584  
*Heteropogoncontortus*6538  
*Hyparrheniaanamesa*4807  
*Pdilatum*6617  
*Zea*  
*Icylindrical*  
*Sacharumofficinarum*  
*Sorghumhalepense*  
*Ttriandra*  
*Pschinzii*

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CGTCGACTTTATA-AGTCGT-GAG????????????????????

*Diheteropogonfillifolius*5872  
*Hyparrheniadissoluta*5145a  
*Hyparrheniafillipendula*5144  
*Sorghumhalepense*9318  
*Hyparrheniapilgeriana*4738  
*Trachypogonspicatus*4846  
*Phacelurusfranksae*4705  
*Hyparrheniaaquarrei*6586  
*Andropogonappendiculatus*6692  
*Elionurusmuticus*6559  
*Elionurusmuticus*6539  
*Elionurusmuticus*6954  
*Hyparrheniahirta*6656f  
*Themedatriandra*6383  
*Themedatriandra*6555  
*Themedatriandra*6525  
*Andropogonschirensis*6581  
*Imperatacylindrica*5114a  
*Bothriochloainsculpta*5681  
*Bothriochloa*7489  
*Heteropogoncontortus*6628  
*Hyparrheniapilgeriana*4603  
*Hemarthriaaltissima*5698  
*Andropogonamethystinus*6679  
*Andropogonamethystinus*6685  
*Andropogondistachyos*6684

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TTTGTA---GACCAAAGT-----CAGATAA---CATG  
TCGGGTC---AATTGCAGG-----GATAA-----ATAG  
TCGGACC---AAGTGCAGG-----GAG-----ATAG  
TCGGGTC---AATAGGACC-----ANGTTTCAG---GGAG  
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TA---TCATAAATAGGAGAAAAGAAAATAGTTACCCA-AATTC  
TCCCTCATCGGAAGAAAAGAAAATAAATATTTACCCA--ATTC  
TCC--CTTTAGGAAACAAGAATA---TAGTTACCCA-CACTC  
TCC--CTTTAGGAAACAAGAATA---TAGTTACCCA-CACTC  
TCC--CATTAGGAAACAAGAATA---TAGTTACCCA-CATTC



*Hyparrheniatamba*6585  
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*Hyparrheniatamba*6537a  
*Cymbopogonexcavatus*6577  
*Cymbopogon*sp5800  
*Andropogonappendiculatus*6677  
*Heteropogoncontortus*6942  
*Andropogoneucomus*6587  
*Cymbopogondieterlenii*6642  
*Cymbopogonprolixus*6682  
*Imperatacylindrica*6632a  
*Trachypogonspicatus*6580  
*Cymbopogonplurinodus*6406  
*Hyparrheniahirta*6618  
*Hemarthria*9319  
*Sorghumbicolour*6588  
*Miscanthuscapensis*6655  
*Miscanthuscapensis*6584  
*Heteropogoncontortus*6538  
*Hyparrheniaaamesa*4807  
*Pdilatum*6617  
*Zea*  
*Icylindrical*  
*Sacharumofficinarium*  
*Sorghumhalepense*  
*Ttriandra*  
*Pschinzii*

*Diheteropogonfillifolius*5872  
*Hyparrheniadissoluta*5145a  
*Hyparrheniafillipendula*5144  
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*Elionurusmuticus*6539  
*Elionurusmuticus*6954  
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*Themedatriandra*6555  
*Themedatriandra*6525  
*Andropogonschirensis*6581  
*Imperatacylindrica*5114a  
*Bothriochloainsculpta*5681  
*Bothriochloa*7489  
*Heteropogoncontortus*6628  
*Hyparrheniapilgeriana*4603  
*Hemarthriaaltissima*5698  
*Andropogonamethystinus*6679  
*Andropogonamethystinus*6685  
*Andropogondistachyos*6684  
*Hyparrheniatamba*6585  
*Hyparrheniarudis*6576  
*Hyparrheniatamba*6537a  
*Cymbopogonexcavatus*6577  
*Cymbopogon*sp5800  
*Andropogonappendiculatus*6677  
*Heteropogoncontortus*6942  
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*Cymbopogondieterlenii*6642  
*Cymbopogonprolixus*6682  
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*Trachypogonspicatus*6580  
*Cymbopogonplurinodus*6406  
*Hyparrheniahirta*6618  
*Hemarthria*9319  
*Sorghumbicolour*6588  
*Miscanthuscapensis*6655

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CCA-TCATTCTGAAACATAGAAA--TAGTAACCCA-ACTTC  
TCA-GCATTCCGACAGTAGAAA--TAGTTACCCA-AACTC  
TCA-ACATTAGTAAACAAGAAA--TAGTTACCCA-CATTC  
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TCA-GCTATCGGAAACGTAGAAA--TAGTTACCCA-AATTC  
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TCT--CTAACCAAAAACAAGAAG--ATAGTTGTCCC-CATTC  
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TCAA-CTTAGGAAAA-AAGAA--ATAGTTACCCA-GATTC  
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AAT--CTGGCGCTCTCCCTATC---TAATGAATA--CTTTA  
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AAA-CCCTCTTTTATTCCTAACC-ATAGTTGTAT-CCTTT  
??

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-CCACTGT--GGCTCTAAATATC----GGACC-ACGTTCAGGGAGAT  
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TAA-GTGA-TC-GAAAGA-GTAAGT--GAGGATGAGCAGTCTCACGC-  
TAG-GTGA-TC-GAAATGAGTAAGATGAGAAATGAGTTTCTCACGC-  
TAA-GTG-ATC-GAATCA-GTAAG-ATGAGAAAGTGTCTCCTCACGC-  
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TGA-GTG-ATC-GAAAGA-GTAAG-AGGAGAAAGTGTTCCTCACGC-  
TAA-GTG-ATC-GAAAGAC-TAAG-AGGAGAAAGTGTTCCTCACGC-  
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TTA-GCG-ATCGAGGAGA-GTAAG-ATGAGAAAGCGTTCCTCACGC-  
TAA-GTG-ATCGAA-AGA-GTAAG-ATGAGAAAG-GTTTCTCACGC-  
TTT-GTG-ATC-TAAAGA-GTAAG-ATGAGTTAGCGTTCCTCACGC-  
TGA-GTG-AGC-GAAAGA-GTAAG-ATGAGAGAGTGTTCCTCACGC-  
TAA-GTG-ATC-GAAAGCA-TAAG-ATGAGAAAGTGTTCCTCACGC-  
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TGTTGTGGAGC-GAAAGAA-TAAG-ATGAGAAATGTGTTCCTCGGC-  
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TCA-GTG-ATC-GAAAGA-GTAAG-ATGAGAGAGTGTTCCTCACGC-  
NAA-GGNA--T-CAAANA-GNAAG-GG-AGAAANAGTTTCTCCTCC-C-  
TGA-GTAAAT-GAAGGA-GTAAC-AACAGCAGCGGTGTCCACAAGC-  
TGA-GTG-ATC-GAAA-ACGTAAG-ATGAGAAAGAGTTTCTCACCT-  
TGA-GTG-ATC-GAAAGA-GTAAG-ATGAGAAAGTGTTCCTCACGC-  
TAA-GTA-ATC-GATAAGTACTA-A-GAGAAAGTGTTCCTCACGC-  
TAA-GGG-ATC-GAAAGATGTAAG-G-GAGAAAGTGTTCCTCACGC-  
TCA-TTG-ATC-GAAATA-GTAAG-AGGAGACCGAGTTTCTCACGC-

Miscanthuscapensis6584  
Heteropogoncontortus6538  
Hyparrheniaanamesa4807  
Pdilatatum6617  
Zea  
Icylindrical  
Sacharumofficinatum  
Sorghumhalepense  
Ttriandra  
Pschinzii

TCA-GTGGATC-GAAATAGATAGG-CGCAGACCGAGTTTCCTCACCC-  
TAA-GTG-ATC-GAAAGAGGTAT---TGAGAAAGTGTCTCCTCGCGC-  
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AAAGCCCATTATAAATGGGCTTAA--GAGTCACTAGCTTTATCATTCT  
TTG-ATTT-TT-TAGTCC-CTTTAA--TTGAAT-AG-----ATGC-  
TTT-TCT-TTT-ATCAAT-GGG---TTT----AAGA--TTC-ACTAGC-  
TTT-TCT-TTT-ATCAAT-GGG---TTT----AAGA--TTC-ACTAGC-  
TTT-TCT-TTT-ATCAAT-GGG---TTT----AAGA--TTC-ACTAGC-  
TTTT-TCT-TTT-ATCAAT-GGG---TTT----AAGA--TTC-ACTAGC-  
??

Diheteropogonfillifolius5872  
Hyparrheniadissoluta5145a  
Hyparrheniafillipendula5144  
Sorghumhalepense9318  
Hyparrheniapilgeriana4738  
Trachypogonspicatus4846  
Phacelurusfranksae4705  
Hyparrheniaquarrei6586  
Andropogonappendiculatus6692  
Elionurusmuticus6559  
Elionurusmuticus6539  
Elionurusmuticus6954  
Hyparrheniahirta6656f  
Themedatriandra6383  
Themedatriandra6555  
Themedatriandra6525  
Andropogonschirensis6581  
Imperatacylindrica5114a  
Bothriochloainsculpta5681  
Bothriochloa7489  
Heteropogoncontortus6628  
Hyparrheniapilgeriana4603  
Hemarthriaaltissima5698  
Andropogonamethystinus6679  
Andropogonamethystinus6685  
Andropogondistachyos6684  
Hyparrheniatamba6585  
Hyparrheniarudis6576  
Hyparrheniatamba6537a  
Cymbopogonexcavatus6577  
Cymbopogonsp5800  
Andropogonappendiculatus6677  
Heteropogoncontortus6942  
Andropogoneucomus6587  
Cymbopogondieterlenii6642  
Cymbopogonprolixus6682  
Imperatacylindrica6632a  
Trachypogonspicatus6580  
Cymbopogonplurinodeus6406  
Hyparrheniahirta6618  
Hemarthria9319  
Sorghumbicolour6588  
Miscanthuscapensis6655  
Miscanthuscapensis6584  
Heteropogoncontortus6538  
Hyparrheniaanamesa4807  
Pdilatatum6617  
Zea  
Icylindrical  
Sacharumofficinatum  
Sorghumhalepense  
Ttriandra  
Pscinzi

---CATCAATA-GGAA-AATAGTTACCCAAATCTAAG-TAA---TCGAA  
---CATCAATA-GGAA-AATAGTTACCCAAATCTAAG-TAA---TCCAA  
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TCCAAAAAATA-AAAAAATACCTACCCAAATCTACT-TCT---TCGAA  
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TAGG--AAAAA-AAGAAAATAGTTACCCAAAGTCTAAG-TAA---TCGAA  
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-TGTTCT-TGA-GTTACTTA-GACTAC-GATAA-GTAATTTAT-TTGAGCT  
-TGTTCT-TGGAGTTACTTA-GAGTAC-GATAA-GTAATTTAT-CTGCTAA  
-TGTTCT-TGAGTTACTTA-GAATAC-GATAA-GTAAGTTAT-CTACTAC  
-TGTTCT-TGAGTTACCGA-GAATAC-GATGA-GTGAGTTAT-CTACGAA  
-TTCTCACGGCGTTACT-A-GAATCCCGATAACGTAATTTCTTCTCTTAA  
-TTCTCT-GGCGTTACTTA-CGAATCCGATAA-GTAATTTAT-CTTCTAA  
-TTCTCT-TGAGTTACCTA-GAAT-----A-GGATCTTAT---ATAA  
-GGCTCT-GGGGTTACTTA-GAATAC-GATGA-GTAATTTAT-CTACTAA  
-TGTTCTG-TGAGTTACTCA-GAATAC-GATCA-GTAATTTAT-CTACTAA  
-TGTTCT-TGAGTTACTCA-GAATAC-GATAA-GTAGTTTAT-CTACTAA  
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-TGTTCT-TGAGTTACTTA-GAATAC-GATAA-GTAATTTAT-CTACTAA  
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-TGTTCTTGA-GTTTCTTA-GAATAC-TATG--CGTAGTCTTCTACTAG  
-TGTTCTATA-GTTTCTTA-GACTGC-GCTC--CGTAGTTTCT-CTACTAA  
-TGTTCTATA-ACTTTTAC-CACTGG-ACTC--CTGAGTTAT-CG-CTGA  
-TGTTCTAGAAATGACTGA-GAATAC-GATCG-CCAAGGTAT-GTACGAA  
-TGTTCTTGA-GTTTCTTA-GAATAC-GATCA-GTAATTTAT-CTACTAA  
-TGTTCTTGA-GTTTCTTA-GAATAC-GATAA-GTAAGTTAT-CTACTAA  
-TGNNCT-TCAGTTTCTTA-GAATCC-GATCA-GTTATTTAT-CTACTAA  
-CGTTCC-CGAGTTTCTTTC-GTATCC-CCTCA-GTCATATAA-TTTCNAA  
-GTT-CT-TGAGTTCTGATGGATAC-GATAA-GTTGTTTAT-CTACTAA  
-TGT-CT-TGAGTTGCTGA-GAATAC-GATAA-GTAATTTAT-CTACTAA  
-CCT-CT-GCGGTTTTTGA-GAATAC-GACAA-CTAATTTAT-CTACTAG  
-CGTTCT-TGGGTTTCTTA-GAATAC-GATAA-GTAATTTAT-CTACTAG  
-TGTTCT-TGCGTTACTTA-GAATACGA-TAG---TATTTTACTAGTCA  
-CGTTCTG-TGCGTTACTTA-TCATACGACTAA---GGTCTGACCGGCTCG  
-AGTTCT-CGCGTTGCTTA-GAATAC-GATAA-GGACTTTATCGTCCAAG  
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CTAT----GGAGCAAAGGA-GAGTGACGAGAACTCAATGTATCTTATGCT  
-AAATA----CATTTACTAA-GAGATG-CACAA-GAAAGGG---TCAGGAT  
--TTTC-TCA--TT-CT-AC-TCTTT-CACAAAGGAGTGCGACAAGAAGT  
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??

Diheteropogonfillifolius5872  
Hyparrheniadissoluta5145a  
Hyparrheniafillipendula5144

AG-----AGTAAGATG-----AGAAAGTGTTC  
AG-----AGTAAATG-----AGAAAGTGTTC  
AT-----AAAAAACCTA--GAATAAGTAATTC

*Sorghumhalepense*9318  
*Hyparrheniampilgeriana*4738  
*Trachypogonspicatus*4846  
*Phacelurusfranksae*4705  
*Hyparrheniaquarrei*6586  
*Andropogonappendiculatus*6692  
*Elionurusmuticus*6559  
*Elionurusmuticus*6539  
*Elionurusmuticus*6954  
*Hyparrheniahirta*6656f  
*Themedatriandra*6383  
*Themedatriandra*6555  
*Themedatriandra*6525  
*Andropogonschirensis*6581  
*Imperatacylindrica*5114a  
*Bothriochloainsculpta*5681  
*Bothriochloa*7489  
*Heteropogoncontortus*6628  
*Hyparrheniampilgeriana*4603  
*Hemarthriaaltissima*5698  
*Andropogonamethestinus*6679  
*Andropogonamethestinus*6685  
*Andropogondistachyos*6684  
*Hyparrheniatamba*6585  
*Hyparrheniarudis*6576  
*Hyparrheniatamba*6537a  
*Cymbopogonexcavatus*6577  
*Cymbopogonssp*5800  
*Andropogonappendiculatus*6677  
*Heteropogoncontortus*6942  
*Andropogoneucomus*6587  
*Cymbopogondieterlenii*6642  
*Cymbopogonprolixus*6682  
*Imperatacylindrica*6632a  
*Trachypogonspicatus*6580  
*Cymbopogonplurinosus*6406  
*Hyparrheniahirta*6618  
*Hemarthria*9319  
*Sorghumbicolour*6588  
*Miscanthuscapensis*6655  
*Miscanthuscapensis*6584  
*Heteropogoncontortus*6538  
*Hyparrheniaanamesa*4807  
*Pdilatum*6617  
*Zea*  
*Icylindrical*  
*Sacharumofficinarum*  
*Sorghumhalepense*  
*Ttriandra*  
*Pschinzii*

AG-----AGCCAGATGAG--CAATAAATCGTTTTCC  
AA-----TGAAGAAAACCT--AGAATAGGGTAAC  
AG-----AGTAAATCCG--AGAGTAAGGGAAGCC  
AG-----AGTAAGA-----AGAGAAAGGGTATCC  
AG-----AGTATGTTG-----AGACAGGGTTTTCC  
GA-----AACAAATAT-----AATCCCATCTCTTAG  
GA-----AAAAATAT-----AATCTCATCTCTTAG  
GA-----AACAAATAT-----AATCTCATCTCTTAG  
GA-----AACAAATAT-----AATCCCATCACTTAG  
AGCCTAGGCT-----CGAGAGAAGAAATCTA--GAATCTCGTCTCTA--  
ACTCAAGGCC-----CGATTGAGGAAACCTA--GAATCTCGTCTCTA--  
ACTCAAGGCC-----CGATTGAGGAAACCTA--GAATCTCGTCTCGA--  
ACTCAAGACC-----CGATTGAG--AAACCTA--GAATCTCGTCTCG--  
AGCCAAGGTC-----TGGATGAAGAAATCTA--TAATCTCGTCTCAA--  
AGG-----AATGATGAAGCCTA--TAATCTCATCTCTCT--  
TGT-----AATGATAAAGATTA--TAATCTCATCTCAA--  
AA-----AACCAATT-----AATCTCAT-----  
AG-----AACAAAG-----AATCTCAT-----  
AG-----AAAAAT-----AATCTCATAGCC--  
AG-----AAGAAATAATC--TAACCCACATAGC--  
AGTAAACT-----AAAGAAAAAAATCAA--TAATCTCATCTCAT--  
AGATATTTCT-----ACAGAAAAAAATCTA--TAATCTCATCTCAT--  
AGCTTATCT-----ACAGAAAAAAATCTA--TAATCTCATCTCAT--  
AGT-----AAAAATAAAA--TAATCTCGTCTCAT--  
AGT-----GAAGAAATAAAA--TAATCGCGTCTCA--  
AGA-----AAAAATAAAA--TAATCTCATCTCAT--  
AT-----ACCTTAATCTA--TAGTCTCAA-----  
AG-----ACAT--AGTATA--TAGTCTCAA-----  
AC-----AGACAAGACCA--TAGTCTCATA-----  
AC-----AGGCGAGCACA--AGCTTCGTA-----  
GG-----AATCTAGCATA--AGGCTCATA-----  
AG-----ACAT--AATATA--TAGTCTCATA-----  
AG-----ACTT--AATCTA--TAGTCTCATA-----  
ATAAA-----GCATGAAGGAAAAAACCTA--CTATATAGTCTCAT--  
AGGAATTATCCACGGCTTAAATAAAAAAATCAAGCAATCTCATCTCAA--  
ACTAAGGCT-----CGAGAGAAGAAACCTA--CAATCTCGTCTCTG--  
GGTTAAGCT-----CTAGAGAAGAAACTAA--GAATCTCGTCTCTA--  
AGAGCTTATA---G--ACGAAAGAAAAAACAAA--TAATCTCGTCTCAA--  
AGAGCCTATA---G--AGGAAAGAAAAATCAA--TAATCTCGTCTCTT--  
ACCTAAGATC-----TAGTGAAGAAATCCA--GAATCTCGTCTCTT--  
ACCTACCCT-----TATTGAAGAAATCCA--GCATCTCATGTCTCT--  
AAGCACTCAAGAA--ACATAAATGATAAAGTTCGCTAAAACATGTAGGATGA  
GGTTACAGACGATACGATGTTGCTTAGCTTCGCTAAAACACTTAGCA  
AC-----ACACGGAA-----AGATGGTATTTT--  
ACTCATGGTA-----GAGCAGAGGAC--TGAA--AATCCTCGTGTCTC--  
A--TGAATCTTATGCTATTTCATTAAATA-----GATGATTTCTTTTTA--  
A--TGAATCTTATGCTATTTCATTAAATA-----GATGATTTCTTTTTA--  
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A--TGAATCTTATGCTATTTCATTAAATA-----GATGATTTCTTTTTA--  
A--TGAATCTTATGCTATTTCATTAAATA-----GATGATTTCTTTTTA--  
A--TGAATCTTATGCTATTTCATTAAATA-----GATGATTTCTTTTTA--  
??

*Diheteropogonfillifolius*5872  
*Hyparrheniadissoluta*5145a  
*Hyparrheniafillipendula*5144  
*Sorghumhalepense*9318  
*Hyparrheniampilgeriana*4738  
*Trachypogonspicatus*4846  
*Phacelurusfranksae*4705  
*Hyparrheniaquarrei*6586  
*Andropogonappendiculatus*6692  
*Elionurusmuticus*6559  
*Elionurusmuticus*6539  
*Elionurusmuticus*6954  
*Hyparrheniahirta*6656f  
*Themedatriandra*6383  
*Themedatriandra*6555  
*Themedatriandra*6525  
*Andropogonschirensis*6581  
*Imperatacylindrica*5114a  
*Bothriochloainsculpta*5681

TTACGCTGCTTTTGAGTTACTTAGAA--TACGATAAGTAAT--TTAT  
TCACCCTTCTCTTGAGTTCTTAAAA--TACGATAAGTAAT--TTAT  
TCTAACTTCTCTAATATCACTTAC--A--TTCATTAGGGATAATTAT  
TCACGCTGCTCTTTTTTCCCTTAGTA--TACGATAAGCATT--TTAT  
TGTTTACTCGTATAATATCATATCCG--TTCATTAGGCATA--ATCT  
TCTCGCCGCGCATGAGTTCCTAACCG--TTCATTAGGTATT--TTAT  
TCACGCGGCGCATGAGTACTTAGCA--TACGATAAGTAAT--ATAT  
TCACACTGCTCTCGTGTGCTTAGAA--ATCGATACGTTAT--TTGC  
CCGTTCCCTTAGAGATAAATAAAGC--TAATAAATTCATAATAC  
CCGTTCCCTTAGAGATAAATAAAGC--TACTAAACTCATAATA  
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AGCCTTCCCGTATAG--CTACTTATCC--GTCCATTAAGATTCTATA  
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*Bothriochloa*7489  
*Heteropogoncontortus*6628  
*Hyparrheniapilgeriana*4603  
*Hemarthriaaltissima*5698  
*Andropogonamethystinus*6679  
*Andropogonamethystinus*6685  
*Andropogondistachyos*6684  
*Hyparrheniatamba*6585  
*Hyparrheniarudis*6576  
*Hyparrheniatamba*6537a  
*Cymbopogonexcavatus*6577  
*Cymbopogonsp5800*  
*Andropogonappendiculatus*6677  
*Heteropogoncontortus*6942  
*Andropogoneucomus*6587  
*Cymbopogondieterlenii*6642  
*Cymbopogonprolixus*6682  
*Imperatacylindrica*6632a  
*Trachypogonspicatus*6580  
*Cymbopogonplurinosus*6406  
*Hyparrheniahirta*6618  
*Hemarthria*9319  
*Sorghumbicolour*6588  
*Miscanthuscapensis*6655  
*Miscanthuscapensis*6584  
*Heteropogoncontortus*6538  
*Hyparrheniaanamesa*4807  
*Pdilatum*6617  
*Zea*  
*Icylindrical*  
*Sacharumofficinarium*  
*Sorghumhalepense*  
*Ttriantra*  
*Pschinzii*

---CTTTATGTAGAC-CTGTAAATCA-ATTTATTANTCTTCATAA  
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---CGTTCCTAAGAT-CTAATAATTAAGCTCATAATTCATAATAA  
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??

*Diheteropogonfillifolius*5872  
*Hyparrheniadissoluta*5145a  
*Hyparrheniafillipendula*5144  
*Sorghumhalepense*9318  
*Hyparrheniapilgeriana*4738  
*Trachypogonspicatus*4846  
*Phacelurusfranksae*4705  
*Hyparrheniaquarrei*6586  
*Andropogonappendiculatus*6692  
*Elionurusmuticus*6559  
*Elionurusmuticus*6539  
*Elionurusmuticus*6954  
*Hyparrheniahirta*6656f  
*Themedatriandra*6383  
*Themedatriandra*6555  
*Themedatriandra*6525  
*Andropogonschirensis*6581  
*Imperatacylindrica*5114a  
*Bothriochloainsculpta*5681  
*Bothriochloa*7489  
*Heteropogoncontortus*6628  
*Hyparrheniapilgeriana*4603  
*Hemarthriaaltissima*5698  
*Andropogonamethystinus*6679  
*Andropogonamethystinus*6685  
*Andropogondistachyos*6684  
*Hyparrheniatamba*6585  
*Hyparrheniarudis*6576  
*Hyparrheniatamba*6537a  
*Cymbopogonexcavatus*6577  
*Cymbopogonsp5800*  
*Andropogonappendiculatus*6677  
*Heteropogoncontortus*6942  
*Andropogoneucomus*6587  
*Cymbopogondieterlenii*6642

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TTCATTC-AGTAAGTGT-----AGGTATAGTAATAG-GACGG  
CTCCTAC-AGTAAGAGTT-----AGCTATCCGTAGATG-GACGG  
CTACTAC-AGAAAATGTT-----AACTATCCTAATGA-GGCGG  
CTACTA-TAGAAA-----TAAGTACTCCTCATCG--CCG  
TTCATTC-GGTAGGTGTT-----ACGTATCCTGATGC-GAGGG  
TTCATGC-GGTAGGTGTT-----ACCTATCCTGATGC-GAGGG  
TTCATTC-CGTAGCTGTT-----ACCTATCCTGATGC-GAGGA  
CTCATTC-GGTAGGTGTT-----ACCTATCCTGATGC-GAGGA  
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TTCATTC-A-GCAATTGTT---GTTTCGTGTCCTCAGCGGAGGG  
TTCATTC-A-GCAATTGTT---GTTTCGTGTCCTCAGCGGAGGG  
TT-ATTC-A-GCAATTGTT---GTTTCGTATCCTCAGCGGAGGG  
TTATTCAT-GCAATAGGT---GTTTCGTATCCTCAGCGGAGGG  
TTCATCAA-TCAATAGGT---GTTTCGTATCCTAAGCGGAGGG  
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TAATTCATTGCAAAACGGT---GTTCCGTATCCTGAAGGGGAGGC  
TACATCAT-GCGTACGGT---GTTACGTATCCTGAGGGGAGGC  
TTCTTTAA-TCAATCGTT---GTTTCGTATCCTAAGCGGAGGG  
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TAAATCAA-TCAATAGTT---GTTTCGTATCCTGAAGCGGAGGG  
TTAATGCAACCTATC-GT---GTTTCATTTACTCAAGCCCATCC  
TTCTTTAAGCGGTAG-TT---GTTTCGTATCCTAAGCGGATCC  
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TAACAA-ACGGGGTGCGT---GTTTCGTACTCTTAGGCTGATCC  
TTACTTTACCCTATCGGT---GTTTCATACACTGAGGACGAGCC  
TAATTC-AGGCGGTAGGT---GTTTCGTATCCTGAGGGGATGG

*Cymbopogonprolixus*6682  
*Imperatacylindrica*6632a  
*Trachypogonspicatus*6580  
*Cymbopogonplurinodus*6406  
*Hyparrheniahirta*6618  
*Hemarthria*9319  
*Sorghumbicolour*6588  
*Miscanthuscapensis*6655  
*Miscanthuscapensis*6584  
*Heteropogoncontortus*6538  
*Hyparrheniaanamesa*4807  
*Pdilatum*6617  
*Zea*  
*Icylindrical*  
*Sacharumofficinarium*  
*Sorghumhalepense*  
*Ttriandra*  
*Pschinzii*

*Diheteropogonfillifolius*5872  
*Hyparrheniadissoluta*5145a  
*Hyparrheniafillipendula*5144  
*Sorghumhalepense*9318  
*Hyparrheniapilgeriana*4738  
*Trachypogonspicatus*4846  
*Phacelurusfranksae*4705  
*Hyparrheniaquarrei*6586  
*Andropogonappendiculatus*6692  
*Elionurusmuticus*6559  
*Elionurusmuticus*6539  
*Elionurusmuticus*6954  
*Hyparrheniahirta*6656f  
*Themedatriandra*6383  
*Themedatriandra*6555  
*Themedatriandra*6525  
*Andropogonschirensis*6581  
*Imperatacylindrica*5114a  
*Bothriochloainsculpta*5681  
*Bothriochloa*7489  
*Heteropogoncontortus*6628  
*Hyparrheniapilgeriana*4603  
*Hemarthriaaltissima*5698  
*Andropogonamethystinus*6679  
*Andropogonamethystinus*6685  
*Andropogondistachyos*6684  
*Hyparrheniatamba*6585  
*Hyparrheniarudis*6576  
*Hyparrheniatamba*6537a  
*Cymbopogonexcavatus*6577  
*Cymbopogonsp*5800  
*Andropogonappendiculatus*6677  
*Heteropogoncontortus*6942  
*Andropogoneucomus*6587  
*Cymbopogondieterlenii*6642  
*Cymbopogonprolixus*6682  
*Imperatacylindrica*6632a  
*Trachypogonspicatus*6580  
*Cymbopogonplurinodus*6406  
*Hyparrheniahirta*6618  
*Hemarthria*9319  
*Sorghumbicolour*6588  
*Miscanthuscapensis*6655  
*Miscanthuscapensis*6584  
*Heteropogoncontortus*6538  
*Hyparrheniaanamesa*4807  
*Pdilatum*6617  
*Zea*  
*Icylindrical*  
*Sacharumofficinarium*  
*Sorghumhalepense*  
*Ttriandra*

TTATTC-ATGCAGTAGGT---GTTTCGTATACTGAGGCGGACGG  
T-TATTCATCTATCGTT---GTTTCGTATACTCAAGCCGAGCG  
T-TAATTCACC-ATCGTT---GGTTCGTATCCCTCAAGCGGAGCG  
TTAAAGTAACCAGTCTGTT---GTTTCGTATACTCAAGCCGAGGG  
T-TATTTATTCAATGAGTT---GTTTCGTATACTCAAGGGGAGGG  
T-AATTCACTCAATGTT---GTTTCGTATCCCTGAGGCGGAGCA  
T-ACTTCATTCGGTAGGT---GTTTCGTATCCCTGATGGGAGGG  
T-AATTTAAGCAGTAGTT---GTTTGGTATCCCTCAAGCGGAG-C  
T-ATTCGAGGCGGAGTG---CTTCCCATCCCTGCCGTGCGAC  
TGGATATATCTATTCGTGAGTCTGTAGTAGTACGTAGCCCATGG  
AATATTCACCTCAGTAAGT---GTCATGTCCAGTAAGG-GAAG  
TTAAGATAGGAAA-----GTAATGTATAATAAAGA-GAGAG  
TGAATAGA-TCAATTGAT---TACAAATATTTTGAGATGGAGGG  
T--CTCGATTATTAATTCGATTTAAGTATTATTAA-GTAAGCC  
T--CTCGATTATTAATTCGATTTAAGTATTATTAA-GTAAGCC  
T--CTCGATTATTAATTCGATTTAAGTATTATTAA-GTAAGCC  
??

-GTAAATG-GTTTAAA--CCTTA----TGA-AAT-ACTAAAA--AATT  
GGTAAAGG-GTTTAAA--CCTTACCTTATGA-AAT-ACTAAAAAT-AATC  
--TATAAG-ATTCAGA--CCTTACCCTTATA-AAT-AAATAAAA--AATC  
---ATACA-GTTCAA--CCCTACCTTTAGA-AAT-ACTAAAA--AATC  
--TGTAT--TTTCAGG--CCATACCTTTAGG-AAT-ACTAAAA--AATC  
--TGTAAG-GTTCACG--TCTTACCTTTTGG-AAT-AATTA AAA--AATC  
--GTAAAG-GTTCAAA--TCTTACCTT-AGA-AAT-ACTAAAA--AATC  
-----TTC-CTTAGAG--CCTTT----TATT-AAT-G-CTAAAA--CTTC  
-GTAAAGG-ATTA AAA--CCTTTCCTTACCGA-AAT-ACTAAAA--AATC  
-GTATAGG-AGTAAAA--CTTATCTTAC-GA-AAT-ACTAAAA--AATC  
-GTAAAGT-ATTA AAA--CTTATCTTAC-GA-AAT-ACTAAAA--AATC  
-GTATAGG-ATTA AAA--CTTTTCCTTAC-GA-AAT-ACTAAAA--AATC  
-GTAAACA-ATTA AAA--CCTTATCTTATGA-AAT-ACTAATCAG--TC  
-GTAAACA-AGTAAAC--CTCATCTTATGA-AAT-ACTAATCAG--TC  
-GTAAACA-AGTAAAC--CTCATCTTATGA-AAT-ACTAATCAG--TC  
-GTAAACA-AGTAAAC--CTCATCTTATGA-AAT-ACTAATCAG--TC  
-GTAAACA-ATTA AAA--CCTTATCTTACGA-ACT-ACTAATAAAA--TC  
-GTAAACA-ATTA AAA--CTTACCTTATGA-AAT-ACTAATCAA--GC  
-GGAAAGG-ATTA AAA--CTCACCTTATGA-AAT-ACTAATAAAA--GC  
-CTCAAT-ATTCATC---TT---CTTCCCTTTT-ACCTAACA--T-  
-GGTAAAG-ATTTAAC-GGCTTACCCTCTGT-AAT-AAATAAAA--TC  
-GTTAAAT-CTTAAAC-TTTATGAAATAAAT---T-ACTAATAAAA--TC  
-TTTAAAA-CTTACGCGTTAAATTTCTAACG-TAT-AAATAAAA--TC  
-GTAAAAG-A-TTAAA--ACTTACCTTATGA--AT-ACTAATAACA--TC  
-GTAAAAG-A-TTACA--TCTTACCTTATCA-CAT-TACTAATAAAA--TC  
-GTAAAAG-AATTA AAA--ACTTACCTTATGA-AAT-AAATAACA--TC  
-GTACAGA-ATTCAAA--CCTTATCTTATGA-AAT-ACTAACAAT--GC  
-GGAAACA-ATTA AAA--CCTTATCTTACGA-AGT-ACTAACAAT-AGC  
-GTAAAAGGATTA AAA--CCTTATCTTACGA-AAT-ACTAATAAAA--TC  
-TTAAACCAATTCAAA--CTCATCTTACGC-AATTACTAACAAT-AGC  
-GTAAAC-AATTA AAA--CCTTATCTTANGC-ANT-ACTAANCCN-AGC  
-GTAAAA--ATATAAC--CCTTATCTTATGC-CAT-ACCTCAACA--AGG  
-TTACTC-TATCTCGC---CCTTATCTTCCGA-ATT-GTCTAATAA--AGG  
-GTAAAG-CATCTTAA--CCTTATCTTGTGA-AAT-TCCTAATCA--GGC  
-GTAAAG-GATTA AAA--CCTTACCTTACGA-AAT-ACTAATAACA--ATC  
-GGAAAC-AATTA AAA--CTCACCTTACGA-AAT-ACTAATAACA--ATC  
-GTAAACA-ATTC AAC--CTCATCTTACGA-AAT-ACTAATAACA--ATC  
-GGAAACA-ATTA AAA--CTCATCTTACGA-AAT-ACTAATAACA--ATC  
-GTAAACA-ATTA AAA--CCTTATCTTACGC-AAT-ACTCAAAAATAGCA  
-GTAAAGA-ATTA AAA--CCTTATCTTATGA-AAT-ACTAATAACA--ATC  
-GTATACG-ATTA AAA--CCTTATCTTATGA-AAT-ACTAATAACA--ATC  
-GTAAAGG-ATTA AAA--CCTTACCTTATGA-AAT-ACTAATAACA--ATC  
-TTATACA-ATTC AAA--CTCATCTTACGCAT-CACTACACAA--GC  
-TTATAGA-AGTTAAC--CTCTTCTTCACT-TAT-AACGACTAAG--TC  
-GTAGGCTAATCGAT---CTCTTCTGACGA-AAT-CACTAATCAG--GC  
--TAAAGG-ATTCAGG---CCTTAACTCTTATGTAT-AAATAAAA--ATC  
-TTATTCAAAGTATAG-TCATTCCTTAC---ATTTCTAAGTTCGAAT  
-GTAAATA-TTTAAAA--ATCTACTTATGTC-TTT-AGCTAGAGT--AA  
-ATCCACA-ATGCATAGGACT-ACCCC-TCCCCATTTCTAATTTGGAAT  
-ATCCACA-ATGCATAGGACT-ACCCC-TCCCCATTTCTAATTTGGAAT  
-ATCCACA-ATGCATAGGACT-ACCCC-TCCCCATTTCTAATTTGGAAT

Pschinzii

??

*Diheteropogonfillifolius*5872  
*Hyparrheniadissoluta*5145a  
*Hyparrheniafillipendula*5144  
*Sorghumhalepense*9318  
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*Cymbopogonsp*5800  
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*Andropogoneucomus*6587  
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*Miscanthuscapensis*6655  
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*Heteropogoncontortus*6538  
*Hyparrheniaanamesa*4807  
*Pdilatum*6617  
*Zea*  
*Icylindrical*  
*Sacharumofficinatum*  
*Sorghumhalepense*  
*Ttriandra*  
*Pschinzii*

AGGGA-TGTTTAAACGGTCT--CTTCGTCT-ACGATTGGCAC-CTTTTGCG  
ACCCA-AATTTACGGTCT--TTATTCCC-ACCGTTTGCCA-CTTTTGTA  
AGCGA-AATTGACCTAT----CTTTTCCC-ACCATGTGCCA-CTTTTGTC  
AGGGA-AATGTAAGTATATTT-CTTTGCC-ATAAAGAGGAC-CTACTAC  
CGGGA-AATT-AACTGTAT--CTACGTTT-ATGAGATGATC-CTACTG-C  
AGGGA-AATT-AACGGTTT--CTTCGTTT-ATGAGATGATC-CTACTG-C  
AGGGA-AATT-AGCGGTAT--CTACGTTT-ATGAGAGGACC-CTACTG-C  
AGGGA-AATT-AACTGTAT--CTACGTTT-ATGAGATGATC-CTACTA-C  
AGGGA-AATT-GACTGTAT--GTTTCGTTT-ACGAAAGGATACCTACTG-C  
AGGGA-AATT-TACTGTAT--GTTTCGTTT-CCGAGATGACAACCTACTA-C  
AGGGC-AATT-ATCTGTAT--GTTTCGTTT-ACCAAAGGATAACTACTA-C  
AGGGA-AACT-GACTGTAT--GTACGTTT-ATGAGATGATC-CTACTA-C  
AGGGA-AACTGTCT-CTAT--CTTCGTTTACCAGGATGATC-CTACTA-C  
AGGGA-A-CTGTCT-GTAT--CTTCGTTTAT-GAGATGATC-CTACTA-C  
AGGGA-A-CTGTCT-GTAT--CTTCGTTTAT-GAGATGATC-CTACTA-C  
AGGGA-A-CTGTCT-GTAT--CTTCGTTTAT-GAGATGATC-CTACTA-C  
AGGGA-ACTAACT-GTAT--CTTCGTTTAC-CAGATGATA-CTACTA-C  
AGGGA-AATTAACGC-TAT--CTTCGTTTACC-ATAGGACA-CTACTG-C  
AGGGA-GATTGACTG-TAT--CTTCGTTTCAAG-AGATGATA-CTACTA-C  
AGAGG-AGGTAATTA-TAT--CTTTATACTTGT--ATGATT-CTATTA-T  
AGGGA-AATTAAGT-TAT--CTACGTTTATGATGAGGATC-CTACTA-C  
AGGGA-AATTAACGGGTAT--CTAGGTTTATGA-GAGGACA-CTTTTA-C  
AGGGA-AATTAAGTGTAT--CTCTGTATCAGGAGAGGACAACCTACTT-C  
AGGGA-AATTAAGT-TAT--CTTCGTTTACC-AGATGATT-CTACTA-C  
AGGGA-AATTAAGT-TAT--CTTCGTTTACC-AGATGATT-CTACTA-C  
AGGGA-AATTAAGT-TAT--CTTCGTTTACC-AGATGATT-CTACTA-C  
ATGGA-AACTAAGT-TAT--CTTCGTTTACC-AGATGATT-CTACTA-C  
AGGGA-AACTGTCTG-TAT--CTTCGTTTACC-AGATGATT-CTACTA-C  
AGGGA-AATTAAGT-TAT--CTTCGTTTACC-AGATGATT-CTACTA-C  
ACTTGAATTTATCT-CTATT-CTTCGTTTACC-AGATGATT-CTACTA-C  
-CNNGGAATTNACT-CTAT--CTTCNTT-----  
AGCGGGCAGTAGCTTATAT--CTTTGTCC-AC-CGCGG-ACACGTTA-T  
AGGGA-CTTTCGCT-ATA--CCTTGTCT-TCGCGCGG-ATTCNTCTN-N  
AGGGA-ACTGTCT-GTAT--CTTCGTTT-ATGAGATG-ATTCTACTA-C  
AGGGAG-ATTAAGT-GTAT--CTACGTTTACCAGATGGATACTACTA-C  
AGGGAGGATTGACT-GTAT--CTACGTTT-CCGAGATG-ATACTACTA-C  
AGGGA-AATTGA-CTGTAT--CTTCGTTT-CCAAATGATAACTACTA-C  
AGGGA-AATTGA-CTGTAT--CTTCGTTT-ACCAGATGATC-CTACTA-C  
CGGGG-CCTTGG-GCGCATATCTTCTTAG-ACCAATGGACA-CTTTTA-C  
AGGGA-ACTGA-CTGTAT--CTTCGTTT-ACCAGATGATT-CTACTA-C  
AGGGA-AATTGGACTGTAT--CTTCGTTT-ACCAGATGATT-CTACTA-C  
AGGGA-AATTAG-CTGTAT--CTACGTTT-ATGAGAGGATT-CTACTA-C  
AGTTG-GACCTGCC-CTAT--CTTCGTTTATC-TATTGTCC-CTACTA-C  
AGGGG-AACTAAGT-CTAT--ATACGTTTACTG-AGTTGCTC-CTACTA-C  
AGGTT-GGCTATCT-GTAT--CTCCGTTTAT-GACATGATC-CTACTA-C  
AGGGA-AATTAAGT-GTAT--CTACGTTT-ATGAGATGATA-CTACTA-C  
AGAGA--ATGCATATTTATTTTATAGTCCCTTTAATTGACA--TAGAAGC  
GTAGA-AAATCTCT-AGAG-TAAGGATAAATC----TCTAC-ACACTT-C  
-GG-AAACTTTTATTG-ATTTTTTATAGTCCCTTTAATTGACA--TAGATGC  
-GG-AAACTTTTATTG-ATTTTTTATAGTCCCTTTAATTGACA--TAGATGC  
-GG-AAACTTTTATTG-ATTTTTTATAGTCCCTTTAATTGACA--TAGATGC  
??

*Diheteropogonfillifolius*5872  
*Hyparrheniadissoluta*5145a  
*Hyparrheniafillipendula*5144  
*Sorghumhalepense*9318  
*Hyparrheniapilgeriana*4738  
*Trachypogonspicatus*4846  
*Phacelurusfranksae*4705  
*Hyparrheniaquarrei*6586  
*Andropogonappendiculatus*6692  
*Elionurusmuticus*6559  
*Elionurusmuticus*6539  
*Elionurusmuticus*6954  
*Hyparrheniahirta*6656f

GCG-CCTTGGTCA  
GCA-CCTTTGTCA  
GTAACCTATGTCC  
GGGTCCTTTCTCC  
GTGTTCTTTGCCA  
GTGTCCTTTGCCA  
GTGTCCTTTGCCA  
GTGTTCTTTCCCA  
GTGTTCTTTGCCA  
CGGTTCTTTGCCA  
GCGATCTTTGTCA  
GTGTTCTTTGCCC  
GTGTTCTTTGCCA

<i>Themadatriandra6383</i>	GTGTTCTTTCCCA
<i>Themadatriandra6555</i>	GTGTTCTTTGCCA
<i>Themadatriandra6525</i>	GCGTTCTTTGCCA
<i>Andropogonschirensis6581</i>	GTGTTCTTTCCCA
<i>Imperatacylindrica5114a</i>	GCGTCCTTTGCCA
<i>Bothriochloainsculpta5681</i>	GCGTTCTTTGTCA
<i>Bothriochloa7489</i>	ATGATCTTTGTCA
<i>Heteropogoncontortus6628</i>	GCGTTCTTTGCCA
<i>Hyparrheniapilgeriana4603</i>	GCGTTCTTTGCCN
<i>Hemarthriaaltissima5698</i>	GTGACTTTTGTCA
<i>Andropogonamethystinus6679</i>	GCGTCCTTTTCCA
<i>Andropogonamethystinus6685</i>	GCGACCTTTGTCA
<i>Andropogondistachyos6684</i>	GTGTCCTTTGCCA
<i>Hyparrheniatamba6585</i>	GTGTTCTTTGTCA
<i>Hyparrheniarudis6576</i>	GTGTCCTTTGTCA
<i>Hyparrheniatamba6537a</i>	GCGATCTTTGTCA
<i>Cymbopogonexcavatus6577</i>	GCGTTCTTTCCCA
<i>Cymbopogonsp5800</i>	-----
<i>Andropogonappendiculatus6677</i>	GTGTCCTTTTCCA
<i>Heteropogoncontortus6942</i>	GTGTTCTTTCCCC
<i>Andropogoneucomus6587</i>	GTGTTCTTTCCCA
<i>Cymbopogondieterlenii6642</i>	GCGTTCTTTCCCA
<i>Cymbopogonprolixus6682</i>	G-GTTCTTTCCCA
<i>Imperatacylindrica6632a</i>	GCGACCTTTGTCA
<i>Trachypogonspicatus6580</i>	GGGACCTTTTCCA
<i>Cymbopogonplurinodus6406</i>	CGGACCTTGGTCA
<i>Hyparrheniahirta6618</i>	GTGACCTTTCTCA
<i>Hemarthria9319</i>	GTGTTCTTTCTCA
<i>Sorghumbicolour6588</i>	GTGTTCTTTCCCA
<i>Miscanthuscapensis6655</i>	G-GTTCTTTCCCC
<i>Miscanthuscapensis6584</i>	G-TGTCTTTCCCA
<i>Heteropogoncontortus6538</i>	GGGTTCTTTGCCA
<i>Hyparrheniaanamesa4807</i>	GTGTCCTTGCCCA
<i>Pdilatum6617</i>	AGTACTCTACTAA
<i>Zea</i>	TTTTTCTTTTAAA
<i>Icylindrical</i>	AAATACTTTCTAA
<i>Sacharumofficinarum</i>	AAATACTTTCTAA
<i>Sorghumhalepense</i>	AAATACTTTCTAA
<i>Ttriandra</i>	AAATACTTTCTAA
<i>P.schinzii</i>	??????????????

**Appendix D:** CLUSTAL W (1.74) multiple sequence alignment data of the Andropogoneae specimens for *tnrL-F* region.

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Aadscen6713      GAA-TCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Aameth6679      ?????????????????????????????????????????????????
Aameth6685      ?????????????????????????????????????????????????
Aappen6543      ----TCGG-TATA-GGTTTTGTGG-----TTTGGGGG-----
Aappen6677      GAA-TCGG-TAG--CGCTACGTT-GTATTCCCG--CCTTCCCAGGA-ACC
Aappen6946      GAA-TGGA-AACCTTGCTAGGTTCTAACTCCCN--AAATCCAGAGCCCC
Adist6684      ?????????????????????????????????????????????????
Aeuco5869      ?????????????????????????????????????????????????
Aschiren6581    GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Bbladh          ?????????????????????????????????????????????????
Binsc5681      GAA-TCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Brachiaria     GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Bserra5074     GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Bserra6654     GAA-TCGGATAGA-CGATACGGT-AATG--GGT--ATATCCCAGCCTGAA
Bserra6954     ?????????????????????????????????????????????????
Cdieter6642    GAAATCGG-TACA-CGCTACGTT-CAAGTCCCT--CTATCCCCACCGAAA
Cexc5877      -----
Cmarg4397     GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Cplur6406     ?????????????????????????????????????????????????
Cplur6406     GAA-TCGG-ACC--GCTACGTTGGAAGTCCCA--ATTCAGAGAACC
Dfilli5872    GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Hanam4841     GAA-TCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Hcon6942     ?????????????????????????????????????????????????
Hdiss5145     GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Sbicolour9319 ?????????????????????????????????????????????????
Hemarthriaaltissima5698 GAA-TGGA-AACCTTGCTAAGTG-TAACTTCCC--AATTCGGGAACCCT
Hfilli5144    ?????????????????????????????????????????????????
Hhirt6550     GAA-TCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCACCAAACC
Hhirt6656     ?????????????????????????????????????????????????
Hlongi6560    GAA-TCGG-TAGA-CGCTACGTT-AAGT--CCC--TCCTCCCAGCCGAAA
Hpilg4603     GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Hpilg4738     GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Hquar5088     ?????????????????????????????????????????????????
Hrud6576      ?????????????????????????????????????????????????
Htamb6585     ?????????????????????????????????????????????????
Icylin5114    GAAATCGG-TAGA-CGCTACGTT-CAATTCCT--CCTTCCCAGCCGAAA
Icylin6632    GAA-TCGG-TAGA-CGCTACGTT--TATTGCCT--CCTTCCCAGAGAACC
Mcape6655     GTA-TGGG-AAC--TGCTAAGTG-TAACTCCCA--ATTCCCAGAAC--
Pdilatatum6617 GAA-TCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Pfrank4705    GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Pschinz6652   GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA
Sbic6589      GAA-T-GG-T-GA-CGCTACGTT--ACTCCCT--CCTTCCCAGCCGAAA
Shalep9318    GAA-TCGG-AACC-CGCTACGTT-AAGT--CCC--ATTC--AGC-GAAA
Snigros6567   GAAATCGG-TAGA-CGCTACGTT-TAATTCCT--CATTCAGCCGAAA
Tspic4846     GAAATCGG-TAGA-CGCTACGTT-CAATTCCT--CCTTCCCAGCCGAAA
Tspic6580     GAA-TGG--AAC--TGCTACGTT-CAAGTGCC--TCCTAGGGGAAA
Ttria6383 b   ?????????????????????????????????????????????????
Ttria6555     GAAAT-GG-TAG--CGGTAGCGTTCAAGTCCCCGCATAAAGTGCCACCT
Zea           TTTCT-AT-CTA--AGTGGAACTCCAATTT-AGAAGTAGT-TAATAACT
Elionurusmuticus6954 GAA-TCGGATAGA-CGATACGGT-AATG--GGT--ATATCCCAGCCTGAA
Icylindrical  -----TTCCA--AATTCAGAGAA-ACC
Sacharumofficinarium -----TTCCA--AATTCAGAGAA-ACC
Sorghumhalepense -----TTCCA--AATTCAGAGAA-ACC
Ttriaandra    -----TTCCA--AATTCAGAGAA-ACC
Pschinzii     ATG-GAAA-CCT--GCTAAGTGG-TAACTTCCA--AATTCAGAGAA-ACC

Aadscen6713    -TGGGT-GGATGCAATGGGCAA-TCCTGAGCCAAATCC-CTCTTTTGAAA
Aameth6679     ?????????????????????????????????????????????????
Aameth6685     ?????????????????????????????????????????????????
Aappen6543     ---GGT---GTGTGTTGGGCAA-TGTTGAGATA-TTCCCTTTTTTTTGAA
Aappen6677     --TGGAATGATA-AATGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA
Aappen6946     --TGGAATGAAA-G-TGGGCAA-TCCTGAGCCAAATCC-CTTTTTTGAAA
Adist6684      ?????????????????????????????????????????????????
Aeuco5869      ?????????????????????????????????????????????????

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*Hanam4841* --ACCA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Hcon6942* ???  
*Hdiss5145* AAACAA-GCGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Sbicolour9319* ???  
*Hemarthriaaltissima5698* AAACAA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Hfilli5144* ???  
*Hhirt6550* AAACAA-GCG-----CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Hhirt6656* ???  
*Hlongi6560* AAACAACGAGGTTCT--CGAACCTAGAA-TACAAAGGAAAAGGATAGGTGC  
*Hpilg4603* AAACCA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAAATGC  
*Hpilg4738* AAACCTA-GCGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Hquar5088* ???  
*Hrud6576* ???  
*Htamb6585* ???  
*Icylin5114* AAACAA-GCGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Icylin6632* AAACAA-GCGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Mcape6655* TGACTA-CATGGTCTACCTAATGATGA-TCCAAAGGTAATGGATAGGTGC  
*Pdilatum6617* AA-CAA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Pfrank4705* AAACAA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Pschinz6652* AAACAA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Sbic6589* AAA----GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Shalep9318* AA-CAA-GTGGTT-T--CCAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Snigros6567* AAACAA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Tspic4846* AAACCTA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Tspic6580* AAACAA-GTGGTTCT--CTAAGTAGAA-TCCAAAGGAAAAGGATAGGTGC  
*Ttria6383 b* ???  
*Ttria6555* GGACAAAGA---TC---CATATTGGTC-----GGTCCCGTAGCG-TCT  
*Zea* TGTATCAT---TC---TGTATGTAA-----GCCACTTAG---CTC  
*Elionurusmuticus6954* AAACAA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Icylindrical* AAACAA-GTG-----CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Sacharumofficinarum* AAACAA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Sorghumhalepense* AAA----GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Ttriandra* AAACAA-GCGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Pschinzii* AAACAA-GTGGTTCT--CAAACCTAGAA-CCCAAAGGAAAAGGATAGGTGC

*Aadscen6713* AGAGACTCAATGGAAGCTATTCTA--ACGAATCGAAGTAATTACGTTGTG  
*Aameth6679* ???  
*Aameth6685* ???  
*Aappen6543* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Aappen6677* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Aappen6946* AGAGACTCTATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Adist6684* ???  
*Aeuco5869* ???  
*Aschiren6581* ???  
*Bbladh* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Binsc5681* ???  
*Brachiaria* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAATTACGTTGTG  
*Bserra5074* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Bserra6654* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAATTACGTTGTG  
*Bserra6954* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Cdieter6642* ???  
*Cexc5877* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Cexc6577* -----  
*Cmarg4397* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Cplur6406* ???  
*Cprolix6682* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Dfilli5872* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Hanam4841* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Hcon6942* ???  
*Hdiss5145* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Sbicolour9319* ???  
*Hemarthriaaltissima5698* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAATTGCGTTGTG  
*Hfilli5144* ???  
*Hhirt6550* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Hhirt6656* ???  
*Hlongi6560* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAGTTAATTACGTTGTG  
*Hpilg4603* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Hpilg4738* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Hquar5088* ???  
*Hrud6576* ???



<i>Ttria6555</i>	-----
<i>Zea</i>	-----
<i>Elionurusmuticus6954</i>	-----
<i>Icylindrical</i>	-----
<i>Sacharumofficinatum</i>	-----
<i>Sorghumhalepense</i>	-----
<i>Ttriandra</i>	-----
<i>Pschinzii</i>	TTGGTAGTGGAACTCCCTCGAAATATAGAAAGAAGGGCTTTATA----
<i>Aadscen6713</i>	CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--
<i>Aameth6679</i>	??
<i>Aameth6685</i>	??
<i>Aappen6543</i>	-----AACGATTAATCACA--
<i>Aappen6677</i>	-----AACGATTAATCACA--
<i>Aappen6946</i>	-----AACGATCAATCACA--
<i>Adist6684</i>	??
<i>Aeuco5869</i>	??
<i>Aschiren6581</i>	??
<i>Bbladh</i>	-----AACGATTAATCACA--
<i>Binsc5681</i>	??
<i>Brachiaria</i>	CATTTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACG--
<i>Bserra5074</i>	-----AACGATTAATCACA--
<i>Bserra6654</i>	CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--
<i>Bserra6954</i>	-----AACGATTAATCACA--
<i>Cdieter6642</i>	??
<i>Cexc5877</i>	-----AACGATTAATCACAAA
<i>Cexc6577</i>	-----
<i>Cmarg4397</i>	-----AACGATTAATCACAAA
<i>Cplur6406</i>	??
<i>Cprolix6682</i>	-----AACGATTAATCACA--
<i>Dfilli5872</i>	-----AACGATTAATCACA--
<i>Hanam4841</i>	-----AACGATTAATCACA--
<i>Hcon6942</i>	??
<i>Hdiss5145</i>	-----AACGATTAATCACA--
<i>Sbicolour9319</i>	??
<i>Hemarthriaaltissima5698</i>	CATCTTATATACACGCATATATACTGACATAGCAAACGATTAATCACG--
<i>Hfilli5144</i>	??
<i>Hhirt6550</i>	-----AACGATTAATCACA--
<i>Hhirt6656</i>	??
<i>Hlongi6560</i>	CACCTAATAAACACGTATAGATACTGGCATAGCAAACGATTAATCACG--
<i>Hpilg4603</i>	-----AACGATTAATCACA--
<i>Hpilg4738</i>	-----AACGATTAATCACA--
<i>Hquar5088</i>	??
<i>Hrud6576</i>	??
<i>Htamb6585</i>	??
<i>Icylin5114</i>	-----AACGATTAATCACA--
<i>Icylin6632</i>	-----AACGATTATTCACA--
<i>Mcape6655</i>	-----GCAGCTTAATCACA--
<i>Pdilatum6617</i>	CATCTAATATACACGCATATATACTGACATAGCAAACGATTAATCACG--
<i>Pfrank4705</i>	-----AACGATTAATCACA--
<i>Pschinz6652</i>	CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCATA--
<i>Sbic6589</i>	-----AACGATTAATCACA--
<i>Shalep9318</i>	CATTTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--
<i>Snigros6567</i>	CATTTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--
<i>Tspic4846</i>	-----TACGATTAATCACA--
<i>Tspic6580</i>	-----AACGATTAATCACA--
<i>Ttria6383 b</i>	??
<i>Ttria6555</i>	-----ACGAGT-TTCAGG--
<i>Zea</i>	-----CCGATT-TTTTTT--
<i>Elionurusmuticus6954</i>	-----AACGATTAATCACA--
<i>Icylindrical</i>	-----AACGATTAATCACA--
<i>Sacharumofficinatum</i>	-----AACGATTAATCACA--
<i>Sorghumhalepense</i>	-----AACGATTAATCACA--
<i>Ttriandra</i>	-----AACGATTAATCACA--
<i>Pschinzii</i>	CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCATA--
<i>Aadscen6713</i>	GAACCCATATCATAATATAGGTTCTTTATTTTATAT-----TTTTTTT
<i>Aameth6679</i>	??
<i>Aameth6685</i>	??
<i>Aappen6543</i>	GAACCCATATTATAATATAGGTTCTTTATTTGATTT-----TT
<i>Aappen6677</i>	GAACCCATATTATAATATAGGTTCTTTATTTGATTT-----TT





Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Pdilatam6617  
Pfrank4705  
Pschinz6652  
Sbic6589  
Shalep9318  
Snigros6567  
Tspic4846  
Tspic6580  
Ttria6383 b  
Ttria6555  
Zea  
Elionurusmuticus6954  
Icylindrical  
Sacharumofficinorum  
Sorghumhalepense  
Ttriandra  
Pschinzii  
  
Aadscen6713  
Aameth6679  
Aameth6685  
Aappen6543  
Aappen6677  
Aappen6946  
Adist6684  
Aeuco5869  
Aschiren6581  
Bblad  
Binsc5681  
Brachiaria  
Bserra5074  
Bserra6654  
Bserra6954  
Cdierter6642  
Cexc5877  
Cexc6577  
Cmarg4397  
Cplur6406  
Cprolix6682  
Dfilli5872  
Hanam4841  
Hcon6942  
Hdiss5145  
Sbicolour9319  
Hemarthriaaltissima5698  
Hfilli5144  
Hhirt6550  
Hhirt6656  
Hlongi6560  
Hpilg4603  
Hpilg4738  
Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Pdilatam6617  
Pfrank4705  
Pschinz6652  
Sbic6589  
Shalep9318  
Snigros6567  
Tspic4846  
Tspic6580

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??  
??  
ATTTTT-TAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
ATTTTT-TAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
TTAAATG-AAGAATTC----ATGT-GAAGCTATTGCTTGCAATATATT  
ATTTTT-TAGAATTA----TTGT-GAATCCATTCCAATCGA-ATATT  
ATTTTT-TAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
ATTTTTTTTAGAATTA----TTGT-GAATCCATTCCAATCGA-ATATT  
ATTTTT-TAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
ATTTTT-TAGAATTA----TTGT-GGATCCATTCCAATCGA-ATATT  
ATTTTT-TAGAATTA----TTGT-GAATCCATTCCAATCTA-ATATT  
ATTTTT-TAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
ATTTTT-TAGAATTA----TTGT-GAATCTATTGCAATGGA-ATATT  
??  
ATTTTT-GAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
ACTCCAT-TTGTCTATA---TATT-CCATATATCACATTCGA-AGATA  
ATTTTT-TAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
ATTTTT-TAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
ATTTTT-TAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
ATTTTT-TAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
ATTTTT-GAGAATTA----TTGT-GAATCTATTCCAATCGA-ATATT  
ATTTTT-TAGAATTA----TTGT-GAATCCATTCCAATCGA-ATATT  
  
-TCATAATCA-AATCCTT--CAATTCATTGTTTTG-----AGATCTTTA  
??  
??  
-GAGTAATCA-AATCCTT--CAA-TCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTC-----GAGATCTTTT  
??  
??  
??  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
??  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTG-----AGATC----  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
??  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
AGGGGAGACGGATCACGCACATATATATATTTCATCA----ATATC-TAT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATC-TTT  
??  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTT-----GAGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
??  
-GAGTAATCA-AATCCTT--CAATTCATTATTTTGTGTTGG-CAGATCTTTT  
??  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
??  
-TAGTAATCA-AATCCTT--CAATTCAGT---TTG-----GAGTCTTTA  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
GACTAATCACAATGGAC--TAATACATTGTGTT-----GAGTCAATTT  
-GAGTAATCA-AATCCTT--CAATTCATTATTTTGTGTTTTCGAGATCTTTA  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTG-----AGATCTTCA  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
-GAGTAATCG-AATCCTT--CAATTCATTGTTTTG-----CGATC----  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTG-----AGATC----  
-GAGTAATCA-AATCCTT--CAATTCATTGTTTTG-----AGATCTTTT  
-GAGTAAT-AGAATGGTT--CAATTCATTATTTTTC-----GAGATGTTTT

*Ttria6383 b* ???  
*Ttria6555* -GAGTTATCA-AATCCTT--CAATTCATTGGTTTCG-----AGATCTTTT  
*Zea* -TCATATTCA-TGGAATA--CGATTCACT---TTCA-----AGATGCCTT  
*Elionurusmuticus6954* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Icylindrical* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Sacharumofficinarum* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Sorghumhalepense* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Ttriandra* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Pschinzii* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTG-----AGATCTTCA

*Aadscen6713* AAAAGTGG-ATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Aameth6679* ???  
*Aameth6685* ???  
*Aappen6543* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Aappen6677* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Aappen6946* CCTCAGTGGATTAATC--GGACGAGGATA--AAGAGA--GACTAACA  
*Adist6684* ???  
*Aeuco5869* ???  
*Aschiren6581* ???  
*Bbladh* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Binsc5681* ???  
*Brachiaria* AAAAGTGG-ATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Bserra5074* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Bserra6654* AAAAGTGGGATTAATC---GACGAGGATA--AAGAGA--GAGTCCCA  
*Bserra6954* TAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Cdieter6642* ???  
*Cexc5877* TAAAAGCGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Cexc6577* ATAAGATGTGATAGTC--GAAAGTATACTGGAAAAGAATGACGTACC  
*Cmarg4397* TAAAAGCGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Cplur6406* ???  
*Cprolix6682* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Dfilli5872* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hanam4841* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hcon6942* ???  
*Hdiss5145* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Sbicolour9319* ???  
*Hemarthriaaltissima5698* AAAAAGTGGATTAAGC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hfilli5144* ???  
*Hhirt6550* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hhirt6656* ???  
*Hlongi6560* AAACAGGTGATTAATCCGGGACGAGGACA--AAGAGA--GAGTCCCA  
*Hpilg4603* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hpilg4738* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hquar5088* ???  
*Hrud6576* ???  
*Htamb6585* ???  
*Icylin5114* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Icylin6632* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Mccape6655* AAAAAGTGGATAAATC--GGCGGAGGATT--AAGCGT--GGTTCATA  
*Pdilatum6617* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Pfrank4705* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Pschinz6652* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Sbic6589* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Shalep9318* GAAAGTGG-TATACTC---GAATCGGACG--AGGATAA--GAGTCCCA  
*Snigros6567* AAAAAGTGG-ATTAATC---GACGAGGATA--AAGAGA--GAGTCCCA  
*Tspic4846* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Tspic6580* AAAAAGTGGATTAATC--GGATGAGGATA--AAGAGA--GAGTAGAA  
*Ttria6383 b* ???  
*Ttria6555* AAAAAGTGGATTAATCC--GGACGAAGATT--AAGAGA--GATTCCCA  
*Zea* GA---TGTTGAAAT---GGTAGAC-----ACGCGA--GACTCAAA  
*Elionurusmuticus6954* TAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Icylindrical* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Sacharumofficinarum* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Sorghumhalepense* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Ttriandra* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Pschinzii* AAAAAGTGGATTAATC--GAACGAGGATA--AAGAGA--GAGTCCCA

*Aadscen6713* TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA  
*Aameth6679* ???  
*Aameth6685* ???



<i>Aappen6543</i>	TTCTA--CATGTCAAT--ACTGA-CAACATT--AATTTCTAGTAAAA
<i>Aappen6677</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Aappen6946</i>	TTTTG--AATCCCAAC--GGAAT-CGGG--ACACATAACTTTCAAAAT
<i>Adist6684</i>	??
<i>Aeuco5869</i>	??
<i>Aschiren6581</i>	??
<i>Bbladh</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Binsc5681</i>	??
<i>Brachiararia</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Bserra5074</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Bserra6654</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Bserra6954</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Cdieter6642</i>	??
<i>Cexc5877</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Cexc6577</i>	GTCAACGANTTTCNGG---NNN-TATAGTTCTTTATTATTTTANANA
<i>Cmarg4397</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Cplur6406</i>	??
<i>Cprolix6682</i>	TTCTA--CATGTCAAT--ACTGA-CGGC--AGATGGATTGGAATCCC
<i>Dfilli5872</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Hanam4841</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Hcon6942</i>	??
<i>Hdiss5145</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Sbicolour9319</i>	??
<i>Hemarthriaaltissima5698</i>	TGCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Hfilli5144</i>	??
<i>Hhirt6550</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Hhirt6656</i>	??
<i>Hlongi6560</i>	TTCTA--CATGTCAAT--ACTG---AACAATGAAATTTCTAGTAAAA
<i>Hpilg4603</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Hpilg4738</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Hquar5088</i>	??
<i>Hrud6576</i>	??
<i>Htamb6585</i>	??
<i>Icylin5114</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Icylin6632</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Mcape6655</i>	TTCCG--TATGTCAAT--GCGAT-CGTC--TGACTTTACTTTCTAAC
<i>Pdilatum6617</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Pfrank4705</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Pschinz6652</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Sbic6589</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Shalep9318</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAA- TTTCTAGTAAAA
<i>Snigros6567</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Tspic4846</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Tspic6580</i>	TTATA--CATGTGNATCTGACAA-CAAGAATATAGGAGGTTTATGAT
<i>Ttria6383 b</i>	??
<i>Ttria6555</i>	TCCTA--CATGTCAAT--ACTGGACAACAATGAAATTTCTAGTTAAA
<i>Zea</i>	ATCT---CGTG-CTAA--AGAGCGTGGAGGTTTCGAGTCCT-CTTCAA
<i>Elionurusmuticus6954</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Icylindrical</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Sacharumofficinatum</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Sorghumhalepense</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Ttriantra</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Pschinzii</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA

<i>Aadscen6713</i>	GGA
<i>Aameth6679</i>	???
<i>Aameth6685</i>	???
<i>Aappen6543</i>	GGA
<i>Aappen6677</i>	GGG
<i>Aappen6946</i>	TCC
<i>Adist6684</i>	???
<i>Aeuco5869</i>	???
<i>Aschiren6581</i>	???
<i>Bbladh</i>	GGA
<i>Binsc5681</i>	???
<i>Brachiararia</i>	GGA
<i>Bserra5074</i>	GGA
<i>Bserra6654</i>	GGA
<i>Bserra6954</i>	GGA

<i>Cdieter6642</i>	???
<i>Cexc5877</i>	GGA
<i>Cexc6577</i>	ATG
<i>Cmarg4397</i>	GGA
<i>Cplur6406</i>	???
<i>Cprolix6682</i>	TCC
<i>Dfilli5872</i>	GGA
<i>Hanam4841</i>	GGA
<i>Hcon6942</i>	???
<i>Hdiss5145</i>	GGA
<i>Sbicolour9319</i>	???
<i>Hemarthriaaltissima5698</i>	GGA
<i>Hfilli5144</i>	???
<i>Hhirt6550</i>	GGA
<i>Hhirt6656</i>	???
<i>Hlongi6560</i>	GGA
<i>Hpilg4603</i>	GGA
<i>Hpilg4738</i>	GGA
<i>Hquar5088</i>	???
<i>Hrud6576</i>	???
<i>Htamb6585</i>	???
<i>Icylin5114</i>	GGA
<i>Icylin6632</i>	GGG
<i>Mcape6655</i>	ACC
<i>Pdilatum6617</i>	GGA
<i>Pfrank4705</i>	GGA
<i>Pschinz6652</i>	GGA
<i>Sbic6589</i>	GG-
<i>Shalep9318</i>	GGA
<i>Snigros6567</i>	GGA
<i>Tspic4846</i>	GGA
<i>Tspic6580</i>	TGG
<i>Ttria6383 b</i>	???
<i>Ttria6555</i>	GGG
<i>Zea</i>	GGC
<i>Elionurusmuticus6954</i>	GGA
<i>Icylindrical</i>	GGA
<i>Sacharumofficinarum</i>	GGA
<i>Sorghumhalepense</i>	GGA
<i>Ttriandra</i>	GGA
<i>Pschinzii</i>	GGA

<i>Aadscen6713</i>	?????
<i>Aameth6679</i>	GACCC
<i>Aameth6685</i>	GACCC
<i>Aappen6543</i>	?????
<i>Aappen6677</i>	GACCA
<i>Aappen6946</i>	GATCC
<i>Adist6684</i>	GACCC
<i>Aeuco5869</i>	GACCC
<i>Aschiren6581</i>	GACCC
<i>Bbladh</i>	GACCC
<i>Binsc5681</i>	GACCC
<i>Brachiaria</i>	?????
<i>Bserra5074</i>	GACTC
<i>Bserra6654</i>	?????
<i>Bserra6954</i>	GACTC
<i>Cdieter6642</i>	GACCC
<i>Cexc5877</i>	?????
<i>Cexc6577</i>	GACCC
<i>Cmarg4397</i>	?????
<i>Cplur6406</i>	GACCC
<i>Cprolix6682</i>	GACCC
<i>Dfilli5872</i>	CCTCC
<i>Hanam4807</i>	GACCC
<i>Hcon6942</i>	GACCA
<i>Hdiss5145a</i>	ACCCC
<i>Sbicolour9319</i>	GACCC
<i>Hemarthriaaltissima5698</i>	GACCA
<i>Hfilli5144</i>	GCCCC

Hhirt6650	?????
Hhirt6656f	GAACC
Hlongi6560	?????
Hpilg4603	GACCC
Hpilg4738	GACCC
Hquar5088	GACCA
Hrud6576	GACCA
Htamb6585	GACCC
Icylin5114a	GACCC
Icylin6632a	GACCA
Mcape6655	GACCC
Pdilatum6617	GACCC
Pfrank4705	GACCA
Pschinz6652	?????
Sbic6588	GACCA
Shalep9318	GACCC
Snigros6567	?????
Tspic4846	GACCA
Tspic6580	GACCC
Ttria6383 b	GACCC
Ttria6555	GACCC
Zea	ATAAT
Elionurusmuticus6954	GACTC
Icylindrical	AAATC
Sacharumofficinatum	AAATC
Sorghumhalepense	AAATC
Ttriandra	AAATC
Pschinzii	AAATC
Aadscen6713	??
Aameth6679	AGTT-CAGGGAGATCGGGGGT-TGGGAGAACTAAGGG-TTGGTA
Aameth6685	AGTT-C-GTGAGATAGGGGT-TGGGAGAAAATAAGGGATTGGTA
Aappen6543	??
Aappen6677	AGGTTTCAGGGAGATAGGGGTT-TGGGAGAAAATAACGGATTGGTA
Aappen6946	ATATTTTGGGACCAAAATCCG---GGAGA---TAGGGG---ATAT
Adist6684	CGTTACAGGAAGACGGGGT-TGGGAGAAAATAAGGGCTTGGTA
Aeuco5869	CGGT--CGGGAGATCGGGGT-TGGGAGAAAATAAGGGTTGGTA
Aschiren6581	A-GTTCAGGGAGATAGGGGTGTGGGAGAAAATAAGGGATTGGTN
Bbladh	AAACTNGNCGAGATACCTTTGTTGGGAGAAAACATCGGGTAGAAT
Binsc5681	AAAATNG-GGAGAGACGGG--TTGGGAGAAAATAAGGCATTGGTA
Brachiaria	??
Bserra5074	CGATTGTCGGACCAAGTTCAG---GGAGA---TAGGGG---AAAA
Bserra6654	??
Bserra6954	CGTTTGTTCGGACCAAGTTCAG---GGAGA---TAGGGGN-TATTT
Cdieter6642	AAGTTCAGGGAGATAGGGGTT-TGGGAGAAAATAAGGGATTGGTA
Cexc5877	??
Cexc6577	AAGT-CAGGGGGATACGGGTT-TGGGAGCAAATAACGGGTCGGTA
Cmarg4397	? ?
Cplur6406	C---TCGGGAAAATAGGGGTGTGCGGAGGAAAATACGGGTTTGGGA
Cprolix6682	CAGT--CGGGAGATAGGGGTT-TGGGAGAAAATAAGGGATTGGTA
Dfilli5872	CNTTTTTAGGACCAAGTTCAG---GGAGA---TAGGGGTTTGGAA
Hanam4807	ACCTCCGCAAA-TAGGCCTCCCGCCCGCCCTTCGGCACACCCA
Hcon6942	AGGTTTCAGGGAGATAGGGGTT-TGGGGGCAAATAACGGATTGGTA
Hdiss5145a	TTATAGTCTACACACGTCCC--TGCACA---AAAGGCACTCCTT
Sbicolour9319	G---TCCGGGAATTAGGGGGT-CCGGAGAACATCAAGGGTTGGGA
Hemarthriaaltissima5698	AGGTTTCGGGAGATA-GGTATGTTGGAAAAA-CAAGGGATTGGTA
Hfilli5144	TCTTAGCCCGACCAAGTTC---GGCACA---TAGGGCCCTATTT
Hhirt6650	??
Hhirt6656f	A-GTTTCGGGAATAGGGTG--GTGGGAGAACATAAGGGATTGGTA
Hlongi6560	??
Hpilg4603	AANTTTAGGGAGATAGGGGTT-TGGGAGAAAATAAGGGATTGAAA
Hpilg4738	TCTTACCGGACCC-AAATTC---GGGAGA---TAGGGGTTATTTT
Hquar5088	AGTTCAGGAGATAGGGGGTG--TGTTT---TAAGGGTCAGGTA
Hrud6576	AAGTTCAGGG-GATAGCGGGTTCGGGAGAAAAGAACCGATTGAT
Htamb6585	AAGTTCAGGG-GATAGCGGGTTCGGGAGAAAATAACCGAATTGTA
Icylin5114a	AANTTTTCAGGAGATAGGGGTTTGGGAGAAAATAAGGGTTTCGGTA
Icylin6632a	--GTTTCAGGGAAAATACGGGGGTT--GGGGGAAAATAAGGGATTGGTA
Mcape6655	ACGTTTCAGGGAAAATAAGGGGGTTGGGAGAAAATAAGGGATTGGTA
Pdilatum6617	AGCTTCCGGGAATTAGGGGCCCTTGGGGCCCAACATTTTAGCCCAA
Pfrank4705	AAGTTTTGAGCACCAAGTTCG--GGGAGA---TAGGGGTCATTGA
Pschinz6652	??

Sbic6588 G---TTCGGGAATTAGGGGGT-TGGGGGAAAATAAGGGATTGGGA  
 Shalep9318 TCTTACCGGACCCCATATA---CCGATT---TCCCGGTTATTTT  
 Snigros6567 ???  
 Tspic4846 TCGTTGTCCANAAAAGTTTC---GCANA---TAGGGGTATTTT  
 Tspic6580 CG-TTCAGGGAAATCCGGGATTCCCGGCAAAAATACGGGCTGGGTA  
 Ttria6383 b AAGTTCAGCAGATAGGGTT--GTGGGAGAAAATAAGGGATTGGTA  
 Ttria6555 AAGTTCAGCAGATAGGGTT--GTGGGAGAAAATAAGGGATTGGTA  
 Zea C-GGAGAATGCTCATTGAT--GAGCATTCCCCTAGAAGTATTC--  
 Elionurusmuticus6954 CGTTTGTTCGGACCAAGTTCAG---GGAGA---TAGGGGN-TATTT  
 Icylindrical CGTCGACTTTATA-AGTCGT-GAGGGTTCAAGTCCCTCTATCCCC  
 Sacharumofficinorum CGTCGACTTTATA-AGTCGT-GAGGGTTCAAGTCCCTCTATCCCC  
 Sorghumhalepense CGTCGACTTTATA-AGTCGT-GAGGGTTCAAGTCCCTCTATCCCC  
 Ttriandra CGTCGACTTTATA-AGTCGT-GAGGGTTCAAGTCCCTCTATCCCC  
 Pschinzii CGTCGACTTTATA-AGTCGT-GAG????????????????????

Aadscen6713 ???  
 Aameth6679 TCC--CTTTAGGAAACAAGAATA---TAGTTACCCA-CACTC  
 Aameth6685 TCC--CTTTAGGAAACAAGAAA---TAGTTACCCA-CACTC  
 Aappen6543 ???  
 Aappen6677 TCA-ACATTAGTAAACAAGAAAC--TAGTTACCCA-CATTC  
 Aappen6946 TCTTCGC---ACACA-AGG-----CA-ATAAAC---CTTG  
 Adist6684 TCC--CATTAGGAAACAAGAATA---TAGTTACCCA-CATTC  
 Aeuco5869 TCN-GTTATTGTTCAAGTAGAAAC--TAGTTACCCATCATCC  
 Aschiren6581 TCA-ACNTTG-GAGATAGGGGA---TAGTTACCC-GGATTC  
 Bbladh CA---CATCA-TCTTACCGAAAGACATAGTTACCCA-TACTC  
 Binsc5681 TCAAATA-TAGGAAA---CAAAGA--TAATAGTTACCAATTC  
 Brachiararia ???  
 Bserra5074 ACTGGCG---ACACG-GGC-----CA-ATAAAC---CTTG  
 Bserra6654 ???  
 Bserra6954 TTTCGGG---TAAAG-AGA-----AA-ATAAAC---CTTC  
 Cdieter6642 TCA-GCTATCGGAAACGTAGAAAA--TAGTTACCCA-AATTC  
 Cexc5877 ???  
 Cexc6577 CCA-TCATTCTGAAACATAGAAAA--TAGTAACCCA-ACTTC  
 Cmarg4397 ???  
 Cplur4606 TCAA-CTTTAGGAAAA-AAGGAA--ATAGGTTTCCC-GATTC  
 Cprolix6682 TCA-GCTATNGGAAAAATAGAAAA--TAGTTACCCC-AATTC  
 Dfilli5872 G-----AAAATAAGG-----GA-----TTGG  
 Hanam4807 GAT-AAAAATCTAGACTCGGA-AAGGTCTAGNCCCA-CAGTT  
 Hdiss6942 TCA-NCTATAGGAAAAATAGAAAA--TAGTTACCCA-CATTC  
 Hdis5145a TTCTCCC---ATCTTTTGG-----CACC-AAA---ATTC  
 Sbicolour9319 GTCA-ACTTAGGAACA-AAGG---ATAGGTTCCCC-A-TTC  
 Hemarthriaaltissima698 TCCCTCATCGGAAGAAAAGAAATAAATTTACCCA--ATTC  
 Hfilli5144 TTTCCGA---CACCATTG-----CAGATAA-----ATAA  
 Hhirt6650 ???  
 Hhirt6656f CCA-ACNTTAGGAGAAAAGAGTA--TAGTTACCCAGTATTC  
 Hlongi6560 ???  
 Hpilg4603 TA--TCATAAATAGGAGAAAAGAAAATAGTTACCCA-AATTC  
 Hpilg4738 TCGGGTC---AATTGCAGG-----GATAA-----ATAG  
 Hquar5088 CATGGGT---TCCGGACC-----AAGGTTTCAG---GGAG  
 Hrud6576 GCA-ACATT-GTAAATCTAG-AAA-GTAGTTAGCCG--ATTC  
 Htamb6585 GCA-CCATTGTAACATAG-AAA-CTAGTTACCCC-CATTC  
 Icylin5114a TCATAGAGTATGCAAAAAAAGAGTAATAGTTACCCA-AATTC  
 Icylin6632a TCTG-CTATGGGAAAAAAGAAA--ATAGTTACCCA-AATTC  
 Mcape6655 TCAA-CTTTAAGAAAAAGAGAC---AAGTTTCGGG-ACTTC  
 Pdilatatum6617 GCCTTGCTCAGTTATTCTGGTAACAGTAGATATCCCAATTA  
 Pfrank4705 TCGGGTC---AATAGGACC-----ANGTTTCAG---GGAG  
 Pschinz6652 ???  
 Sbic6588 TCAA-CTTTAGGAAAA-AAGG---NTTGTACCCC-AATTC  
 Shalep9318 TTTGTAA---GACCAAAGT-----CAGATAA-----CATG  
 Snigros6567 ???  
 Tspic4846 TCGGACC---AAGTGCAGG-----GAG-----ATAG  
 Tspic6580 TCT--CTAACCAAAAACAAGAAG--ATAGTTGTCCC-CATTC  
 Ttria6383 b TCC--CATTCGCAATAAAGAAA---TAGTTACCA--AATTC  
 Ttria6555 TCC--CATTCGCAACAAGAAA---TAGTTACCA--AATTC  
 Zea AAT--CTGGCGCTCTCCTCTATC---TAATGAATA--CTTTA  
 Elionurusmuticus6954 TTTCGGG---TAAAG-AGA-----AA-ATAAAC---CTTC  
 Icylindrical AAA-CCCTCTTTTATCCCTAACC-ATAGTTGTTAT-CCTTT  
 Sacharumofficinorum AAA-CCCTCTTTTATCCCTAACC-ATAGTTGTTAT-CCTTT  
 Sorghumhalepense AAA-CCCTCTTTTATCCCTAACC-ATAGTTGTTAT-CCTTT  
 Ttriandra AAA-CCCTCTTTTATCCCTAACC-ATAGTTGTTAT-CCTTT

*Pschinzii* ???

*Aadscen6713* ???

*Aameth6679* TGA-GTG-ATC-GAAAGAG-TAAG-AGGAGAAAG-TGTTTCCTCACGC-

*Aameth6685* TAA-GTG-ATC-GAAAGAC-TAAG-AGGAGAAAG-TGTTTCCTCACGC-

*Aappen6543* ???

*Aappen6677* TAA-GTG-ATC-GAAAGCA-TAAG-ATGAGAA-AGTGTTCCTCACGC-

*Aappen6946* ACCCCGGT--G-CTCCTAAATATC----GGACC--ACGTTTCAGGGAGAT

*Adist6684* TAATGTG-ATC-GAAAGACGTAAG-ATGAGAAAG-TGTTTCCTCACGC-

*Aeuco5869* TGGTGTGGAGC-GAAAGAA-TAAG-ATGAGAAAGTGTGTTTCTTCGCGC-

*Aschiren6581* TAG-GTGA-TC-GAAATGAGTAAGATGAGAAATG-AGTTTCCTCACGC-

*Bblad* TAATGTA-ATCTNAAANAC-TAATCATGCCATCGGTCTTTCCTCCCC

*Binsc5681* TAA-GTG-ATC-GAATCAG-TAAG-ATGAGATAG-TGTCTCCTCACGC-

*Brachiaria* ???

*Bserra5074* --CACTGT--G-CTCA-AAATATC----GGACC--ACGTTTCAGGGAGAT

*Bserra6654* ???

*Bserra6954* A-CNCNGT--G-CTCATAAAAATC----AGACC--ACGTTTCAGGGAGAT

*Cdieter6642* TGA-GTG-ATC-GAAAGAG-TAAG-ATGAGAA-AGTGTTCCTCACGC-

*Cexc5877* ???

*Cmarg4397* TTT-GTG-ATC-TAAAGAG-TAAG-ATGAGTT-AGCGTTTCTCACGC-

*Cplur6406* ???

*Cprolix6682* TGA-GTG-ATC-GAAA-ACGTAAG-ATGAGAAA-GAGTTTCCTCACCT-

*Dfilli5872* TCA-GTG-ATC-GAAAGAG-TAAG-ATGAGAG-AGTGTTCCTCACGC-

*Hanam4807* TATAAAATAAGGGATTGGTATGGA----TTG----GTATT-----AT

*Hcon6942* TAAGGACAATAGGAGTTATTTTTTC--GGGAAATATATATCCTAGGTCT

*Hdiss5145a* TAATGTG-ATC-GAAAGAA-TAAG-ATGAGAA-TGTGTTTCCTCGCGC-

*Hemarthriaaltissima5698* AAGGAAATAGGGGTTTGGGACAAA----ATAAC--GGATTGGT----AT

*Hfilli5144* TAA-GTA-ATC-GATAAAGTACTA-A-GAGAAA-GTGTACCTCACGC-

*Hhirt6650* TA--AGA-ATC-GATAGCG-TAAG-ATGAGAACG-TGTTTCCCCGCGC-

*Hhirt6656f* CAGTGTGT--GAGATAGAAATAAT----TACCA--AGTTTCAT---CTA

*Hlongi6560* ???

*Hpilg4603* TGA-GTGGATC-GAAAGACGTAAGATGGAGAAAG-TGTTTCCTCAC-C-

*Hpilg4738* ???

*Hquar5088* TA-AGTA-ATC-GAAAGAG-TAAG-ATGAGAAAG-TGTTTCCTCACGC-

*Hrud6576* CATGGGTTGGAGGATAAAAATTAA----GTACC--AAGTTTGT---CCT

*Htamb6585* CGTAGGTGTGGGAGAGAACTAAG----GGATA--GGTATCAT---CAT

*Icylind5114a* TTA-GCG-ATCGAGGAGAG-TAAG-ATGAGAAAG-CGTTTCCTCACGC-

*Icylind6632a* TAA-GTT-ATCGAA-AGCG-TAAG-ATGAGATTG-CGTTCCTCACGC-

*Mcape6655* TAA-GTG-ATC-GAATCAG-TAAG-ATGAGAAAG-TGTCTCCTCACGC-

*Pdilatum6617* NAA--GNNA--T-CAAANA-GNAAG-GG-AGNAAANAGTTTCCTCCC-C-

*Pfrank4705* TCA-TTG-ATC-GAAATAG-TAAG-AGGAGACCG-AGTTTCCTCACGC-

*Pschinz6652* AAAGCCCATATAAATGGGCTTAA--GAGTCACTAGCTTTATCATTCT

*Sbic6588* ATAGGGGTTGGGAGAAAA-TAAG----GGATT--GGTACCAT---CAA

*Shalep9318* ???

*Snigros6567* TAA-GGG-ATC-GAAAGATGTAAG-G-GAGAAA-GTGTTCCTCACGC-

*Tspic4846* GTGGGCGTT-GGGAAAGAAATAAC----AGGTT--TGTCTCAC---CT-

*Tspic6580* ???

*Ttria6383 b* CCGGGGGGAGGGAGAAAA--TAG----GTACC--GGTATCGT---CAT

*Ttria6555* TGA-GTAAATT-GAAGGA-GTAAC-AACAGCAGTCGGTGTCCACAAGC-

*Zea* TTG-GTGG-TC-GAAAGA-GTAAGT--GAGGATG-AGCAGTCTCACGC-

*Elionurusmuticus6954* TTG-GTGG-TC-GAAAGA-GTAAGT--GAGGATG-AGCAGTCTCACGC-

*Icylindrical* TTG-ATTT--TT-TAGTCC-CTTTAA--TTGAAT--AG-----ATGC-

*Sacharumofficinarum* A-CNCNGT--G-CTCATAAAAATC----AGACC--ACGTTTCAGGGAGAT

*Sorghumhalepense* TTT-TCT-TTT-ATCAAT-GGG--TTT----AAGA--TTC-A-CTAGC-

*Ttriandra* TTT-TCT-TTT-ATCAAT-GGG--TTT----AAGA--TTC-A-CTAGC-

*Pschinzii* TTT-TCT-TTT-ATCAAT-GGG--TTT----AAGA--TTC-A-CTAGC-

???

*Aadscen6713* ???

*Aameth6679* -TGTTTCG-TGAGTTACTCA-GAATAC-GATCA-GTAATTTAT-CTACTAA

*Aameth6685* -TGTTCT-TGAGTTACTCA-GAATAC-GATAA-GTAGTTTAT-CTACTAA

*Aappen6543* ???

*Aappen6677* -TGTTCTATA-GTCTCTA-GACTGC-GCTC--CGTAGTTT-CTACTAA

*Aappen6946* GGGGTGTGGGAGAAAAATAAGGGATTGGTATCAACTTTCGGAA---AC-AA

*Adist6684* -TGTTCT-TGAGTTACTTA-GAATAC-GATAA-GTAAGTTAT-CTTCTAA

*Aeuco5869* -TGTTTCAGAAATGACTGA-GAATAC-GATCG-CCAAGGTAT-GTACGAA

*Aschiren6581* -TGTTTC-TGGAGTTACTTA-GAGTAC-GATAA-GTAATTTAT-CTGCTAA

*Bblad* -TTCTCACGGCGTTACT-A-GAATCCCGATAACGTAATTTCTTCTCTTAA

*Binsc5681* -TGTTCT-TGAGTTACCGA-GAATAC-GATGA-GTGAGTTAT-CTACGAA

*Brachiaria* ???

*Bserra5074* GGGGTTGGGGAGAAAATAAGGGATTGGTATCACCAATAGGAA---ACGAA  
*Bserra6654* ???  
*Bserra6954* GGGGTTGGGGAGAAAATAAGGGATTGGTATCACCTATCGGAA---ACTAA  
*Cdieter6642* -TGTTCTTGA-GTTACTTA-GAATAC-GATCA-GTAATTTAT-CTACTAA  
*Cexc5877* ???  
*Cexc6577* -TGTTCTTGA-GTTACTTA-GA-TAC-TATC--CTAAGTTT-CTTCTAG  
*Cmarg4397* ???  
*Cplur6406* -GTT-CT-TGAGTTTCTGATGGATAC-GATAA-GTTGTTTAT-CTACTAA  
*Cprolix6682* -TGTTCTTGA-GTTACTTA-GAATAC-GATAA-GTAAGTTAT-CTACTAA  
*Dfilli5872* ---CATCAATA-GGAA-AATAGTTACCCAAATTCTAAG-TAA---TCGAA  
*Hanam4807* AGGTTTGGAAATAAGAAGA-TAATACCGAGTTTGTCTCGGTGCT-CGAG  
*Hcon6942* -TGTTCTATA-AACTTTAC-CACTGG-ACTC--CTGAGTTAT-CG-CTGA  
*Hdiss5145a* ---CATCAATA-GGAA-AATAGTTACCCAAATTCTAAG-TAA---TCCAA  
*Sbicolour9319* -CCT-CT-GCGGTTTTTTA-GAATAC-GACAA-CTAATTTAT-CTACTAG  
*Hemarthriaaltissima5698* -GGCTCT-GGGTTACTTA-GAATAC-GATGA-GTAATTTAT-CTACTAA  
*Hfilli5144* TAGTTACAAA-AAAATAACTAACCAGTTCC-CT-AAG---CCTAA  
*Hhirt6650* ???  
*Hhirt6656f* -TGTTCTGACGTTGCTGA-AGATAC-GATAAAGTAAATTTAT-CT-AGTA  
*Hlongi6560* ???  
*Hpilg4738* -TTCTCT-TGAGTTACCTA-GAAT-----A-GGATCTTAT---ATAA  
*Hquar5088* AAGTGATAGA-AAGAATA-AGATGCCAAAGTGTAGCCTTG---GCTAG  
*Hrud6576* TAGGAAAAAGAGAAGACATAGTTTCCAGATTCTACG-TAG---TCGAA  
*Htamb6585* -TGTTCT-TGAGTTACTTA-GAATAC-GATAG-GTAATTTT--CTGCTAG  
*Icylin5114a* -TGTTCT-TGAGTTACTTA-GAATAC-GATAA-GTAATTTT-CTACTAA  
*Icylin6632a* -TGTTCT-TGAGTTACTTA-GAATAC-GATAA-GTAAGTTAT-CTACTAC  
*Mcape6655* -TGNCT-TCAGTTTCCCTA-GAATCC-GATCA-GTTATTTAT-CTACTAA  
*Pdilatum6617* -TGCTCT-TGCGTTACTTA-GAATACGA-TAG---TATTTTACCTATGCTA  
*Pfrank4705* CTAT---GGAGCAAAGGA-GAGTGACGAGAACCTCAATGTATCTAGTCT  
*Pschinz6652* TAGG--AAAA-AAGAAATAGTTACCCAAAGTCTAAG-TAA---TCGAA  
*Sbic6588* ???  
*Shalep9318* -CGTTCT-TGGGTTTCTTA-GAATAC-GATAA-GTAATTTAT-CTACTAG  
*Snigros6567* TCCAAAAATA-AAAAAATACCTACCCAAATTCTACT-TCT---TCGAA  
*Tspic4846* ???  
*Tspic6580* AGGA--AAAA-AAGAATATAGTTACCCAAATTCTAAG-TAG---TCGAG  
*Ttria6383 b* -CGTTCC-CGAGTTTCTTC-GTATCC-CCTCA-GTCATATAA-TTTCNAA  
*Ttria6555* -AGTTC-TGA-GTTACTTA-GAGTAC-GATAA-GTAATTTAT-TTGAGCT  
*Zea* -AGTTC-TGA-GTTACTTA-GAGTAC-GATAA-GTAATTTAT-TTGAGCT  
*Elionurusmuticus6954* -AAATA---CATTACTAA-GAGATG-CACAA-GAAAGGG---TCAGGAT  
*Icylindrical* GGGGTTGGGGAGAAAATAAGGGATTGGTATCACCTATCGGAA---ACTAA  
*Sacharumofficinatum* --TTTC-TCA--TT-CT-AC-TCCTT-CACAAAGGAGTGCACAAAGAACT  
*Sorghumhalepense* --TTTC-TCA--TT-CT-AC-TCCTT-CACAAAGGAGTGCACAAAGAACT  
*Ttriandra* --TTTC-TCA--TT-CT-AC-TCCTT-CACAAAGGAGTGCACAAAGAACT  
*Pscinzi* ???

*Aadscen6713* ???  
*Aameth6679* AGTTAAACT-----AAAGAAAAAAATCAA-TAATCTCATCTCAT--  
*Aameth6685* AGATATCTCT-----ACAGAAAAAAATCTA-TAATCTCATCTCAT--  
*Aappen6543* ???  
*Aappen6677* AC-----AGACAAGACCA-TAGTCTCATA-----  
*Aappen6946* GA-----ACAATAT-----AATCCCATCTCTTAG  
*Adist6684* AGCTTATCT-----ACAGAAAAAAATCTA-TAATCTCATCTCAT--  
*Aeuco5869* GG-----AATCTAGCATA--AGGCTCATA-----  
*Aschiren6581* AGCCAAGGTC-----TGGATGAAGAAATCTA-TAATCTCGTCTCAA--  
*Bbladh* AA-----AACCAATT-----AATCTCAT-----  
*Binsc5681* TGT-----AATGATAAAGATTA-TAATCTCATCTCAA--  
*Brachiaria* ???  
*Bserra5074* GA-----AAAAATAT-----AATCTCATCACTTAG  
*Bserra6654* ???  
*Bserra6954* GA-----ACAATAT-----AATCCCATCACTTAG  
*Cdieter6642* AG-----ACAT-AATATA-TAGTCTCATA-----  
*Cexc5877* ???  
*Cexc6577* AT-----ACCTTAATCTA-TAGTCTCAA-----  
*Cmarg4397* ???  
*Cplur6406* ACTAAGGCT-----CGAGAGAAGAACTCA-CAATCTCGTCTCTG--  
*Cprolix6682* AG-----ACTT-AATCTA-TAGTCTCATA-----  
*Dfilli5872* AG-----AGTAAGATG-----AGAAAGTGTTC  
*Hanam4807* GGTTACAGACGATACGATGTGTGCCTTAGCTGCATAAAAAAACCCTAGCA  
*Hcon6942* AC-----AGCGAGCACA--AGCTTCGTA-----  
*Hdiss5145a* AG-----AGTAAATG-----AGAAAGTGTTC  
*Sbicolour9319* AGAGCTTATA---G-ACGAAAGAAAAACAAA-TAATCTCGTCTCAA--

*Hemarthriaaltissima*5698 AG-----AAGAAATAATC-TAACCACATAGC----  
*Hfilli*5144 AT-----AAAAAACCTA--GAATAAAGTAATTC  
*Hhirt*6650 ???  
*Hhirt*6656f AGCCTAGGCT-----CGAGAGAAGAAATCTA-GAATCTCGTCTCTA--  
*Hlongi*6560 ???  
*Hpilg*4603 AG-----AAAAAAT-----AATCTCATAGCC---  
*Hpilg*4738 AA-----TGAAGAAACCT--AGAATAGGGTAACCTA  
*Hquar*5088 AG-----AGTATGTTG-----AGACAGGGTTTTCC  
*Hrud*6576 AGT-----GAAGAAATAAA-TAATCGCGTCTCA---  
*Htamb*6585 AGT-----AAAAAATAAA-TAATCTCGTCTCAT--  
*Icylin*5114a AGG-----AATGATGAAGCCTA-TAATCTCATCTCAT--  
*Icylin*6632a ATAAA-----GCATGAAGGAAAAAACCTA-CTATATAGTCTCAT--  
*Mcape*6655 ACCTAAGATC-----TAGTGAAGAAATCCA-GAATCTCATCTCCT--  
*Pdilatum*6617 AC-----ACACGGAAT-----AGATGGTATTTT--  
*Pfrank*4705 AG-----AGTAAGA-----AGAGAAAGGGTATCC  
*Pschinz*6652 ???  
*Sbic*6588 AGAGCCTATA---G-AGGAAAGAAAAATCAA-TAATCTCGTGTCTT--  
*Shalep*9318 AG-----AGCCAGATGAG--CAATAAATCGTTTTCC  
*Snigros*6567 ???  
*Tspic*4846 AG-----AGTAAATCCG--AGAGTAAGGGAAGCC  
*Tspic*6580 AGGAATTATCCACGGCTTTAATAAAAAAATCAAGCAATCTCATCTCAA--  
*Ttria*6383 b ACTCAAGGCC-----CGATTGAGGAAACCTA-GAATCTCGTCTCTA--  
*Ttria*6555 ACTCAAGGCC-----CGATTGAGGAAACCTA-GAATCTCGTCTCGA--  
*Zea* A-TCATGGTA-----GAGCAGAGGAC-TGAA-AATCCTCGTCTCAC--  
*Elionurusmuticus*6954 GA-----ACAATAT-----AATCCCATCACTTAG  
*Icylindrical* A-TGAATCTTATGCTATTTCATTAATA----GATGATTTCTTTTTTA--  
*Sacharumofficinorum* A-TGAATCTTATGCTATTTCATTAATA----GATGATTTCTTTTTTA--  
*Sorghumhalepense* A-TGAATCTTATGCTATTTCATTAATA----GATGATTTCTTTTTTA--  
*Ttriandra* A-TGAATCTTATGCTATTTCATTAATAAAATAGATGATTTCTTTTTTA--  
*Pschinzii* ???

*Aadscen*6713 ???  
*Aameth*6679 AGCCTTTCCTTAGAG-CTACTAATTC-ATCCATTAATAATTCATAA  
*Aameth*6685 AGCCTTTCCTTAGAG-CTACTAATTC-ATCTATTAATAATTCATAA  
*Aappen*6543 ???  
*Aappen*6677 -GCCTTTCCTTAAGCTCTAACAATTT-AGCTCTTTACCGTCATAA  
*Aappen*6946 CCGTTCCTTAGAGATAATAATTAAGC-TAATAAAATTCATAATAC  
*Adist*6684 AGCCTTTCCTTAGAG-CTAATAATTC-ATCCAAAAAATTCATAA  
*Aeuco*5869 --TCTTTCCTTAGAG-CTAACATTTT-CGCTCATTAGGGTCATAA  
*Aschiren*6581 AGCCTTTCCTTAGAG-CTACTAATCC-GGCTATTAACATTCATAA  
*Bblad* ---CTTTATGTAGAC-CTGTAATCA-ATTTATTANTCTTCATAA  
*Binsc*5681 AGCCTTTCCTTAGAG-CTAATTATCA-GACTATTAACATTCATAA  
*Brachiaria* ???  
*Bserra*5074 CCGTTCCTTAGAGATACTAATTAAGC-TACTAAAATTCATAATAA  
*Bserra*6654 ???  
*Bserra*6954 CCGTTCCTTAGAAATAATAATTAAGC-TACTAAAATTCATAATAA  
*Cdieter*6642 -GCCTTTCCTTAGAG-CTAATAATTA-AGCTATTAACATTCATAA  
*Cexc*5877 ???  
*Cexc*6577 -GCTTACCTTAGACTCTACTAATTT-ATCTATTACATTCATAA  
*Cmarg*4397 ???  
*Cplur*6406 AGTCTTCCCTTAGAG-CTACTTATCC-GTCTATTAGCATTCATAA  
*Cprolix*6682 -GCCTTTCCTTAGAG-CTACTAATCC-AGCTATTAACATTCATAA  
*Dfilli*5872 TTACGCTGCTTTTGAGTTACTTAGAA-TACGATAAGTAAT-TTAT  
*Hanam*4807 CGGCACAAGTACGTTCTTAGTACAACAGTCTTTCCGTTCTCGGAT  
*Hcon*6942 -GTCTTCCCTGAAAC-CTCACAATC-TGCTCTAATCCGTGATAA  
*Hdi*5145a TCACCCTTCTTTGAGTTTCTTAAAA-TACGATAAGTAAT-TTAT  
*Sbicolour*9319 AGTCTTCCCTTAGAG-CTACTAATTA-AGCCAATAAAATTCATAA  
*Hemarthriaaltissima*5698 ---CGTTCCTAAGAT-CTAATAATTAAGCTCATAATTCATAATAA  
*Hfilli*5144 TCTAACTTCTTAATATCACTTAC-A-TTCATTAGGGATAAATTAAT  
*Hhirt*6650 ???  
*Hhirt*6656f AGCCTTCCCGTATAG-CTACTTATCC-GTCCATTAAGATTCATAA  
*Hlongi*6560 ???  
*Hpilg*4603 ---CTTTCTTAGGGT-CAATAAGTGAGATAAAAATAGTCATAATAA  
*Hpilg*4738 TGTTTACTCGTATAATATCATATCCG-TTCATTAGGCATA-ATCT  
*Hquar*5088 TCACACTGCTCTCGTGTGCTTAGAA-ATCGATACGTTAT-TTGC  
*Hrud*6576 AGCCTTTTCCTTAGAG-CAACTTATTT-GTTTATTAGCATTCATAA  
*Htamb*6585 AGCCTTTTCTTAGAG-CTACT-ATTA-ATCTATTAACCTTCATAA  
*Icylin*5114a AGCCTTCCCTTAGAG-CTACTTATTC-ATCCATTAAGATTCATAA  
*Icylin*6632a AGCCGTCCCTTAGAG-CTACTTATCA-ATCTATAAACATTCATAA  
*Mcape*6655 AGCCGTTTCTTAGAG-CTACTCATTA-GTCTATAAGGTTTCATAA  
*Pdilatum*6617 ----TTGGATAGAGACGCGT--TTA-GGC---AAGGGATCATGA

*Pfrank4705* TCACGCCGCGCATGAGATACTTAGCA-TACGATAAGTAAT-ATAT  
*Pschinz6652* ???  
*Sbic6588* AGCCGTTCCCTTAGAG-CTACTAATA-AGCCAATAAGATTCATAA  
*Shalep9318* TCACGCTGCTCTTTTTTTCCTTAGTA-TACGATAAGCATT-TTAT  
*Snigros6567* ???  
*Tspic4846* TCTCGCCGCGCATGAGTTCCTAACCG-TTCAATAGGTATT-TTAT  
*Tspic6580* AGCCGTTCCCTTATAC-CTACTTATCC-ATCTATTAACATTCATAA  
*Ttria6383 b* AGCCTTCCCGTAGAG-CTACTTATCC-GTTCATTAGGGATCATAA  
*Ttria6555* AGCCTTCCCGTAGAG-CTACTTATCC-GTTCATTAGGGATCATAA  
*Zea* CAGTTCAAA-TCTGG-TTCCTCAGA-----AAAAAAGGATCATA  
*Elionurusmuticus6954* CCGTTCCTTAGAAATAATAATTAAGC-TACTAAAATTCATAATA  
*Icylindrical* TTTCTTTTTTTTTTTTATTT-ATTAGAGT--ATCGGCAA--GGAA  
*Sacharumofficinatum* TTTCTTTTTTTTATTTATTAGAGTAGAGT--ATCGGCAA--GGAA  
*Sorghumhalepense* TCTCTTTTTTTTATTTATTAGAGTAGAGT--ATCGGCAA--GGAA  
*Ttriantra* TTTCTTTTTTTT---T-ATTAGAGTAGAGT--ATCGGAAA--GGAA  
*Pschinzii* ???

*Aadscen6713* ???  
*Aameth6679* TAAATCAT-GCATGAGGT---GTTCCGTATCCTGAAGGGGAGG  
*Aameth6685* TAATTCATTGCAAACGGT---GTTCCGTATCCTGAAGGGGAGG  
*Aappen6543* ???  
*Aappen6677* TAACAT-ATCTAGTCGGT---GTTGCATACTCTGAGGAGAAGCT  
*Aappen6946* TTCATTG-GGTAGGTGTT-----ACGTATCCTGATGC-GAGGG  
*Adist6684* Adist6684 GCATCAT-GCGTAGCGT---GTTACGTATCCTGAGGGGAGG  
*Aeuco5869* TTACTTTACCCATCGGT---GTTTCATACACTGAGGACGAGCC  
*Aschiren6581* TTATTCAT-GCAATAGGT---GTTTCGTATCCTGAGGGGAGG  
*Bbladh* CACTTACTTCCATGGGT---TTTCCGTCTGGGCTAATACCTA  
*Binsc5681* TTAATCAT-TCAATAGGT---GTTTCGTATCCTGAGGGGAGG  
*Brachiaria* ???  
*Bserra5074* TTCATGC-GGTAGGTGTT-----ACCTATCCTGATGC-GAGGG  
*Bserra6654* ???  
*Bserra6954* CTCATTC-GGTAGGTGTT-----ACCTATCCTGATGC-GAGGA  
*Cdieter6642* TAATTC-AGGCGGTAGGT---GTTTCGTATCCTGAGGGGATGG  
*Cexc5877* ???  
*Cexc6577* TTAATGCAACCTATC-GT---GTTTCATTTACTCAAGCCCATCC  
*Cmarg4397* ???  
*Cplur6406* TTAAAGTAACCAGTCGTT---GTTTCGTATCCTCAAGCCGAGGG  
*Cprolix6682* TTATTC-ATGCAGTAGGT---GTTTCGTATCCTGAGGGGACGG  
*Dfilli5872* CTACTA-AAGAAAAA-----TAACATCCTGATGGGGACGG  
*Hanac4807* AATATTCACTCAGTAAGT---GTCATGTCCAGTAAGGG-GAAGG  
*Hcon6942* TAACAA-ACGGGGTGCCT---GTTGCGTACTCTTAGGCTGATCC  
*Hdiss5145a* CTACTA-AAAAAA-----GAAACTATCCTTATGGGGGAG  
*Sbicolour9319* T-AATTCACTCAATTGTT---GTTTCCTATCCTGAGGGGAGGA  
*Hemarthriaaltissima5698* CTCATTCTGTTGAGTTGAGC-CATCGTGAACGTGGGGGTACAG  
*Hfilli5144* TCATTC-AGTAACGTGTT-----ACGTATACCAATAG-GACTG  
*Hhirt6650* ???  
*Hhirt6656f* TTCTTTCT-GCGGGAGTT---GGTTCGTGTCCTCAGGGGAGGG  
*Hlongi6560* ???  
*Hpilg4603* TTCTTTTCGGTACATGTGTAGGTATCTTGATGGGGGGGTTAAAAG  
*Hpilg4738* TTCATTC-AGTAAGTGT---AGGTATAGTAATAG-GACGG  
*Hquar5088* CTACTA-TAGAAA-----TAAGTACTCTCATCG---CCG  
*Hrud6576* TTAATTAG-GCCATCGTT---GGTTCGTTTACTCAAGCGGAGGG  
*Htamb6585* TTCTTTAA-TCAATCGTT---GTTTCGTATCCTCAAGCGGAGGG  
*Icylin5114a* TTCATCAA-TCAATAGGT---GTTTCGTATCCTAAGCGGAGGG  
*Icylin6632a* T-TATTCATCTATCGTT---GTTTCGTATCCTCAAGCGGAGGG  
*Mcape6655* T-AATTTAAGCAGTAGTT---GTTTGGTATCCTCAAGCGGAG-C  
*Pdilatum6617* TTAAGATAGGAAA-----GTAATGTATAATAAAGA-GAGAG  
*Pfrank4705* CTACTAC-AGAAAATGTT-----AATATCCTAATGA-GGCGG  
*Pschinz6652* ???  
*Sbic6588* T-ACTTCATTCGGTAGGT---GTTTCGTATCCTGATGGGGAGGG  
*Shalep9318* CTGCTA-AAGGAAGCTT-----AATCCATCCGAGTGA-AACCG  
*Snigros6567* ???  
*Tspic4846* CTCCTAC-AGTAAGAGTT-----AGCTATCCTGATAG-GACGG  
*Tspic6580* T-TAATTCACC-ATCGTT---GGTTCGTATCCTCAAGCGGAGGG  
*Ttria6383 b* TTCATTCA-GCAATTGTT---GTTTCGTGTCCTCAGGGGAGGG  
*Ttria6555* TTCATTCA-GCAATTGTT---GTTTCGTGTCCTCAGGGGAGGG  
*Zea* TGAATAGA-TCAATTGAT---TACAAATATTTGAGATGGAGGG  
*Elionurusmuticus6954* CTCATTC-GGTAGGTGTT-----ACCTATCCTGATGC-GAGGA  
*Icylindrical* T--CTCGATTATTAATTCGATTTTAAAGTATTATTA-GTAAGCC  
*Sacharumofficinatum* T--CTCGATTATTAATTCGATTTTAAAGTATTATTA-GTAAGCC  
*Sorghumhalepense* T--CTCGATTATTAATTCGATTTTAAAGTATTATTA-GTAAGCC



*Ttriandra* T--CTCGATTATTAATTCTATTTTAAAGTATTATTAA-GTAAGCC  
*Pschinzii* ???

*Aadscen6713* ???  
*Aameth6679* -GTAAGG-A-TTAAA--ACTTACCTTATGA--AT-AACTAAACAA--TC  
*Aameth6685* -GTAAGG-A-TTACA--TCTTACCTTATCA-CAT-TACTAAACAA--TC  
*Aappen6543* ???  
*Aappen6677* -GTAAGG--ATATAAC---CTTATCTTATGC-CAT-ACCTCAACA--AGG  
*Aappen6946* -GTAAGG-ATTAAAA--CTTTTCTTACCGA-AAT-AACTAAAA--AATC  
*Adist6684* -GTAAGG-AATTAAA--ACTTACCTTATGA-AAT-AAATAAACAA--TC  
*Aeuco5869* -GTAAGG-CATCTTAA---CTTATCTTGTGA-AAT-TCCTAATCA--GGC  
*Aschiren6581* -GTAACA-ATTAAAA---CTTATCTTACGA-ACT-AACTAAACAA--TC  
*Bbladh* -CTCAATT-ATTTCATC---TT---CTTTCCCTTTT-ACCTAAACAA--T-  
*Binsc5681* -GGAAAGG-ATTAAAA---CTCACCTTATGA-AAT-AACTAAAA--GC  
*Brachiaria* ???  
*Bserra5074* -GTATAGG-AGTAAAA--CTTATCTTAC-GA-AAT-AACTAAAA--AATC  
*Bserra6654* ???  
*Bserra6954* -GTATAGG-ATTAAAA--CTTTTCTTAC-GA-AAT-AACTAAAA--AATC  
*Cdieter6642* -GTAAGG-CATCTTAA---CTTACCTTACGA-AAT-AACTAAACA--ATC  
*Cexc5877* ???  
*Cexc6577* -TTAAACCAATTCAAAA---CTCATCTTACGC-AATTAACATAACAT-AGC  
*Cmarg4397* ???  
*Cpilg4606* -GTAACA-ATTAAAA---CTTATCTTACGC-AAT-AACTCAAAATAGCA  
*Cprolix6682* -GGAAAC-AATTAAAA---CTCACCTTACGA-AAT-AACTAAACA--ATC  
*Dfilli5872* -GTAATG-GTTTTAAA--CCTTA-----TGA-AAT-AACTAAAA--AATT  
*Hanam4807* --TAAAGG-ATTTCAGG---CTTAACTCTTATGTAT-AAATAAAAA--ATC  
*Hcon6942* -TTACTC-TATCTCGC---CTTATCTTCCGA-ATT-GTCTAAACA--AGG  
*Hdiss5145a* GGTAAGG-GTTTTAAA--CCTTACCTTATGA-AAT-AACTAAAT-AATC  
*Sbicolour9319* -GTATACG-ATTAAAA---CTTATCTTATGA-AAT-AACTAAACA--ATC  
*Hemarthriaaltissima5698* -TTAAAA-CTTACGCTTAAATTTCTAACG-TAT-AAATAAAAA--TC  
*Hfilli5144* --TATAAG-ATTTCAGA---CCTTACCTTATA-AAT-AAATAAAAA--AATC  
*Hhirt6650* ???  
*Hhirt6656f* -GTAACA-ATTAAAA---CTTATCTTATGA-AAT-AACTAATCAG--TC  
*Hlongi6560* ???  
*Hpilg4603* -GPTAAAT-CTTAAAC-TTATGAAATAAAT---T-AACTAAAA--TC  
*Hpilg4738* --TGATG-CTTCAGG--CCATACCTTTAGG-AAT-AACTAAAA--AATC  
*Hquar5088* -----TTC-CTTAGAG--CCTTT-----TATT-AAT-G-CTAAAA--CTTC  
*Hrud6576* -GGAAACA-ATTAAAA---CTTATCTTACGA-AGT-AACTAAACAT-AGC  
*Htamb6585* -GTACAGA-ATTCAAAA---CTTATCTTATGA-AAT-AACTAAACA--GC  
*Icylin5114a* -GTAACA-ATTAAAA---CTTACCTTATGA-AAT-AACTAATCAA--GC  
*Icylin6632a* -GTAACA-ATTCAAC---CTCATCTTACGA-AAT-AACTAAACA--ATC  
*Mcape6655* -TTATACA-ATTCAAAA---CTCATCTTACGCTAT-CACTACACAA--GC  
*Pdilatum6617* -TTATTCAAGTATAG-TCATTCCTTAC---ATTTCCCTAAGTTGCAAT  
*Pfrank4705* --GTAAG-GTTCAAAA--TCTTACCTT-AGA-AAT-AACTAAAA--AATC  
*Pschinz6652* ???  
*Sbic6588* -GTAAGG-ATTAAAA---CTTACCTTATGA-AAT-AACTAAACA--ATC  
*Shalep9318* ---ATACA-GTTCAAAA---CCCTACCTTATGA-AAT-AACTAAAA--AATC  
*Snigros6567* ???  
*Tspic4846* --TGTAAG-GTTCAGG--TCTTACCTTTTGG-AAT-AAATAAAA--AATC  
*Tspic6580* -GGAAACA-ATTAAAA---CTCATCTTACGA-AAT-AACTAAACA--ATC  
*Ttria6383 b* -GTAACA-AGTAAAC---CTCATCTTATGA-AAT-AACTAATCAG--TC  
*Ttria6555* -GTAATA-TTTAAAA---ATCTACTTAGTC-TTT-AGTCTAGAGT--AA  
*Zea* -GTATAGG-ATTAAAA--CTTTTCTTAC-GA-AAT-AACTAAAA--AATC  
*Elionurusmuticus6954* -ATCCACA-ATGCATAGGACT-ACCCC-TCCCCATTTCCCTAATTTGGAAT  
*Icylindrical* -ATCCACA-ATGCATAGGACT-ACCCC-TCCCCATTTCCCTAATTTGGAAT  
*Sacharumofficinatum* -ATCCACA-ATGCATAGGACT-ACCCC-TCCCCATTTCCCTAATTTGGAAT  
*Sorghumhalepense* -ATCCACA-ATGCATAGGACT-ACCCC-TCCCCATTTCCCTAATTTGGAAT  
*Ttriandra* -ATCCACA-ATGCATAGGACT-ACCCC-TCCCCATTTCCCTAATTTGGAAT  
*Pschinzii* ???

*Aadscen6713* ???  
*Aameth6679* AGGGA-AATTAACGT-TAT--CTTCGTTACCG-AGATGATT-CTACTA-C  
*Aameth6685* AGGGA-AATTAGCTC-TAT--CTTCGTTACCG-AGATGATT-CTACTA-C  
*Aappen6543* ???  
*Aappen6677* AGCGGGCAGTAGCTTATAT--CTTTGTCC-AC-CGCGG-ACACGTCTA-T  
*Aappen6946* AGGGA-AATT-GACTGTAT--GTTCTGTT-ACGAAAGGATACCTACTG-C  
*Adist6684* AGGGA-AATTAACGT-TAT--CTTCGTTACCGAGATGATA-CTACTA-C  
*Aeuco5869* AGGGA-AACTGTCT-GTAT--CTTCGTT-ATGAGATG-ATTCTACTA-C  
*Aschiren6581* AGGGA-AACTAACT-GTAT--CTTCGTTTAC-CAGATGATA-CTACTA-C  
*Bbladh* AGAGG-AGTAATTA-TAT--CTTTATACTTGT--ATGATT-CTATTA-T

*Binsc5681* AGGGA-GATTGACTG-TAT--CTTCGTTCAAG-AGATGATA-CTACTA-C  
*Brachiaria* ???  
*Bserra5074* AGGGA-AATT-TACTGTAT--GTTTCGTTT-CCGAGATGACAACACTACTA-C  
*Bserra6654* ???  
*Bserra6954* AGGGA-AACT-GACTGTAT--GTACGTTT-ATGAGATGATC-CTACTA-C  
*Cdieter6642* AGGGAG-ATTAAC-TGAT--CTACGTTTACCGAGATGGATACTACTA-C  
*Cexc5877* ???  
*Cexc6577* ACTTGAAATATCT-CTATT-CTTCGTTT-ACCAAATG-ATCCTACTA-C  
*Cmarg4397* ???  
*Cplur6406* CGGGG-CCTTGG-GCGCATATCTTCTTAG-ACCAATGGACA-CTTTTA-C  
*Cprolix6682* AGGGAGGATGACT-GTAT--CTACGTTT-CCGAGATG-ATACTACTA-C  
*Dfilli5872* AGGGA-TGTTAACGGTCT--CTTCGTTT-ACGATTGGCAC-CTTTTGGC  
*Hanam4807* AGGGA-AATTAAC-TGAT--CTACGTTT-ATGAGATGATA-CTACTA-C  
*Hcon6942* AGCGA-CTGTGCT-ATA---CCTTGTCT-TCGCGCGG-ATTCTNCTN-N  
*Hdiss5145a* ACCCA-AATTCACGGTCT--TTATTCCT-ACCGTTTGCCA-CTTTTGTG  
*Sbicolour9319* AGGGA-AATTGGACTGTAT--CTTCGTTT-ACCAAAGGATA-CTACTA-C  
*Hemarthriaaltissima5698* AGGGA-AATTAAC-TGAT--CTTCGTTTACCGAGAGGACAACACTACTT-C  
*Hfilli5144* AGCGA-AATTGACCTAT---CTTTTCCC-ACCATGTGCCA-CTTTTGTG  
*Hhirt6650* ???  
*Hhirt6656f* AGGGA-AACTGTCT-CTAT--CTTCGTTTACCGAGATGATC-CTACTA-C  
*Hlongi6560* ???  
*Hpilg4603* AGGGA-AATTAACGGGTAT--CTAGGTTTATGA-GAGGACA-CTTTTA-C  
*Hpilg4738* CGGGA-AATT-AACTGTAT--CTACGTTT-ATGAGATGATC-CTACTG-C  
*Hqar5088* AGGGA-AATT-AACTGTAT--CTACGTTT-ATGAGATGATC-CTACTA-C  
*Hrud6576* AGGGA-AACTGTCTG-TAT--CTTCGTTT-CCAAATGATA-CTACTA-C  
*Htamb6585* ATGGA-AACTAATG-TAT--CTTCGTTT-ACGATATGATA-CTCTTA-C  
*Icylin5114a* AGGGA-AATTAACGC-TAT--CTTCGTTTACAG-ATAGGACA-CTACTG-C  
*Icylin6632a* AGGGA-AATTGA-CTGTAT--CTTCGTTT-CCAAATGATAACTACTA-C  
*Mcape6655* AGTTG-GACCTGCC-CTAT--CTTCGTTTATC-TATTGTCC-CTACTA-C  
*Pdilatum6617* AGAGA--ATGCATATTTATTTTTAGTCCCTTTAATTGACA--TAGAAGC  
*Pfrank4705* AGGGA-AATT-AGCGGTAT--CTACGTTT-ATGAGAGGACC-CTACTG-C  
*Pschinz6652* ???  
*Sbic6588* AGGGA-AATTAG-CTGTAT--CTACGTTT-ATGAGAGGATT-CTACTA-C  
*Shalep9318* AGGGA-AATGTAACATATAT-CTTTGCC-ATAAGAGGAC-CTACTAC  
*Snigros6567* ???  
*Tspic4846* TSGGA-AATT-AACGGTTT--CTTCGTTT-ATGAGATGATC-CTACTG-C  
*Tspic6580* AGGGA-AATTGA-CTCTAT--CTTCGTTT-ACCAGATGATC-CTACTA-C  
*Ttria6383 b* AGGGA-A-CTGTCT-GTAT--CTTCGTTT-CAT-GAGATGATC-CTACTA-C  
*Ttria6555* AGGGA-A-CTGTCT-GTAT--CTTCGTTT-CAT-GAGATGATC-CTACTA-C  
*Zea* GTAGA-AAATCTCT-AGAG-TAAGGATAAATC---TCTAC-ACACTT-C  
*Elionurusmuticus6954* AGGGA-AACT-GACTGTAT--GTACGTTT-ATGAGATGATC-CTACTA-C  
*Icylindrical* -GG-AATACTTTATTG-ATTTTTTAGTCCCTTTAATTGACA--TAGATGC  
*Sacharumofficinatum* -GG-AATACTTTATTG-ATTTTTTAGTCCCTTTAATTGACA--TAGATGC  
*Sorghumhalepense* -GG-AATACTTTATTG-ATTTTTTAGTCCCTTTAATTGACA--TAGATGC  
*Ttriandra* -GG-AATACTTTATTG-ATTTTTTAGTCCCTTTAATTGACA--TAGATGC  
*Pschinzii* ???

*Aadscen6713* ??????????????  
*Aameth6679* GCGTCCTTTCCA  
*Aameth6685* GCGACCTTTGTCA  
*Aappen6543* ??????????????  
*Aappen6677* GTGTCCTTTTCA  
*Aappen6946* GTGTCCTTTGCCA  
*Adist6684* GTGTCCTTTGCCA  
*Aeuco5869* GTGTCCTTTCCCA  
*Aschiren6581* GTGTCCTTTCCCA  
*Bbladh* ATGATCTTTGTCA  
*Binsc5681* GCGTCTTTGTCA  
*Brachiaria* ??????????????  
*Bserra5074* GCGTCTTTGCCA  
*Bserra6654* ??????????????  
*Bserra6954* GTGTCCTTTGCC  
*Cdieter6642* GCGTCTTTCCCA  
*Cexc5877* ??????????????  
*Cexc6577* GCGTCTTTCCCA  
*Cmarg4397* ??????????????  
*Cplur6406* CGGACCTTTGGTCA  
*Cprolix6682* G-GTTCCTTTCCCA  
*Dfilli5872* GCG-CCTTGGTCA  
*Hanam4807* GTGTCCTTTGCCA  
*Hcon6942* GTGTCCTTTCCCA

Hdi55145a	GCA-CCTTTGTCA
<i>Sbicolour</i> 9319	GTGTTCTTTCTCA
<i>Hemarthriaaltissima</i> 5698	GTGACTTTTGTC
<i>Hfilli</i> 5144	GTAACTATGTCC
<i>Hhirt</i> 6650	????????????
<i>Hhirt</i> 6656f	GTGTTCTTTGCCA
<i>Hlongi</i> 6560	????????????
<i>Hpilg</i> 4603	GCGTCTTTGCCN
<i>Hpilg</i> 4738	GTGTTCTTTGCCA
<i>Hquar</i> 5088	GTGTTCTTTCCCA
<i>Hrud</i> 6576	GTGTCCTTTGTCA
<i>Htamb</i> 6585	GTGTTCTTTGTCA
<i>Icylin</i> 5114a	GCGTCCTTTGCCA
<i>Icylin</i> 6632a	GCGACCTTTGTCA
<i>Mcape</i> 6655	G-GTCTTTCCCC
<i>Pdilatum</i> 6617	AGTACTCTACTAA
<i>Pfrank</i> 4705	GTGTCCTTTGCCA
<i>Pschinz</i> 6652	????????????
<i>Sbic</i> 6588	GTGTTCTTTCCCA
<i>Shalep</i> 9318	GGGTCCTTTCTCC
<i>Snigros</i> 6567	????????????
<i>Tspic</i> 4846	GTGTCCTTTGCCA
<i>Tspic</i> 6580	GGGACCTTTTCA
<i>Ttria</i> 6383 b	GTGTTCTTTCCCA
<i>Ttria</i> 6555	GTGTTCTTTGCCA
<i>Zea</i>	TTTTCTTTTAAA
<i>Elionurusmuticus</i> 6954	GTGTTCTTTGCCC
<i>Icylindrical</i>	AAATACTTTCTAA
<i>Sacharumofficinarum</i>	AAATACTTTCTAA
<i>Sorghumhalepense</i>	AAATACTTTCTAA
<i>Ttriandra</i>	AAATACTTTCTAA
<i>P.schinzii</i>	????????????

**Appendix E:** CLUSTAL W (1.74) multiple sequence alignment data of the Andropogoneae specimens for *ITS1* region.

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Aameth6685      TCGTGACC-GCGTCGATGACCG-T--AACAAAGT-TTCCCTCAGTT--G
Aameth6679      TCGTGACCCGCGTGAAAAATATG-AC-AACAAAGGCTTCCCTCAGTT--G
Dfili5874       TCGTGACCCG-TGAAAAAACC-CA-AACAAGAGCTTCCCTCAGTT--C
Ttria6555       TCGTGACC-GT-GAAGGTAACG-CC-AACAAGAGTTTCCCTCAGTT--G
Ttria6525       TCGTGACC-CAG-GAAGGAATT-CG-TAACAAAGCTTCCCTCAGTT--G
Tspic5205       TCGTGACA-----ATAAACGA-AT-AACAAACGTTTCCCTCTCTT--C
Tspic6579       TCGTGACA--CGG-AGGAATTC-GT-AACAAA-GTTTCCCTCAGTT--G
Tspic6580       TCG--ACG----AAAGGTTATC-GT-AACAAA-GTTTCCCTCAGTT--G
Tspic5873       TCGTGACC--CGGAGGAAATTC-GT-AACAAA-GCTTCCCTCTGTT--G
Adist6684       TCGTGACA-----CGAAAGAAA-C-GT-AACAAACGCTTCCCTCAGTT--C
Mcape6584       TCGTGAG----CGATCGTAC-C-GT-AACAAA-GCTTCCCTCNGTT--C
Aappen6543      TCGTGAC-----GAAGGAATTC-GT-AACAAA-GTTTCCCTCAGTT--N
Mcape6655      TCGTGAGG---CGAAGGATT-C-GT-AACAAG-GTTTCCCTCAGTT--G
Aappen6946      TCGTGACC-----GAGGAATTC-GT-AACAAA-GTTTCCCTCAGTG--C
Sbic6589        TCGTGACC--CAGGAGGAATTC-GT-AACACA-GTTTCCCTCAGTT--C
Cplur6406       TCGTGACC--ATGTAGATATTC-GT-AACAAA-GCTTCCCTCAGTT--C
Icylin6632      TCGTGACC--G-GAAGA-ATTC-GT-AACAAA-GCTTCCCTCAGTT--C
Hpilg4738       TCGTGACC--CGGAAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--C
Icylin5090      TCGTGAGC--CGGAAGG-ATTC-GT-AACAAA-GCTTCCCTCAGTT--G
Emuti6715       TCGTGAGC-----GAGGAATTC-GT-AACAAA-GCTTCCCTCAGT--G
Emuti6559       TCGTGAGG-----GAGGAATTC-GT-AACAAA-GCTTCCCTCCGT--G
Emuti5074       TCGTGACC----GGAGGAATTC-GT-AACAAA-GTTTCCCTCAGT--G
Emuti6539       TCGTGAACA--CGGAGGAATTG-CT-AACAAA-GTTTCCCTCAGT--G
Pfrank4705      TCGTGAC-----GAGGAATTC-GT-AACAAA-GTTTCCCTCAGTT--G
Icylin5114      TCGTGACCT---GAAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G
Aeuco5869       TCGTGACCCG---GAAA-AATCC-GT-AACAAG-GCTTCCCTCATGCATG
Dfili5872       TCGTGACCGA--GGA---ATTC-GT-AACAAG--CTTCCCTCAGTT--G
Aeuco6587       TCGTGACCCCTT--GGAAGGATTC-GT-AACAAA-GCTTCCCTCACT--G
Bradi4784       TCGTGAACCT--GAAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G
Binsic6581      TCGTGACC---GGAGGAATTC-GT-AACAAA-GCTTCCCTCAGT---
Tspic4846       TCGTGAT----GGAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G
Cprolix6682     TCGTGACCCG---TTAAACCG--GT-AACAAA-GCTTCCCTCCCT--G
Csp5800         TCGTGACCCG---CGGAATT-GT-AACAA--GCTTCCCTCAGT---
Cprolix5101     TCGTGACCCG---AGGAAATC-GT-AACAAA-GCTTCCCTCACT--G
Cdieter6642    TCGTGACCCCG--ACGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G
Cmarg4397       TCGTGACAATTGAAAGGAAATCG-GT-AACAAGAGCTTCCCTCAGTT--
Hpilg4603       TCGTGAGCGG---AGGAAATTC-GT-AACAAA-GTTTCCCTCCGT--G
Htamb6585       TCGTGACC---CGGAGGAAATCCGT-AACAAA-GCTTCCCTCTGAT--G
Hdiss5145       TCGGTACT---TGGAGGAATTC-GT-AACAAAG-CTTCCCTCTGAT--G
Htamb6537       TCG-----NGAAAGAAATTCGT-AACAAAG-CTTCCCTCTGTT--G
Hanam4807      TCGTGAGC---CGGAGGAATTC-GT-AACAAAG-TTTCCCTCTCAT--G
Hfilli5144     TCGTGATAG---CGGAGGAATTC-GT-AACAAAG-CTTCCCTCTCAT--G
Hanam4841       TCGTGACC---CGAGGAAATC-GT-AACAAAG-CTTCCCTCTCAT--G
Hhirt6656       TCGTGACG---AGGAAATTC-GT-AACAAAG-CTTCCCTCAGTT--G
Hhirt6618      TCGTGACC---CGGAGGAATTC-GT-AACAAANGCTTCCCTCAGTT--G
Shalep9318     TCGTGACCC---CGAAGGAAATC-GT-AACAAAG-CTTCCCTCTCAT--G
Hcon4741       TCGTGACCC--TGAAGGAAATC-GA-AACAAA-GCTTCCCTCTGTT--G
Evill5085      TCGTGACCC--TGAAGGAAATCAG--ACCGAAAGCTTCCCTCTAAT--G
Bbladh5173     TCGTACCC---TGAAGGAAACAGA-AACAAA-GCGTCCCTCTCAT--G
Hquar5088      TCGTGACC---TTAAACAAAACAGACCGTGAACATGTCTCTCATGT---
Sbic6588       TCGTGACCC--TTAAACAAAACAGACCGTGAACATGTCTCTCATGT---
Sbicolour9319  TCGTGACCC--TTAAACAAAACAGACCGCGAACGCGTCTCTCATGT---
Hrud6576       TCGTGACCC--TTAAACAAAACAGACCGTGAACGCGTCCCTCATGT---
Andropogondistachyos
Cleistachnesorghoides
CoixBuckler
Arundinellanepalensis
Pennisetumsetaceum
Panicumbisulcatum
Danthoniacalifornica
PentaschistisaristifoliaSpies6
Eragrostisdielsii
Aristidaharvardii
Zeadiplo
ZmaysGfl

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*Evillosa*  
*Saccharumofficinatum*  
*Imperatocylindrical*  
*Bothriochloainsculpta*  
*Dichanthiumannulatum*  
*Cplurnodus*  
*Cexcavatus*  
*Hyparrhrietamba*  
*Hanamesa*  
*Heteropogoncontortus*

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TCGTGACC---TTAAACAAAACAGACCGCGAACGCGTCCCTCGTG--CC  
TCGTGACCC--TTAAACAAAACAGACCGCGAACGAGTCTCTCGTG--CC  
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*Ttria6525*  
*Tspic5205*  
*Tspic6579*  
*Tspic6580*  
*Tspic5873*  
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*Binsc5681*  
*Tspic4846*  
*Cprolix6682*  
*Csp5800*  
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*Cdieter6642*  
*Cmarg4397*  
*Hpilg4603*  
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*Hanam4807*  
*Hfilli5144*  
*Hanam4841*  
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*Hcon4741*  
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*Bblad5173*  
*Hquar5088*  
*Sbic6588*  
*Sbicolour9319*  
*Hrud6576*  
*Andropogondistachyos*  
*Cleistachnesorghoides*  
*CoixBuckler*  
*Arundinellaneapalensis*  
*Pennisetumsetaceum*  
*Panicumbisulcatum*

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*PentasthisaristifoliaSpies6* CGGCCACACGA-----GCTCACGCCGTGTGGC-CTAGGCCGCCGAC  
*Eragrostisdielsii* CGTCCGG-GTGATGGGGTTG-CCTCGCTCCCGAC-TTAGGCCCCAGAC  
*Aristidaharvardii* CGTCCGG-GCGTC-----TCCCACGACGCCCGC-CAAGGCCCCGAC  
*Zeadiplo* CGCCGG-GCTCC-----GGCCCGGCACGCTGCCCCGCC-  
*ZmaysGfl* CGCCGG-GCTCC-----GGCCCGGCACGCTGCCCCGCC-  
*Evillosa* -GCCGG-GCTTC-----GGCTCGGC-CAAGGCACCYGAG  
*Saccharumofficinatum* -GCCGG-GCTTC-----GGCCCGGCACGAGGTCCCGGAG  
*Imperatocylindrical* -GCCGG-GCTCC-----GGCCCGGC-CAAGGCCCCGAG  
*Bothriochloainsculpta* -GACGG-GCTCC-----GGATCGGA-CAAGGCCCCGAG  
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*Cplurnodus* -GCCGG-GCTTC-----GGCTCGGA-CAAGGCCTCCGAA  
*Cexcavatus* -GCCGG-GCCTC-----GGCCTGAGCAAGGTCCCGAA  
*Hyparrhrietamba* -GCCGG-GCTCC-----GGCTCGGC-CAAGGCCTCCGAA  
*Hanamesa* -GTCGG-GCTCC-----GGTCCGGC-CAAGGCCTCCGAA  
*Heteropogoncontortus* -GTCGA-GCCTC-----GGCTCGGC--AAGGCCCCCGAT

*Aameth6685* -CCTCGTCCC-----GGGGCGGAGGTGCCGC-AAAAGAACCACGGC  
*Aameth6679* -CCCCGTCCC-----GGGGCGGAGGTGCCGC-AAAAGAACCACGGC  
*Dfili5874* -CCCCGTCCC-----GGGGCGGAGGGGCCGC-AAAAGAACCACGGC  
*Ttria6555* -CTCCGTCCC-----GGGGCGGAGGGGCCAC-AAAAGAACCACGGC  
*Ttria6525* -CTCCGTCCC-----GGGGCGGAGGGGCCAC-AAAAGAACCACGGC  
*Tspic5205* -CTCCGTCCC-----GGGGCGGAGGGGCCAC-AAAAGAACCACGGC  
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*Tspic5873* -CTCCGTCCC-----GGGGCGGAGGGGCCAC-AAAAGAACCACGGC  
*Adist6684* -TCCCTCCC-----TGGTAATCGCCGAA-AA-ACAACCCACGGC  
*Mcape6584* -CCCCGTCCG-----GGGGCGGACGGGCCAC-AACACAACCACGGT  
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*Cplur6406* -CCCCNTCTT-----GGGGTGGAGGGGCCGC-AAAATAACCACGGC  
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*Hpilg4738* -CCCCGTCCAC-----GGGGCGGAGGGGCCGC-AAAAGAACCACGGC  
*Icylin5090* -CTCCGTCCC-----GGGGCGGAGGGGCCGC-AACAGAACCACGGC  
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*Emuti6559* -CTCCGGA-----GAGGGGCCGC-AACAGAACCACGGC  
*Emuti5074* -CTCCGGA-----GAGGGGCCGC-AACAGAACCACGGC  
*Emuti6539* -CTCCGGA-----CAAGGGCCGC-AACAGAACCACGGC  
*Pfrank4705* -CACCGTCAA-----GGGGC-GAGGGGCCGC-AACAGAACCACGGC  
*Icylin5114* -CTCCGTCCC-----GGGGCGGAGGGGCCGC-AACAGAACCACGGC  
*Aeuco5869* -CCCCGTCCC-----GGGGCGGAGGGGCCGC-AAAAGAACCACGGC  
*Dfili5872* -CCCCGTCCA-----GGGGCGGAGGGGCCGC-AAAAGAACCACGGC  
*Aeuco6587* -CCCCGTCCA-----GGGGCGGAGGGGCCGC-AAAAGAACCACGGC  
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*Binsc5681* -CTCCGTCTT-----GGGGCGGAGGGGCCGC-AAAAGAACCACGGC  
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*Hhirt6656* -CCCCGTCCG-----GGGGCGGAGGTGCCGC-AAAAGAACCACGGC  
*Hhirt6618* -CCCCGTCCC-----GGGGCGGGGGCCAC-AAAACAACCACAGG-C  
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*Hcon4741* -CTTC-----GGCAGAGGGGCCGC-CACAGAACCACGGC  
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*Bblad5173* -CCCCGTCCC-----GGGGCGGAGGAGCCGC-CAAAGAACCACGGC  
*Hquar5088* -CCCCGTCCC-----GGGGCGGAGGTGCCAGCCAAAAGAACCACAAAGC  
*Sbic6588* -CTCCAAC-CT-----GGGGCAGAGGGGCCAC-AAAAGAACCACGGC  
*Sbicolour9319* -CTCCAAC-CT-----GGGGCAGAGGGGCCAC-AAAAGAACCACGGC  
*Hrud6576* -CCCCGTCTCTCATGTGCGGGCGGAGGGGCCGC-AAAAGAACCACGGC

*Andropogondistachyos* -CCCCGTC-CC-----GGGNGGAGGTGCCGCCAAAAGAACCCACGGC  
*Cleistachnesorghoides* -CCNNGTCNT-----GGGGCGGAGGGGCCAC-AAAAGAACCCACGGC  
*CoixBuckler* -CTCCGTCCC-----GGGGCGGAGGGGCCGC-AACAGAACCCACGGC  
*Arundinellaneपालensis* -CTCCGTCCC-----GGGGCAGAGGGGCCCTT-AACAGAACCCACGGC  
*Pennisetumsetaceum* -TTCTTTAT-----GAGGGGGA--GGGCCGC-AAAAGAACCCACGGC  
*Panicumbisulcatum* -CACCT-AT-----TAGGAGGATGGGGCCGC-AAAAGAACCCACGGC  
*Danthoniacalifornica* -CTCCCTCCTCC----GGGAGGGGAGCGGCCGT-AAAAGAACCCACGGC  
*PentastichisaristifoliaSpies6* -CTCCGCAA-----GGACGGGAGCGGCCGC-AAAAGAACCCACGGC  
*Eragrostisdielsii* -TTTCTTAC-----GGAGGGCTGGGGCCGC-AACAGAATCCACGGC  
*Aristidaharvardii* -CTCCCTGT-----GGAGGGGAGGGGCCGC-AACAGAACCCACGGC  
*Zeadipl* -GAACCTCCCGC----GGGGAAGGGGGGCCGCAAAAAGAACCCACGGC  
*ZmaysGfl* -GAACCTCCCGC----GGGGAAGGGGGGCCGCAAAAAGAACCCACGGC  
*Evillosa* TCCAGTCCC-----GGGCGCGAGGGTGCGCGCAAAAAGAACCCAYGGC  
*Saccharumofficinarium* CTCCGTCCC-----GGAGCGGAGGGGCCAC--AACAGAACCCACGGC  
*Imperatacylindrical* CTCCGTCCC-----GGGCGGAGGGGCCGC--AACAGAACCCACGGC  
*Bothriochloainsculpta* CTCCGTCT-----GGGGGGGGGGA-GTCGCAAAAAGAACCCACGGC  
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*Cexcavatus* CCCCTTCCC-----GGGGGGGAGG-TTCCCC-AAAAGAACCCACGGC  
*Hyparrheniatamba* CCCCTTCCC-----GGGGGGGAGG-TTCCCCCAAAGAACCCACGGC  
*Hanamesa* CCCCTTCCC-----GGGGGGGAGG-TTCCCCCAAAGNAACC-ACGGC  
*Heteropogoncontortus* CTTCGA-CA-----GAGGGG-----CCACCACAGAACCCACGGC

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Aameth6679 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Dfili5874 GCCTT---AGGCGTCAAGGAACACATG-TATTGCCTT--GC  
Ttria6555 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Ttria6525 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Tspic5205 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCAT--GC  
Tspic6579 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCAT--GC  
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Adist6684 GCCTTTCGTAAACATTCAGGAACACTTA-TATTGCCTT--GC  
Mcape6584 GCCTT---ATGCGTCAAGGAACACTTA-TATTGCCTT--GC  
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Mcape6655 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
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Sbic6589 GCCTT---AGGCGTCAAGGAACACTCA-TGTTGCCTT--GC  
Cplur6406 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Icylin6632 GCCTA---AGGCGTCAAGGAACACTTC-TATTGCCTT--GC  
Hpilg4738 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTC--GC  
Icylin5090 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Emuti6715 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTC--GC  
Emuti6559 GCCTT---AGGCGTCAAGGAACACTTA-TATCGGCTC--GC  
Emuti5074 GCCTT---AGGCGTCAAGGAACACTTA-TATTGGCTC--GC  
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Pfrank4705 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Icylin5114 GCCTA---AGGCGTCAAGGAACACTTC-TATTGCCTT--GC  
Aeuco5869 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Dfili5872 GCCTT---AGGCGTCAAGGAACACTTG-TATTGCCTT--GC  
Aeuco6587 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Bradi4784 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Binsc5681 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Tspic4846 GCCTT---AGGCGTCAAGGAACACTG-TATTGCCTT--GC  
Cprolix6682 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Csp5800 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Cprolix5101 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Cdieter6642 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Cmarg4397 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Hpilg4603 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Htamb6585 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Hdiss5145 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Htamb6537 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Hanam4807 GCCTT---AGGCGTCAAGGAACACTTAATATTGCCTT--GC  
Hfilli5144 GCCTT---AGCC-TCA-GGAACACTTA-TATTGCCTT--GC  
Hanam4841 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Hhirt6656 GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
Hhirt6618 GCCTT---AGGCGTCAAGGAACACTTGT-TATTGCCTT--GC  
Shalep9318 GCCGA---AGGCGTCAAGGAACACTGA-TATTGCTTT--GC  
Hcon4741 GCCTT---AGGCGTCAAGGAACACTTG-TATTG-CCTT--GC

Evill5085  
Bbladh5173  
Hquar5088  
Sbic6588  
Sbicolour9319  
Hrud6576  
Andropogondistachyos  
Cleistachnesorghoides  
CoixBuckler  
Arundinellanepalensis  
Pennisetumsetaceum  
Panicumbisulcatum  
Danthoniacalifornica  
PentaschistisaristifoliaSpies6  
Eragrostisdielsii  
Aristidaharvardii  
Zeadiplo  
ZmaysGfl  
Eviolosa  
Saccharumofficinatum  
Imperatacylindrical  
Bothriochloainsculpta  
Dichanthiumannulatum  
Cplurnodus  
Cexcavatus  
Hyparrhrietamba  
Hanamesa  
Heteropogoncontortus

GCCTT---ACGAGTCAAGGAA-ACTTG-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTAC-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTCAATATTGCCTT--GC  
GC-TT---AGGCGTCAAGGAACACTCA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTCA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTA---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTAA-TGTTGCCTT--GC  
GCCAT---AGGCGTCAAGGAACACTCA-TATTGCCTT--GC  
GCCGA---AGGCGTCAAGGAACACTGA-TATTGCCTT--GC  
GCCAA---AGGCGTCAAGGAACACTA-TATTGCCTT--GC  
GCCGAC---CGGCGCAAGGAACACTGTTATTGCCTT--GC  
GCCGTA---CGGCGTCAAGGAAAACACTGT-TATTGCCTT--GC  
GCCGA---CGGCGTCAAGGAACACTAA-TATTGCCTT--GC  
GCCGAA---CGGCGTCAAGGAACACTGTCTATTGCCTT--GC  
GCCTC---GGGCGCAAGGAACACCAG-TACTACCTCCTGC  
GCCCC---GGGCGCAAGGAACACCAG-TACTACCTCCTGC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTTGCCTC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTTGCCTC  
GCCT---AGGGGTCAAGGAACACTTC-TATTGCCTTGCCTC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTTGCCTC  
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GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTTGCCTC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTTGCCTC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTTGCCTC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTTGCCTC

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Aameth6679  
Dfili5874  
Ttria6555  
Ttria6525  
Tspic5205  
Tspic6579  
Tspic6580  
Tspic5873  
Adist6684  
Mcape6584  
Aappen6543  
Mcape6655  
Aappen6946  
Sbic6589  
Cplur6406  
Icylin6632  
Hpilg4738  
Icylin5090  
Emuti6715  
Emuti6559  
Emuti5074  
Emuti6539  
Pfrank4705  
Icylin5114  
Aeuco5869  
Dfili5872  
Aeuco6587  
Bradi4784  
Binsc5681  
Tspic4846  
Cprolix6682  
Csp5800  
Cprolix5101  
Cdieter6642  
Cmarg4397  
Hpilg4603  
Htamb6585  
Hdi58145  
Htamb6537  
Hanam4807

ACGGCGGGCGGCTGGC-TTGCTTCCGCTCCCCGCGCAGC---GATCAT  
ACGGCGGGCGGCTGGC-TTGCTTCCGCTCCCCGCGCAGC---GATCAT  
CCGGCGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
TCGGCGGAGCGGTGGC-TTGCTTCCGCTCCCCGCGCAGC---GATGAC  
TCGGCGGAGCGGTGGC-TTGCTTCCGCTCCCCGCGCAGC---GATGAT  
TCTGTGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
TCGGCGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
TCTGTGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
TCGGCGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
ACTTCCGAGCGGTGGC-TTGCTTCCCTCCCCGCGCAGC---GATCAT  
TCGGCGG-GACGCTGC-CTGCCTTCCGCTCCCCGCGCAGC---ATATGAT  
TCGGCGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
TCG-CGG-GCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
ACA-CGGAGTGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
CCGGTGA-CGCTC-C-CTGCCTTCCACTCCCCCGCA-C---GATGAT  
TCGGCGGAGCGGTGGC-C-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
CCGGCGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
CCGGCGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
TCGGCGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
TCGGCGGAGCGGTGGC-ATGCCTTCTGCTCCCCGCGCAGC---GATGAT  
TCGGCGGGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
TCGGCGGGGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
CCGGTGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC-C---GATGAT  
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GCCGTGGAGCGTCAACTCTTCCCTCC-CTCCCCGCGCAGC---GATGAT  
CCGGTGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
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CCGGTGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
AAGGTGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
CAGGTGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
CCGTGGGAGCGGTGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT



*Hfilli5144* CCGTGGAA--CGGTCCGGC-CTGCCTTTCGCTCCCCGTGCAGC---GATGAT  
*Hanam4841* CCGGTGGAGCGGTGCAGC-TTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Hhirt6656* CCGGTGGAGCGGTCTGC-TTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Hhirt6618* CAGGTGTACTCCTCCGGC-CTGCCTTCCGCTCCCCGTGCAAC---GATGAT  
*Shalep9318* TCGGGGGTTTGGTTGGC-TTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Hcon4741* TCGGTGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Evill5085* TCGGTGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Bblad5173* CCGGCGGAGCGGTCCNNC-CTGCCTTCCGCTCCCCG-GCAGC---GATGAT  
*Hquar5088* CCGGTGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Sbic6588* GCAGTGGAG-GCTCCGGC-CTCCGTTGGTGCCGCGATGAAGAACGCAAGAT  
*Sbicolour9319* ACAGTGGAGTGGTCCGGC-NTGCGTTGGGGTCCCGATGAAGAACAAAAGAT  
*Hrud6576* AAGGTTGGAACCAAGGC-TTCCGTGGGGGCACCGATGAAGA-CGGATGAT  
*Andropogondistachyos* CCGGTGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Cleistachnesorghoides* NCGGCGGAGCGGTCCGGC-NTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*CoixBuckler* TCGACGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Arundinellaneपालensis* ACGGTGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Pennisetumsetaceum* GCGGGGCTGTGGTTGGC-CTGCCGATCGCACCTCGTGCAGC---GATACT  
*Panicumbisulcatum* TCGGGGTTGTGGTTGGC-TTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Danthoniacalifornica* GCGGCGCGCGGCCAGC-CTGCCGGTCTCACACGCGTA-----  
*PentstemonisaristifoliaSpies6* GCGTGGCGGTCCGGTCCGGC-CTGCCAGCGCGCGCGCGCAGC---GATTCT  
*Eragrostisdielsii* CCAATGACTCGACCGGC-CTGCCGGGCGGTCTTGCAGCAGC---GATACC  
*Aristidaharvardii* GCGGGGCTGCGGCCGGC-CTGCCGAGCGCGCCCCAGCCAGC---GATGCT  
*Zeadiplo* CCGGCGGAGCGGTCCGGC-CCGCCTTCCGCTCCCCGGGCGAGC---GGTTA-  
*ZmaysGfl* CCGGCGGAGCGGTCCGGC-CCGCCTTCCGTTCCCGAGGCGAGC---GGTTA-  
*Evillosa* --GGCGGAGCGGTCCGGC-TTGCCTTCCGCTCCCCACGCGGT---GATCAT  
*Saccharumofficinarum* --GGTGGAGCGGTCCGGC-CCGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Imperatacylindrical* --GGCGGA-CGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GTGCT  
*Bothriochloainsculpta* --GGCGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Dichanthiumannulatum* --GGTGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Cplurnodus* --GGTGGAGNGGCTGGC-CTGCCTTCCGCTCACCAGCGCAGC---GATGAT  
*Cexcavatus* --GGTGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Hyparrhrietamba* --GGTGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Hanamesa* --GGTGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Heteropogoncontortus* --GGTGGAGCGGTCCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT

*Aameth6685* ATAC-TAATC  
*Aameth6679* ATAC-TAATC  
*Dfili5874* ATCT-TAATC  
*Ttria6555* TTAT-TAATC  
*Ttria6525* TTAT-TAATC  
*Tspic5205* ATCA-TAATC  
*Tspic6579* ATCA-TAATC  
*Tspic6580* ATCT-TAATC  
*Tspic5873* ATCA-TAATC  
*Adist6684* ATCTCTAATC  
*Mcape6584* ATCT-TAAAC  
*Aappen6543* ATCT-TAATC  
*Mcape6655* ACCT-TAAAC  
*Aappen6946* ATCT-TAATC  
*Sbic6589* ATCT-TAATC  
*Cplur6406* ATCT-TAATC  
*Icylin6632* ATCT-TAATC  
*Hpilg4738* ATCT-TAATC  
*Icylin5090* ATCT-TAATC  
*Emuti6715* ATCT-TAATC  
*Emuti6559* ATCT-TAATC  
*Emuti5074* ATCT-TAATC  
*Emuti6539* ATCT-TAATC  
*Pfrank4705* ATCT-TAATC  
*Icylin5114* ATCT-TAATC  
*Aeuco5869* ATCT-TAATC  
*Dfili5872* ATCT-TAATC  
*Aeuco6587* ATCT-TAATC  
*Bradi4784* ATCT-TAATC  
*Binsc5681* ATCT-TAATC  
*Tspic4846* ATCT-TAATC  
*Cprolix6682* ATCT-TAATC  
*Csp5800* ATCT-TAATC  
*Cprolix5101* ATCT-TAATC  
*Cdieter6642* ATCT-TAATC

<i>Cmarg4397</i>	ATCT-TAATC
<i>Hpilg4603</i>	ATCT-TAATC
<i>Htamb6585</i>	ATCTTTAATC
<i>Hdiss5145</i>	ATCT-TAATC
<i>Htamb6537</i>	ATCT-TAATC
<i>Hanam4807</i>	ATCT-TAATC
<i>Hfilli5144</i>	ATCT-TAATC
<i>Hanam4841</i>	ATCT-TAATC
<i>Hhirt6656</i>	ATCT-TAATC
<i>Hhirt6618</i>	ATCT-TAATC
<i>Shalep9318</i>	ATCT-TAATC
<i>Hcon4741</i>	ATCT-TAATC
<i>Evil15085</i>	ATCT-TAATC
<i>Bbladh5173</i>	ATCTTTAATC
<i>Hquar5088</i>	ATCT-TAATC
<i>Sbic6588</i>	ATCT-TAATC
<i>Sbicolour9319</i>	ATCA-TCC--
<i>Hrud6576</i>	ATCT-TAATC
<i>Andropogondistachyos</i>	ATCT-TAATC
<i>Cleistachnesorghoides</i>	ATCT-TAATC
<i>CoixBuckler</i>	ATCT-TAATA
<i>Arundinellanepalensis</i>	ATGT--AAC-
<i>Pennisetumsetaceum</i>	ATCT-TAATC
<i>Panicumbisulcatum</i>	ATCT-TAATC
<i>Danthoniacalifornica</i>	-----
<i>PentaschistisaristifoliaSpies6</i>	ATAC-TAATC
<i>Eragrostisdielsii</i>	TTTTGTAATC
<i>Aristidaharvardii</i>	ATCA-TAAC-
<i>Zeadiplo</i>	-----
<i>ZmaysGfl</i>	-----
<i>Evillosa</i>	ATAC-TAATC
<i>Saccharumofficinarum</i>	ATCT-TAATC
<i>Imperatacylindrical</i>	ATCT-TAATC
<i>Bothriochloainsculpta</i>	ATAC-TAATC
<i>Dichanthiumannulatum</i>	ATAC-TAATC
<i>Cplurnodus</i>	ATCC-TAACC
<i>Cexcavatus</i>	ATCT-TAATC
<i>Hyparrhrietamba</i>	ATCC-TAATC
<i>Hanamesa</i>	ATCT-TAATC
<i>Heteropogoncontortus</i>	ATCT-TAATC

**Appendix F:** CLUSTAL W (1.74) multiple sequence alignment data of the Andropogoneae specimens for *ITS2* region.

<i>B. radicans</i> 4831	TGCG-ATACCTGGTGT-GAATTGC
<i>H. quarrei</i> 6586	TGCG-ATACCTGGCGT-TAATTGC
<i>H. dissoluta</i> 5154	TACGGATACCTAGCGCTCAATTGC
<i>H. rudis</i> 6576	TACG-ATACCTGGTGTGACTTGC
<i>H. anamesa</i> 4841	TGCT-ATACCTCGAGT-GAAGTGC
<i>E. muticus</i> 6954	TACGGATACCTGGTG-GGAATTGC
<i>CoixBuckler</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>Arundinellanepalensis</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>Pennisetumsetaceum</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>Panicumbisulcatum</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>Chimonobambusabrevinoda</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>Danthoniacalifornica</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>Eragrostisdielsii</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>Aristidaharvardii</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>Zeadiplo</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>ZmaysGfl</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>Cleistachnesorghoides</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>I. cylindrica</i> 6632	TGCGGATACCTGGTGTGAATTGC
<i>D. filliformis</i> 5874	TGCGGATACCTGGGCT-AAATTGC
<i>T. spicatus</i> 4846	TGCG-ATACCTGGTGT-GAATTGC
<i>P. franksae</i> 4705	-----
<i>C. plurinoides</i> 6404	-----
<i>A. eucomus</i> 5869	TGCG-ATACCTGGTGT-GAATTGC
<i>D.filliformis</i> 5872	TGCG-ATACCTGGTGT-GAATTGC
<i>H. contortus</i> 4271	TGCGGATACCTGGTGT-GAATTGC
<i>H. contortus</i> 4741	TGCGGATACCTGGTGT-GAATTGC
<i>B. radicans</i> 4831	TCG--ATACCTGGTGT-GAATTGC
<i>A. appendiculatus</i> 6638	TGCG-ATACCTGGTGT-GAATTGC
<i>A. appendiculatus</i> 6677	TGCGGATACCTGGTGT-GAATTGC
<i>A. appendiculatus</i> 6543	TGCGGATACGTGGCGT-GAATTGC
<i>H. hirta</i> 6656	TGCGGATACCTGGTGT-GAATTGC
<i>H. hirta</i> 6550	TGCGGATACCTGGTGT-GAATTGC
<i>H. quarrei</i> 5088	TGCGGATACCTGGTGT-GAATTGC
<i>B. insculpta</i> 5681	TGCGGATACCTGGTGT-GAATTGC
<i>T. triandra</i> 6555	---GGATACTGGTGTGAATTGC
<i>Spies</i> 6620	TGCGGATACCTGGTGT-GAATTGC
<i>H. pilgeriana</i> 4738	TGCGNATACCTGGTGT-GAATTGC
<i>A. appendiculatus</i> 6625	TA--GATACTGGTGT-GAATTGC
<i>Bothriochloa</i> 7489	TGCGGATACCTGGTGT-GAATTGC
<i>H. longofolium</i> 6560	TGCGGATACCTGGTGT-GAATTGC
<i>H. longofolium</i>	TGCGGATACCTGGTGT-GAATTGC
<i>Panicum schinzii</i> 6652	TGCG-ATACCTGGTGT-GAATTGC
<i>B. serata</i>	TGCGGATACCTGGTGT-GAATTGC
<i>S. halepense</i> 9318	TGCG-ATACCTGGTGT-GACTTGC
<i>Papalum dilatatum</i> 6617	TCG--ATACCTGGTGT-GAATTGC
<i>Paspalum</i>	TGCGGATACCTGGTGT-GAATTGC
<i>Brachiaria</i>	TGCG-ATACCTGGTGT-GAATTGC
<i>Aristida adscencianis</i> 6713	TGCG-ATACCTGGTGT-GAATTGC
<i>H. contortus</i> 6628	TGCGGATACCTGGTGT-GAATTGC
<i>T. leucothrix</i> 6627	TGCG-ATACCTGGTGT-GAATTGC
<i>S. halepense</i> 9319	TGCG-ATACCTGGTGT-GAATTGC
<i>S. bicolour</i> 6588	TGCG-ATACCTGGTGT-GAATTGC
<i>S. bicolour</i> 6589	TGCGGATACCTGGTGT-GAATTGC
<i>E. muticus</i> 5074	TACGGATACCTGGTGT-GAATGCC
<i>E. muticus</i> 6715	TGCGGATACCTGGTGT-GAATTGC
<i>M. capensis</i> 6655	TGCGGATACCTGGTGT-GAATTGC
<i>C. prolixus</i> 6682	TGCGGATACCTGGTGT-GAATTGC
<i>C. excavatus</i> 5877	TAG-GANACCTAGCGT-CAATTCC
<i>C. plurinodus</i> 6406	TGCGGATACCTAGCGGTGAATTCC
<i>H. anamesa</i> 4807	TGCGGATACCTGGTGT-GAATTGC
<i>A. amethystinus</i> 6685	TGCG-ATACCTAGTGT-GAATTGC
<i>I. cylindrica</i> 5114	TGCGGATACCTAGTGT-GAATTGC
<i>I. cylindrica</i> 6582	TACT-ATACCTAGAAT-TAATTGC
<i>Monocymbium</i>	T-CG-ATACCTAGTGT-CAATTGC
<i>A. distachyos</i> 6684	TNCCGATACCTAGCGGCAATTCC

*B. bladhii* 5173  
*S. nigrostis* 6567  
*H. fillipendula* 5144  
*H.tamba* 6585  
*A. schirensis* 6581  
*E. villosa* 5085  
*C. excavatus* 6577  
*C. marginatus* 4397  
*Evillosa*  
*Saccharumofficinatum*  
*Imperatocyliculata*  
*Bothriochloainsculpta*  
*Dichanthiummannulatum*  
*Cytopogonprolixus*  
*Cplurnodus*  
*Cmarginatus*  
*Cexcavatus*  
*Hyparrhietamba*  
*Hanamesa*  
*Heteropogoncontortus*

TGCCGT--CCTAGCGT-NATTCCC  
TTCCACGTCGACCTATGACACTGC  
TGGCCCCACATGCCTATAC---GC  
TCCCGATACCTAGCGGCTA-TTGC  
ATGCGATACCTAGCGT-CAATTCC  
ATNCGATACCTAGTNT-GAATTGC  
GTGCCATACCTAGTGT-CAA-TTC  
-ATCGATACCTAGTGT-GAA-TTC  
TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC  
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TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC

*B. radicans* 4831  
*H quarrei* 6586  
*H. dissoluta* 5154  
*H. rudis* 6576  
*H. anamesa* 4841  
*E. muticus* 6954  
*CoixBuckler*  
*Arundinellaneapalensis*  
*Pennisetumsetaceum*  
*Panicumbisulcatum*  
*Chimonobambusabrevinoda*  
*Danthoniacalifornica*  
*Eragrostisdielsii*  
*Aristidaharvardii*  
*Zeadiplo*  
*ZmaysGfl*  
*Cleistachnesorghoides*  
*I. cylindrica* 6632  
*D. filliformis* 5874  
*T. spicatus* 4846  
*P. franksae* 4705  
*C. plurinoides* 6404  
*A. eucomus*5869  
*D.filliformis* 5872  
*H. contortus* 4271  
*H. contortus* 4741  
*B. radicans* 4831  
*A. appendiculatus* 6638  
*A. appendiculatus* 6677  
*A. appendiculatus* 6543  
*H. hirta* 6656  
*H. hirta* 6550  
*H. quarrei* 5088  
*B. insculpta* 5681  
*T. triandra* 6555  
*Spies* 6620  
*H. pilgeriana* 4738  
*A. appendiculatus* 6625  
*Bothriochloa* 7489  
*H. longofolium* 6560  
*H. longofolium*  
*Panicum schinzii* 6652  
*B. serata*  
*S. halepense* 9318  
*Papalum dilatatum* 6617  
*Paspalum*  
*Brachiaria*  
*Aristida adscencianis* 6713  
*H. contortus* 6628

ATTAATCCCG--CGAAACCATCG--AGTTTTTTAA-CGCAACTTGCCCC  
AT-AATCCCG--CTAATC-ATCG--AGTTGTTGAA-CTCAAGTTGCCCCC  
ATTAATCCCG--GTCAACCATCG-ATTTTTTGCAACTCAACTTGCCCC  
ATAAATCCCT--C-CAACCATCGA-AGTTTTTG-AACGCAACTTGCCCC  
AGAAATCCCG--ACGAACCATCGA-GATTCCTC-AACGCAAGTTGCCGCC  
CTGAATCCCN--C-CAACCGTCGC-AGTTGTTG-AACGCAAGTTGCCCTC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AT-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATTAATCCCG--C-CAACGATCC--AGTTTTTTAA-CGCAAGTTGCCGCC  
AT-AATCCCG--CGCAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
-----  
A--AATCCCG--CGAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAAATCCCN--CCAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCCCCC  
ATAAATCCCG--CGAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAAATCCCG--CGAA-CCATCG-AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAA-TCCCG--CGAAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAA-TCCCG--CGAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAA-TCCCG--CGAA-CCATCG-AGTTTTTGAA-CGCAAGTTGCCGCC  
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ATAA-TCCCG--CGAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAAATCCCG--CGAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAAATCCCG--CGAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AGAA-TCCCG--CGCAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AGAA-TCCCG--CG-AACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAA-TCCCG--CG-AACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAA-TCCCG--CG-AACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAAATCCCG--CG-AACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAA-TCCCG--CGAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAA-TCCCG--CGAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCCCT  
ATAA-TCCCG--CGAA-CCATCC--AATTTTTGAA-CGCAAGTTGCTCCC  
AGAA-TCCCG--CGAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
ATAAATCCCG--CGAAACCATCGT-AGTTGTTGAA-CGCAAGTTGCCGCC  
ATAAATCCCG--TGTAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC  
AGCAATCCCG--CGGAACCATCG--AGTTTTTGAA-CGCAAGTTGCCGCC

*T. leucothrix* 6627  
*S. halepense* 9319  
*S. bicolor* 6588  
*S. bicolor* 6589  
*E. muticus* 5074  
*E. muticus* 6715  
*M. capensis* 6655  
*C. prolifix* 6682  
*C. excavatus* 5877  
*C. plurinodus* 6406  
*H. anamesa* 4807  
*A. amethystinus* 6685  
*I. cylindrica* 5114  
*I. cylindrica* 6582  
*Monocymbium*  
*A. distachys* 6684  
*B. bladhii* 5173  
*S. nigrostis* 6567  
*H. fillipendula* 5144  
*H.tamba* 6585  
*A. schirensis* 6581  
*E. villosa* 5085  
*C. excavatus* 6577  
*C. marginatus* 4397  
*Evillosa*  
*Saccharumofficinarium*  
*Imperatacylindrica*  
*Bothriochloainsculpta*  
*Dichanthiumannulatum*  
*Cyobopogonprolixus*  
*Cplurnodus*  
*Cmarginatus*  
*Cexcavatus*  
*Hyparrhietamba*  
*Hanamesa*  
*Heteropogoncontortus*

AGAAATCCCG--CG-AACCATCG--AGTTTTTGAA--CGCAAGTTGCGCCC  
ATAA-TCCCG--CA-AACCATCGG--AGTTTTTGAA--CGCAAGTTGCGCCC  
ATAA-TCCCG--CG-AACCATCGG--AGTTTTTGAA--CGCAAGTTGCGCCC  
ATAA-TCCCG--CG-AACCATCGG--AGTTTTTGAA--CGCAAGTTGCGCCC  
ATAA-TCCCG--C-GAACCATCG--AGTTTTTGAA--CGCAAGTTGCGCCC  
ATAA-TCCCG--C-AAACCATCGG--AGTTTTTGAA--CGCAAGTTGCGCCC  
ATAA-TCCCG--CGCAACCATCN--AGTTTTTGAA--CGCAAGTTGCGCCC  
ATAA-TCCCG--C--AACCATCGG--AGTTTTTGAA--CGCAAGTTGCGCCC  
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AT-AATCCCG--C-NAATCATCN--ATTTTTTGAA--CGCAAGTTGCGCCC  
ATAA-TCCCG--C--AACCATCG--AGTTTTTGAA--CGCAAGTTGCGCCC  
ATACATCCCG--CTGAACCATCT--AGGTTTTTAAACGCAAGTTGCGCCC  
ATAA-TCCCG--C-CAATCATCC--ATTTTTTGAA--CGCAAGTTGCGCCC  
ATAA-TCTCG--C-CAACCATCC--ATTTTTTGAA--CGCAACTTGCGCCC  
ATAA--TCTG--GTCAATGGTCC--ATTTTTTCAA--CTCAACTTGCGCCC  
ATTATGTGCTCGAATGGATTCCATTATTTCCGACTGCAACTTGCGCCC  
--TAATGTGC-TC-AATGGGTCC--TTATGGCCGAATACACCG--GCCGCG  
--TAATTCG--TC-AATGGATTGC---ATTTGTCAATAAACCA--GCTACA  
--TACTGTGC-TC-CATAGTTC---TTATGTCCACGAATCTAGCCCC  
--TAATCCCG-TC-AATCGATTCC---TTTTTTGAACCAAGTTGCGCCC  
ATAAATCCCG--TCGAATGCATCCC--ATTTTTTCAACTCAACTTGCGCCC  
ATGAATCCCG--TCCAATC--ATCGCAGTTTTTGCAACTGCAACTTACGCC  
A-TAATCCCG--TC--AACCATCG--AGTTTTTGAA--CGCAAGTTGCGCCC  
CATAATCCCG--CGAACC--ATCGA---GTTTTTGAACGCAAGTTGCGCCC  
AG-AATCCCG--CG-AACCATCG--AGTTTTTGAA--CGCAAGTTGCGCCC  
AG-AATCCCG--CG-AACCATCG--AGTTTTTGAA--CGCAAGTTGCGCCC  
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AG-AATCCCG--CG-AATCATCG--AGTTTTTGAA--CGCAAGTTGCGCCC  
AG-AATCCCG--CG-AACCATCG--AGTTTTTGAA--CGCAAGTTGCGCCC

*B. radicans* 4831  
*H quarrei* 6586  
*H. dissoluta* 5154  
*H. rudis* 6576  
*H. anamesa* 4841  
*E. muticus* 6954  
*CoixBuckler*  
*Arundinellaneapalensis*  
*Pennisetumsetaceum*  
*Panicumbisulcatum*  
*Chimonobambusabrevinoda*  
*Danthoniacalifornica*  
*Eragrostisdielsii*  
*Aristidaharvardii*  
*Zeadiplo*  
*ZmaysGfl*  
*Cleistachnesorghoides*  
*I. cylindrica* 6632  
*D. filliformis* 5874  
*T. spicatus* 4846  
*P. franksae* 4705  
*C. plurinoides* 6404  
*A. eucomus*5869  
*D.filliformis* 5872  
*H. contortus* 4271  
*H. contortus* 4741  
*B. radicans* 4831  
*A. appendiculatus* 6638  
*A. appendiculatus* 6677  
*A. appendiculatus* 6543  
*H. hirta* 6656  
*H. hirta* 6550  
*H. quarrei* 5088

T-ATGCC-TTCTGNTCAA--GGGCACCTCTGCCTGGGCG--TCACGCC--  
G-AGGCC-TTCTGGTCAA--GGGCACCTCTGCCTGGGCG--TCACGCC--  
TGACGCC-TTCTGTTCAA--CGACACATCTGATTGGGCA--TCACAGCC--  
GGA-GCC-TTCTGGTCAA--CGGCACGCTGCCTGGGCG--TCAC-GCC--  
C-ACGTC-TTCTGGTCAA--CGGCACCTCTGCCTGAGCG--TCAA-TCT--  
GGA-GCC-TTCTGGTCAA--CGGCACCTCTGCCTGGGCG--TCCC--C--  
G-AGGCC-TTCTGGCCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AAGCC-TCCAGCCGA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AGGCC-TTCTGGCTGA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AGGCC-TTCTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AGGCC-CTCGGGCCGA--GGGCACGCTGCCTGGGCG--TCACGCC--  
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G-AGGCC-TTCTGGCTGA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AGGCC-TTCTGGCCGA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AAGCC-TTCTGGCCGA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AAGCC-TTCTGGCCGA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AGGCC-TTCTGGCCGA--GGGCACGCTGCCTGGGCG--TCACGCC--  
GTAGGCC-TTCTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
GA-GGCC-TTTTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
GAAGGCC-TTTTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
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G-AAGCC-TTTTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AAGCC-TTTTGGTCAA--GGGCAC-TCTGCCTGGGCG--TCACGCC--  
G-ATGCC-TTCTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AGGCC-TTCTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AGGCC-TTCTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-AG-CC-TTCTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-GAGCC-TTCTGGTCAA--GGGCACCTCTGCCTGGGCG--TCACGCC--  
G-AGGCC-TTCTGGTCAA--GGGCACCTCTGCCTGGGCG--TCACGCC--  
G-A-GCC-TTCTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-A-GCC-TTCTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--  
G-ATGCC-TTCTGGTCAA--GGGCACGCTGCCTGGGCG--TCACGCC--



*I. cylindrica* 6632  
*D. filliformis* 5874  
*T. spicatus* 4846  
*P. franksae* 4705  
*C. plurinoides* 6404  
*A. eucomus* 5869  
*D. filliformis* 5872  
*H. contortus* 4271  
*H. contortus* 4741  
*B. radicans* 4831  
*A. appendiculatus* 6638  
*A. appendiculatus* 6677  
*A. appendiculatus* 6543  
*H. hirta* 6656  
*H. hirta* 6550  
*H. quarrei* 5088  
*B. insculpta* 5681  
*T. triandra* 6555  
*Spies* 6620  
*H. pilgeriana* 4738  
*A. appendiculatus* 6625  
*Bothriochloa* 7489  
*H. longifolium* 6560  
*H. longifolium*  
*Panicum schinzii* 6652  
*B. serata*  
*S. halepense* 9318  
*Papalum dilatatum* 6617  
*Paspalum*  
*Brachiaria*  
*Aristida adscencianis* 6713  
*H. contortus* 6628  
*T. leucothrix* 6627  
*S. halepense* 9319  
*S. bicolor* 6588  
*S. bicolor* 6589  
*E. muticus* 5074  
*E. muticus* 6715  
*M. capensis* 6655  
*C. prolixus* 6682  
*C. excavatus* 5877  
*C. plurinodus* 6406  
*H. anamesa* 4807  
*A. amethystinus* 6685  
*I. cylindrica* 5114  
*I. cylindrica* 6582  
*Monocymbium*  
*A. distachyos* 6684  
*B. bladhii* 5173  
*S. nigrostis* 6567  
*H. fillipendula* 5144  
*H. tamba* 6585  
*A. schirensis* 6581  
*E. villosa* 5085  
*C. excavatus* 6577  
*C. marginatus* 4397  
*Evillosa*  
*Saccharum officinarum*  
*Imperata cylindrical*  
*Bothriochloa insculpta*  
*Dichanthium annulatum*  
*Cytopogon prolixus*  
*Cplurnodus*  
*Cmarginatus*  
*Cexcavatus*  
*Hyparrhnia tamba*  
*Hanamesa*  
*Heteropogon contortus*

AAAAGAC-ACTCCC-AACCCACCC-TCGGGG-----AGGGACATGGT-  
AAAACAC-ACTCCC-AAACCGAAA-TCCGGG-----AGGGACGTGGT-  
AAAAGAC-ACTCCC-AACCCACCC-TCGGGGG-----AGGGACGTGGT-  
--GTACC-CGCTCAGATACTAG---CAGCAA---ATAGGACTTCCT-  
-----TCGGG-----GTAGTA-TTGGC-  
AAAAGAC-ACGTCCCAAGCCAC---AATCAAG---GAGGTA-CTGGC-  
AAAAGAC-A-CTCCCAAGCCAC---CCTCGGG-----GAGGACTTGGC-  
AAAAGAC-A-CTCCCAACCCAT---CCTCGGG-----CGGGGACTTGGC-  
AAAACAC-A-CTCCCAACCCAA---AATCGGG-----CGGGGATGTGGT-  
AAAAGAC-ACTCCC-AACCCAA---CCTCGGG-----AGGG-ATGTGGT-  
AAAAGAC-ACTCCC-AACCCAA---CCTCGGG-----AGGG-ATGTGGT-  
AAAAGAC-ACTCCC-AACCCAA---CCTCGGG-----AGGG-ATGTGGT-  
AAAACAC-ACTCCC-AACCCAA---AATCGGG-----AGGG-ATGTGGT-  
AAAA-AC-ACTCCC-AACCCAC---CCTCGGG-----ACGG-CAGTGGT-  
AAAAGAC-ACTCCC-AACCCAC---CATCGGG-----ACGG-ACGTGGT-  
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AAAAGAC-ACTCCAACCCAC---CCTCGGG-----AGGG-ACGTGGT-  
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AAAAGAC-ACTCCAACCCGT---ACTCGGG-----AGGG-ACGTGGT-  
AAAATAC-TCTCCAACCCGG---AATCGGG-----ACGG-ACGTGGT-  
AAAATAC-ACTCCAACCCGG---ACAAGGG-----AGGG-ACATGGT-  
-AAACAC-GCTCCCGACCCCACTAACCTGTGAT---CTGC-ATGCCAC-  
-AAACAC-GCTCCC-ACCCCACTAACCTGTGGT---GGG---ATCGCGC-  
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AAAAGAC-GCTCCC-AACCCA--ACCAAGGGG-----AGGG-ACGTGGT-  
ACATTAC-ACTCCC-AACCCA---CCCTCGGG-----AGGG-ATGTGGT-  
ACAGTAC-ACTCCC-AACCCA---GCCTCGGG-----AGGG-ATGTGGT-  
ACAG-AC-ACTCCC-AACCCA---GGCTCGGG-----AGGG-ATGTGGT-  
AAAANAC-ACTCCC-AAACCAC---AATCGTGG---AGGGAC-TGGT-  
AAAACACTACTCCC-AACCCAC---NCTCGTGG---AGGGACATGTT-  
AAAACAC-ACTCCAACCCAC---ACAAGG-----AAAGGACATTTT-  
AAAATGAC-ACTCCC-AACCCAT---CCTCGGG-----ACGGACGTGGT-  
AAAATAC-ACTCCAACCCCAA---CCCTCGGG-----AGG--ACCTTGC-  
AAAAGAC-ACTCCAACAGCA--CCCTCGAGCGGCATAATGGACCTGAC-  
AAAATAC-ACTCCC-AACCCAT---CCTCGGG-----AGGACATGGT-  
ATAGCAC-ACTCACCAGATGAC---CCTCGGCT---ACAGATTTGGT-  
AAATGAT-ACTCGCAACGCGCAT---CCTCGGG-----AGGGACAT-TC  
AAATGAC-ACTCCC-AACCCAT---CCTCGGG-----ACGGACATTTT-  
TAATGAT-ACTCCG-AACCCCTTCG-CAAGGGG-----ACGGACATTTT-  
GAATGAG-ACGTGCACAGATTCTTCGGCAGACGG---CACACTTGG-  
GC-TGTG-CC---CGGCACGGCTATCCGGCAACC-----CAAACCTCGG-  
GC--GGC-GCTCGAACAAGTTTTGCTGGCAGCC-----CAAACCTCG--  
TC-TGTG-CC---CAATCCACTTGTGCCAGGC-----CCCACTTTTG-  
GCTGTAC-ACTGCCACAAC-CACTATGGCAGTAG---CTACCTTCTG-  
-T--GAC-ACT---CACAAAACACCCCTTGCAGAC-----CCACCTTCTG-  
-AAAGAC-ACTCCAGTCCCTACCCCTGACAGGGCA--TCTACTTCTT-  
AAATATG-ACTCGAGACCATTCCTCGCAGCA-----CGCACCTTCG-  
-AAATAC-ACT---CCCAACCAATCCTCGGAGAAG---GACGCTTGG-  
AAAAGAC-ACTCCAACCCGA---CCTCGGG-----AGGG-ACGTGGT-  
AAAAGAC-ACTCCAACCCAC---CCGAGGGG-----AGGG-ACGTGGT-  
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AAAAGAC-ACTCCAACCCAC---CCTCGGGGGGG---AGNTGGN-  
AAAAGAC-ACTCCATCCCAN---CCTCGGGGGGG---ACGTGGT-  
AAAANAA-NCTCCAACCCAN---CTTGGGACGGG-----ACGTGGT-

*B. radicans* 4831

-TTTCGCCTCCCTTCC-GGCAAGGGCGAGGCTGAACACAACCTTC

*H. quarrei* 6586  
*H. dissoluta* 5154  
*H. rudis* 6576  
*H. anamesa* 4841  
*E. muticus* 6954  
*CoixBuckler*  
*Arundinellaneपालensis*  
*Pennisetumsetaceum*  
*Panicumbisulcatum*  
*Chimonobambusabrevinoda*  
*Danthoniacalifornica*  
*Eragrostisdielsii*  
*Aristidaharvardii*  
*Zeadiplo*  
*ZmaysGfl*  
*Cleistachnesorghoides*  
*I. cylindrica* 6632  
*D. filliformis* 5874  
*T. spicatus* 4846  
*P. franksae* 4705  
*C. plurinoides* 6404  
*A. eucomus* 5869  
*D. filliformis* 5872  
*H. contortus* 4271  
*H. contortus* 4741  
*B. radicans* 4831  
*A. appendiculatus* 6638  
*A. appendiculatus* 6677  
*A. appendiculatus* 6543  
*H. hirta* 6656  
*H. hirta* 6550  
*H. quarrei* 5088  
*B. insculpta* 5681  
*T. triandra* 6555  
*Spies* 6620  
*H. pilgeriana* 4738  
*A. appendiculatus* 6625  
*Bothriochloa* 7489  
*H. longofolium* 6560  
*H. longofolium*  
*Panicum schinzii* 6652  
*B. serata*  
*S. halepense* 9318  
*Papalum dilatatum* 6617  
*Paspalum*  
*Brachiaria*  
*Aristida adscencianis* 6713  
*H. contortus* 6628  
*T. leucothrix* 6627  
*S. halepense* 9319  
*S. bicolor* 6588  
*S. bicolor* 6589  
*E. muticus* 5074  
*E. muticus* 6715  
*M. capensis* 6655  
*C. prolixus* 6682  
*C. excavatus* 5877  
*C. plurinodus* 6406  
*H. anamesa* 4807  
*A. amethystinus* 6685  
*I. cylindrica* 5114  
*I. cylindrica* 6582  
*Monocymbium*  
*A. distachyos* 6684  
*B. bladhii* 5173  
*S. nigrostis* 6567  
*H. fillipendula* 5144  
*H. tamba* 6585  
*A. schirensis* 6581  
*E. villosa* 5085  
*C. excavatus* 6577  
  
GTTTGGCCGTCCTCCCTGTC-GGCAAGGGCGAGGCTTGACACAACCTTG  
CTTTGCCC--CCCCGTAC-CATACGACGGGGGTGAACACAAACTT-  
GTTTGGCC--CCCCCTGTC-CATACGGCGGGCC-GGACACGAAATT-  
NTT-----  
TTGGACCC--CTCCCCCC-NNGAGGGCAGGGGTGGGAA-NAAGTT-  
ATCTGGCC--CCCCGCGC-CGCAAGGCGCGG-TGGGCCG-AAGTT-  
GTATGGCC--CCCCGTGC-CGCATGGCGCGG-TGGGCCG-AAGTT-  
GTTTGGCT--CCTCGTGC-TGCAAAGCGCGG-TGGGCCG-AAGTT-  
GTTTGGCT--CCCCGTGC-CTCAGGTGCGG-TGGGCCG-AAGTT-  
GTTTGGCC--CCCCGCGC-CGCACGGCGCGG-TGGGCCG-AAGTT-  
GTTTGGCT--CCCCGTGC-CGCAGGGCGCGG-TGGGCCG-AAGTT-  
ATTTGGCT--CCCCGTGC-AACACGACACGG-TGGGCCG-AAGTT-  
GTATGGCC--CCCCGTGC-CGCGAGGCGCGG-TGGGCCG-AAGTT-  
GTCTGGCC--CCCCACGC-CGCAGGGCGAGG-CGGGCCG-AAGCA-  
GTCTGGCC--CCACGCGC-CGCAGGGCGAGG-TGGGCCG-AAGCA-  
GTTTGGCC--TCCCGTGC-CTCACGGCGCGG-TGGGCCG-AAGTT-  
GTTTGGAC--CCCCGTGC-CGCAGGGCGCGG-TGGGCCA-AACTT-  
GTTTGGCC--CCCCGTGC-CGCAGGGCGCGA-TGGGCCGAAAGTT-  
GTTTGGCT--CCCCGTGC-CGCAGGGCGCGG-TGGGCCA-AACTT-  
GTTTGGAT--CCCCGTGC-CGCATGGGCACAGTCGGGGCGAAAGTT  
GTCTGCC--CCCCCTGC-CTCATGACGCGGATGGTCG--AATAT-  
GTATGGCC--CCCCGTGC-CGCACGGCGCGG-TGGGCCG-AAATT-  
GTATGGCC--CCCCGTGC-CGCACGGGGCGGTTGGGCC--AAATT-  
GTTTGGCC--CCCCGTGC-CGCAGGGCGCGG-TGGGCCA-AAATA-  
GTTTGGCC--TCCCGTGC-CGCAAGGCGCGG-TAGGCCG-AAGTT-  
GTTTGGCC--TCCCGTGC-CGCAGGGCGCGG-TGGGCCG-AAGTT-  
GTTTGGCC--TCCCGTGC-CGCAGGGCGCGG-TGGGCCG-AACTT-  
GTTTGGCC--TCCCGTGC-CGCAGGGCGCGG-TGGGCCG-AACTT-  
GTTTGGCC--CCCCATGC-CGTATGGCGCGG-TGGGCCG-AACTT-  
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GTTTGGCT--CCCCGTGC-CGCAGGGCGCGG-AGGGCCA-AAATT-  
GTTTGGCC--CCCCGTGC-CGCAGGGCGCGG-TGGGCCG-AAGTT-  
GTTTGGCC--CCCCGTGC-CGCAGGGCGCGG-TGGGCCG-AAAGAT-  
GTTTGGCC--CCCCGTGC-CGCAGGGCGCGG-TGGGCCA-AACTT-  
GTTTGGCC--CCCCGTGC-CGCAGGGCGCGG-TGGGCCG-AAGTT-  
ATGTGGCT--CCCCGTACTCGCAAGCGCGGTGTGCGGCTCAAGAT-  
ATGTGGCC--CCCCGT-CTCGCAAGGGCGCGG-TGGGCCA-AAAGAT-  
GTTTGGCT--CCCCCTGC-CTTGCNGATGG-TGGGCTA-AAAGTT-  
GTTTGGCT--CCCCCTAC-CGCAAGGGCGCGG-TGGGCCG-AAATT-  
GTTTGGCT--CCCTGTGC-TGCAAAGCGCGG-GGGGCTG-AAATT-  
GTTTGGCC--TCTCGTGC-CGTTGTGCACAGGTGGGCCA-AAATC-  
GTTTGGCT--CCTCGTAC-TGCATGGTGCAGG-TGGGCCG-AAGTT-  
GTTTGGCT--CCCCGTGC-CGCAAGGAGCGG-TGGGCCA-AAAGTT-  
GTTTGGCT--CCTCGTGC-CGTAAGGTGCGG-TGGGCCG-AAGTT-  
GTTTGGCT--CCCCGTGC-CGCAGGGCGCGG-TGGGCCG-AAAGAT-  
GTTTGGCT--CCCCGTGC-CGCAGGGCGCGG-TGGGCCG-AAAGAT-  
TTTTGGCT--CCCCGATGCCGATGGCGTGG-TGGGCCG-AAGTT-  
TTTTGGCT--CCCCGATGCCGATGGCGCGG-TGGGCCG-AAAGTT-  
TTTTGGCT--CCCCGA-GCCGATGGCGCGG--GGGCCG-AAGTT-  
GTTTGGCC--CCTCGCCTCAGG--CGCGG-TGGGCC-AAATTT-  
GTTTGGCC--CCTC-CGCCTCANGG-CACGG-GGGGCC-AAATTT-  
GTTTCGTC--CCCCCTTCCGCAAGG-CGCGA-TGGGCCA-AAATTT-  
GTCTGGCC--CCCCGTGCCGCAAGG-CGGGG-TGGGCCA-AAAGTT-  
GTCTGGCC--CCCCGTGCCGACAGG-GGGGG-TGGACCCCAAGTT-  
GTCTGGCC--CCCCGCCCCGAGG-CGAAG-TGGGCCCAATAT-  
GTTTGGCC--CCCCGTGCCGCAAGGCGGGG-TGACCACAACCTT--  
GTTTCGCC--CCTCGTGGCACAGGGGGCGAGC-ACCACAAGTT--  
GTTTGGAC--CCCCCTGCCGCGAGGGGGCGGTTGGGACACAAGTTG  
GTTTCGAC--CCCCCTGCCGCGAGG-GGGTTACAGACACAATTTG-  
GTTTGGAC--CCCCGTGCCGGAAGG-GGGGGTCAGACAC-AACTG-  
-TTTCACC-CCTCTTGGCGCAAGGAGGGGCCGAACAC-AACGTG-  
-CT-CACC-TGCGCCTGCGGCA-GGGGGAGCCAAACACC-AC-TG-  
-TTCCACC-CCACCATCTAGCT-CGAGCAGCCAAACACC-ACGTG-  
-GCCAACC-CCACCTATTGAACGTTG--GGCAAACACC-CCCTT-  
-CTCTACC-ACCCCGTGGCGCACGAGG-GGCATACACC-AACAT-  
-TT-TACC-GCCCCGTGCCGACGGGGG-GCCGAACACC-AAGTC-  
-TCCTTCC-TCCCCGTGCCGAGGACGA-GGAACCAACACC-ATGAT-  
-GTCTAGC-GCCCCCTGCCGACAGG-GGGGCCAGACCC-AACTT-



*C. marginatus* 4397 -TT-GACC-GCGCCCTGCGGCAAGGGGGGCCAGACACACACTG--  
*Evillosa* GTTTGGCC-CCTCGTGC-CGCAAGG-CGCGG-TGGGCCG-AAGTT-  
*Saccharumofficinatum* GTTTGGTC-CCCCGTGC-CGCAGGG-CGCGG-TGGGCCG-AAGTT-  
*Imperatocylindrical* GTTTGGAC-CCCCCGC-CGCAGGG-CGCGG-TGGGCCG-AAGTT-  
*Bothriochloainsculpta* GTTTGGCC-CCTCGTNC-CTCAAGG-CACGG-TGGGCCG-AAGTT-  
*Dichanthiumannulatum* GTTTGGCCCCCNCNTC-CNNAAGG-GGNGGGGGCCC--AAGTT-  
*Cyobopogonprolixus* GTTTGGCCCCCCTTCCC-CAAGGGG--GAGG-TGGCCCCG-AAGTT-  
*Cplurnodus* GTTGNCCCCCNCNTGCC-CAAGGGG--GAGG-TGGCCCC-AAGTA-  
*Cmarginatus* GTTTGGCCCCCCTTCTACAANAN--NACGG-TGGGCCG-AAGTT-  
*Cexcavatus* GTTTGGCCCCCCTGCC-GAATGGG-GAGG--TGGTACCNAAGTT-  
*Hyparrhinetamba* -TTTGGCCCCCTCCGTCC-AAAGGGG-GGGGG--GGCCC--AAGNT-  
*Hanamesa* TTTTGGCCCCCCTTCC-CCATTGG-CGCGG-TGGGGCGGGAATT-  
*Heteropogoncontortus* GTTTGNCCCCCCTNCC--GCAGGG-CGCGG-TAGGCCG-AAGTT-

*B. radicans* 4831 CGGTCTGCCGAGCGTAATCTTGCCGAGGTCACA-TTACG-TGGTTGACG  
*H. quarrei* 6586 CCGTCT-CCGAG-GTAATCTTGCCGAGGTGACA-TTTGG-TGGTGAGCC  
*H. dissoluta* 5154 CGGGCTTCCAGCA-TTATCTATCTTA-GTGAATTTTT---TCGGTATGAC  
*H. rudis* 6576 GCCTCTCCCATGC-CTATCTGTGGCACGTGCACTTTTAGCTAGTACGGCN  
-----  
*H. anamesa* 4841 GGGGCTGCCGAA-CGATCGGCTTTCGGACACATTTT-----TGGTGGAC  
*E. muticus* 6954 GGGGCTGCCGT---CGAATCGTG-TCGGG-CACA-GCACG-TGGTGGGCG  
*CoixBuckler* GGGGCTGCCGG---CAAACCGTG-TCGGG-CACA-GCATG-TGGTGGGCG  
*Arundinellaneapalensis* GGGGCTGCCGG---CATAACATG-CCGGG-CACC-GCACG-TGGTGGGCG  
*Pennisetumsetaceum* TGGGCTGTGG---CATAACCGTG-CCGGG-CACC-GCACG-TGGTGGGCG  
*Panicumbisulcatum* CGGGCTGCCGG---CGCAGCACG-CCGGG-CACA-GCACA-AGGTGGGCG  
*Chimonobambusabrevinoda* GGGGCTGCCGG---CGTACCGTG-CCGGG-CGCA-GCACA-AGGTGGGCG  
*Danthoniacalifornica* GGGGCTGCCGG---CTTACGGTA-CCGGT-CACA-GCACA-AGGTGGGCG  
*Eragrostisdielsii* GGGGCTGCCGG---CGTATCGCG-TCGGG-CACA-GCACG-TGGTGGGCG  
*Aristidharvardii* GGTGCTGCCGG---CGAACCGCG-CCGGG-CGCA-GCACG-TGGTGGGCG  
*Zeadiplo* GGGGCTGCCGG---CGAACCGCG-CCGGG-CGCA-GCACG-TGGTGGGCG  
*ZmaysGfl* GGGGCTGCCGG---CGAACCGCG-CCGGG-CGCA-GCACG-TGGTGGGCG  
*Cleistachnesorghoides* GGGGCTGCCGG---CGAATCGTG-TCGGG-CACA-GCACG-TGGTGGGCG  
*I. cylindrica* 6632 GGGGCTGCCGG---CGAA-CGTG-TCGGG-CACA-TGACG-TGGTGGGCG  
*D. filliformis* 5874 GGGGCTGCCATC--TGAATCATA-CCGGTACACA-TCACG-CGGTGGGCG  
*T. spicatus* 4846 GGGGCTGCCAG---TGAATCACAGCCGGTACACA-TCACG-TGGTGGGCG  
*P. franksae* 4705 GGGGCTGCCGG---ATAACCG-TGTCGGG-CACA-GCATT-TGGTGGGC-  
*C. plurinoides* 6404 GGGGCTGCCGGC--G-AATCG-CGCCGGG-TACT-GGACG-TGGTGGACC  
*A. eucomus* 5869 GGGGCTGCCGGC--T-AATCC-TGCCAGG-CACA-GCACC-TGGTGGGC-  
*D. filliformis* 5872 GGGGCTGCCAGC--A-AATCT-TGCCAGG-CACA-TCACG-TGGTGGGC-  
*H. contortus* 4271 GGGGCTGCCGGC--A-AATCT-TGCCGGG-CACA-CCACT-TGGTGGGC-  
*H. contortus* 4741 GGGGCTGCCGGC--G-AATCG-TGTCGGG-CACA-GCACG-TGGTGGGC-  
*B. radicans* 4831 GGGGCTGCCG-G--CGAATCG-TGCCGGG-CACA-GCACG-TGGTGGGCG  
*A. appendiculatus* 6638 GGGGCTGCCG-G--CGAATCA-TGCCGGG-CACA-TCACG-TGGTGGGCG  
*A. appendiculatus* 6677 GGGGCTGCCGAG--CTAATCN-TGCCGGG-CACA-TCACG-TGGTGGGCG  
*A. appendiculatus* 6543 GGGGCTGCCG-G--CTAATCG-TGCCGGG-CACA-TCACG-TGGTGGGCG  
*H. hirta* 6656 GGGGCTGTGCG-A--CGAATCG-TGTCGGG-CACA-TCACG-TGGTGGGCG  
*H. hirta* 6550 GGGGCTGTGCG-A--CGAATCG-TGTCGGG-CACA-GCACG-TGGTGGGCG  
*H. quarrei* 5088 GGGGCTGCCGGC--G-AATCG-TGTCAGG-CACA-GCACG-TGGTGGGCG  
*B. insculpta* 5681 GGGGCTGCCGGC--G-AATCG-TGTCAGG-CACA-GCACG-TGGTGGGCG  
*T. triandra* 6555 GGGGCTGCCGGC--G-AATCG-TGTCG-G-CACA--CAC--TGGTGGGCG  
*Spies* 6620 GGGGCTGCCGGC--G-AACCG-TGTCGGG-CACA-GCACG-TGGTGGGCG  
*H. pilgeriana* 4738 GGGGCTGCCGGC--GGAATCG-CGCCGGG-CACA-GCACG-TGGTGGGCG  
*A. appendiculatus* 6625 GGGGCTGCCGAC--GGAATCGTCGCCGAG-CACA-TTACG-TGGTGGGCG  
*Bothriochloa* 7489 GGGGCTGCCGGC--GGAATCG-TGCCGGG-CACA-GCACG-TGGTGGGCC  
*H. longifolium* 6560 CCAGGTGCCGGT---GTATCGT-ACCAGG-CACA-GCAGG-TGGCACCGG  
*H. longifolium* CCGGCTGCCGGT---GTATCGTGACCGGA-CACA-GCAGG-TGGTACCGG  
*Panicum schinzii* 6652 GTGGCTGCCGGC---ATACCGN-GTCGGG-CACCTGCATG-TGNGGGCC  
*B. serata* GGAGCTGCCGGC---GTACCTT-GCCGGG-CACC-GCACG-TGGTGGGCG  
*S. halepense* 9318 GGGGCTGCCGGCA-TATACATT-GCCGGG-CACC-GCATG-TGGTGGGCG  
*Papalum dilatatum* 6617 TTGGCTGCCGG---TGTATCTT-GCTGGG-CACC-TCATA-TGGTGGGCG  
*Paspalum* GGGGCTGCCGGC---ATAACTT-GCCGAG-CACC-GCACG-TGGTGGGCG  
*Brachiaria* GGGGCTGCCGGC---ATAACTT-GACTGG-CACC-TCACG-TGGTGGGCA  
*Aristida adscencianis* 6713 GGGGCTGCCGGC---ATATCTT-GCCGGG-CACC-GCACG-TGGTGGGCG  
*H. contortus* 6628 GGGGCTGCCGGC---GGGACCGT-GCCGGG-CACC-GCACG-TGGTGGGCG  
*T. leucothrix* 6627 GGGGCTGCCGGC---GGGACCGT-GCCGGG-CACC-GCACG-TGGTGGGCG  
*S. halepense* 9319 GGGGCTGCCGGC---GATTCGT-GTCGGG-CACA-TCACG-TGGTGGGCG  
*S. bicolor* 6588 GGGGCTGCCGGC---GATTCGT-GTCGGG-CACC-TCACG-TGGTGGGCG  
*S. bicolor* 6589 GGGGCTGCCGGC---GATTCGT-GTCGGG-CACA-TCACG-TGGTGGGCG  
*E. muticus* 5074 GGGGCTGCCGG---CATATCG-CGTGGG-CACA-TCACG-TGGTGGGCG  
*E. muticus* 6715 GGGGCTGCCGGA--CAGATCG-CGTGGG-CACA-TCTT--TGGTGGGCG  
*M. capensis* 6655 GGGGCTGCCCTCG---GAATCGT-GTCAGA-CACA-TTATG-TGGTGGACC

*C. prolixus* 6682  
*C. excavatus* 5877  
*C. plurinodus* 6406  
*H. anamesa* 4807  
*A. amethystinus* 6685  
*I. cylindrica* 5114  
*I. cylindrica* 6582  
*Monocymbium*  
*A. distachyos* 6684  
*B. bladhii* 5173  
*S. nigrostis* 6567  
*H. fillipendula* 5144  
*H. tamba* 6585  
*A. schirensis* 6581  
*E. villosa* 5085  
*C. excavatus* 6577  
*C. marginatus* 4397  
*Evillosa*  
*Saccharumofficinorum*  
*Imperatacylindrica*  
*Bothriochloainsculpta*  
*Dichanthiumannulatum*  
*Cytopogonprolixus*  
*Cplurnodus*  
*Cmarginatus*  
*Cexcavatus*  
*Hyparrhnieta*  
*Hanamesa*  
*Heteropogoncontortus*

GGGGCTGCCGGC--GGAATCGTCGCCGGGCACA-TCACG-TGGTGGGCC  
CCCTCTCCCAGCG--TAATCT-CGCCAG-CACA-TTACG-TGGTGAGCC  
GGGGCTGCCAGC---TAATCT-CGCCAG-CACA-TTACC-TGGTCGGCC  
GCGGCTCCCGAC--GGAATCGTTGCCGAGTGACA-TTACG-TGGTGAGCC  
CTAGCTGACGGGTGATCTCGT-GACTGT-CTTTGGACG-TGACTGGCC  
GGGTCTGCCGAG---GGAGCCTTGGCGAGTGACTTTT--GGTGGTTGACG  
GGGTCTGCCGAG---GGAGCCGTGGCGAGTGACA-TTTCG-TGGT-GACG  
-GGGTTGCCCTG--GGAGTCGTTGGCGAGTGACTTTT--GGTGGT-GACG  
CCCTCTCCCAGC--GTAG-TCGTTGTCGAGTG-ACCTTTGATGGT-GAGC  
CCCTC-CCCTG--GGAG-GCTTGG--GAGTG-ACCTTTTGGCGGT-GACG  
CT-TC-GCCATG--GATG-CGTTGG--GAGTG-TCTTTAGTCG-T-GACG  
CCTTC-CCCAAG--GTAC-CCTTGG--AATTT-ACCTTTGTCATT-AACC  
CTCTCTGCCATG--CATGGCCGTGTCAGGTGCACCTTTTAGCT-T-AACG  
CC-TCCTGCAAG--GCTAATCTGGG---AGTG-TATTTTAGCG-T-GACG  
CCCTC---CCGG--GGAGTGCTGGC---AGTG-TCTTTTCGCG-T-GACG  
CCCTCTCCCAGG--GTAG-TGTTGC--GAGTG-TCTTTTGGCGTT-GACG  
CCCTT-GCCGAG--GGAG-TGTTGCG-GAGTG-ACCTTTTCGCGT-GACG  
GGGGCTGCCGGC---ATACCG-TGTCGGG-CACA-GCACG-TGGTGGGCG  
GGGGCTGCCGGC---GAATCG-TGTCGGG-CACA-GCACG-TGGTGGGCG  
GGGGCTGCCGGC---GAACCG-GCCGGG-CACA-GCACG-TGGTGGGCG  
GGGGCTGCTCCGGCGAATCG-TGTCGGG-CACA-GCACG-TGGTGGGCG  
GGGGCTGCTCCGGCGAATCG-TGTCGGG-CGCA-GCACG-TGGTGGGCG  
GGGGCTGCGNGC---GANTCG-TGTTGGG-CACA-GCACG-TGGTGGGCG  
GGGGCTGCTGGC---GATTCG---TTTCGG-TAGCAAAAAG-TGGTGGGCG  
GGGGCTGCCGGC---GAATCG-TGTTGGG-CACA-GCACG-TGGTGGGCG  
GGGGCTGCCGGC---GATTCG---TTTCGG-TAACAGAACG-TGGTGGGCG  
GGGTCT-CTNCCGGCGAATCG-TTTCGGG-TACA-ACACA-TGGTGGGCG  
GGGGCTGCCGNC---GAATCG-TATCGGG-CACA-AAAAA-TGGTGGGCG  
GGGNTGCCGNC---GAATCG-TGTCGGG-CACA-GCACG-TGGTGGGCG

*B. radicans* 4831  
*H. quarrei* 6586  
*H. dissoluta* 5154  
*H. rudis* 6576  
*H. anamesa* 4841  
*E. muticus* 6954  
*CoixBuckler*  
*Arundinellaneapalensis*  
*Pennisetumsetaceum*  
*Panicumbisulcatum*  
*Chimonobambusabrevinoda*  
*Danthoniacalifornica*  
*Eragrostisdielsii*  
*Aristidaharvardii*  
*Zeadiplo*  
*ZmaysGfl*  
*Cleistachnesorghoides*  
*I. cylindrica* 6632  
*D. filliformis* 5874  
*T. spicatus* 4846  
*P. franksae* 4705  
*C. plurinoides* 6404  
*A. eucomus* 5869  
*D. filliformis* 5872  
*H. contortus* 4271  
*H. contortus* 4741  
*B. radicans* 4831  
*A. appendiculatus* 6638  
*A. appendiculatus* 6677  
*A. appendiculatus* 6543  
*H. hirta* 6656  
*H. hirta* 6550  
*H. quarrei* 5088  
*B. insculpta* 5681  
*T. triandra* 6555  
*Spies* 6620  
*H. pilgeriana* 4738  
*A. appendiculatus* 6625  
*Bothriochloa* 7489  
*H. longofolium* 6560

CCACATC---AGATGTT---CTC---TCTGACATACA-CTCCTCAC---G  
CAAAATC---AGATGTT---CTC---TCTGACACATAACTCCTCAC---A  
C-ACAACA---TTTGTTCCTCGSCGATAAAA-ATCCCTCCCGCGCT---  
C-ACGTCA---AACTTCTCNTCGCAGAGACTCCATCATGGCTGCCAGT-TA  
-----  
G-ACATGA---AGTTGTTT-TCGGTGACAAGACTCGTCACGCAACCAG-TC  
--ACACCT---AGTTGTT---CTC---GGTG-CAG-CGCCCCAGCAC---G  
--ACAAC---AGTTGTT---CTC---GATG-CAG-TGGCTCGACAC---G  
--ACATTT---AGTTGTT---CTC---GGTG-CAG-CGTCTCGGTA---G  
--ACTTAC---AGTTGTT---CTC---GGTG-CAG-CGTCCCGGTAC---G  
--ATCTCGCACTCTCTT---A-C---G-TG-CAG-TGCCCCGGCGT---G  
--ACCA---ACGTTGTC---A-C---GGTG-CTG-CGCCCCGGGCC---G  
--ACAC---ACGGTGT---CTC---GCTG-CTG-CGAACTGGTCCA-AG  
--ACCT---AGTTGTC---ATC---GGTG-CCG-TGACCCGACGC---G  
--ACATC---AAGTTGTT---CTC---GGTG-CAG-CGTCCCGGGCG---G  
--ACATC---AAGTTGTTGTTCTC---GGTG-CAG-CGTCCCGGGCG---G  
--ACACCT---AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCAC---G  
--ACATC---AGTTT---CTC---GGTG-CA---CACCCGGCGC---G  
--ACATC---AATT-TC---T-C---GNG-CAA-TGCCNCGNAA---G  
--ACATC---AGTTGTC---T-C---GGTGACAC-TGCCCCGGCAC---G  
G-AGATCT---ACTGTT---GCC---GGTG-CAG-CGCCCCGGCAC---G  
G-ATATC---AGTTGTT---CTC---GGTG-CAT-TGCCCCGGCGC---G  
G-ACATC---AGTTGT---CTC---CGTG-CAG-TGACCCGTCAA---A  
--ACATC---AATTGT---CTC---GGTG-CAA-TACCTCAA---G  
A-ACGCA---AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCAC---G  
G-ACATT---AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCAC---G  
--ACATC---AGTTGTT---CTC---TGTG-CAA-TGCCCTCGGCAC---G  
--ACATC---AGTTGTT---CTC---TGTG-CAA-TGCCCTCGGCAC---G  
--ACATC---AGTTGTT---CTC---TGTG-CAA-TGCCCTCGGCAC---G  
--ACATC---AGTTGTT---CTC---GGTG-CAT-TGCCCTCGGCAC---G  
--ACATT---AGTTGTT---CTC---GGTG-CAT-TGCCCTCGGCAC---G  
C-ACATT---AGTTGTT---CTC---GGTG-CAT-TGCTTCGGAGC---C  
C-ACATCC---G-TTGT---CTC---GGTG-CAG-CGCCCCGGCAT---G  
--ACATCT---AATTGTT---CTC---GGTG-CA---CGCCCCGGCAC---G  
--ACATCT---AGTTGTT---CTC---GGTG-CAA-CGCCCCGGCAC---G  
--ACATCA---G-TTGT---CTC---GGTG-CAG-TGCCCCGGCAC---G  
C-ACATCA---A-TTGT---CTC---GGTG-CAG-TGCCCCGGCAC---G  
--ACTCCC---G-TTGT---CTC---GGTG-CAG-CGACCCGGCAC---G  
--ATCT---CGCTATA---CTA---AACA-CAAATGCATCCGGGG---C

*H. longofolium*  
*Panicum schinzii* 6652  
*B. serata*  
*S. halepense* 9318  
*Papalum dilatatum* 6617  
*Paspalum*  
*Brachiaria*  
*Aristida adscencianis* 6713  
*H. contortus* 6628  
*T. leucothrix* 6627  
*S. halepense* 9319  
*S. bicolour* 6588  
*S. bicolour* 6589  
*E. muticus* 5074  
*E. muticus* 6715  
*M. capensis* 6655  
*C. prolixus* 6682  
*C. excavatus* 5877  
*C. plurinodus* 6406  
*H. anamesa* 4807  
*A. amethystinus* 6685  
*I. cylindrica* 5114  
*I. cylindrica* 6582  
*Monocymbium*  
*A. distachyos* 6684  
*B. bladhii* 5173  
*S. nigrostis* 6567  
*H. fillipendula* 5144  
*H. tamba* 6585  
*A. schirensis* 6581  
*E. villosa* 5085  
*C. excavatus* 6577  
*C. marginatus* 4397  
*Evillosa*  
*Saccharumofficinorum*  
*Imperatacylindrical*  
*Bothriochloainsculpta*  
*Dichanthiumannulatum*  
*Cytopogonprolixus*  
*Cplurnodus*  
*Cmarginatus*  
*Cexcavatus*  
*Hyparrhnetamba*  
*Hanamesa*  
*Heteropogoncontortus*

--ATCT----CGCTATA---CTA---AACG-CAA-TGCATCCGGGG---C  
--ACAT-T---AGTTGTT---CTC---GGTG-CAA-TGTCCTGCAC---G  
--ACAC-T-CAGGTGTT---CTC---TGTG-TAA-TGCCTTGGCAA---G  
--ACATCG-TAGGTGTT---CTC---TGTG-CAA-TATCGCCACAT---G  
--ACAT-G-AAGTTGTT---CTC---ACTG-CAA-CATCGCATCAT---G  
--ACGT-T-TAGTTGTT---CTC---GGTG-CAG-TGTCTCGGOTT---G  
--ACAT-C-T-GTTGTT---CTC---AGTG-TAA-TGTCTCGCAA---G  
--ACAT---TAGTTGTT---CTC---GGTG-CAA-CATCCCGGTAT---A  
--ACTC-A-AAGTTGTT---CTCGTTAGTG-CAG--GCCTCGGCAC---G  
--ACTC-A-AAGTTGTT---CTCGTTAGTG-CAG--GCCTCGGCAC---G  
--ACAC-T-TAATTGTT---CTC---TGTG-CAA-CACCCC-ACAC---G  
--ACAC-T-TAATTGTT---CTC---TGTG-CAG-CGCCCC-ACAC---G  
--ACAC-T-TAATTGTT---CTC---TGTG-CAG-CGCCCCGACAC---G  
--ACATCA---AGTTGTT---CTC---GGTG-CAN-CGCCTCNGCAC---G  
C-ACATCA---AGTTGTT---CTC---TGTG-CAN-CGACTCGGCAC---G  
--ACACTC---ATTGTT---CTC---NTG-CAG-CGCCCCGGCAC---G  
--ACATC---AGATGTT---CTC---TGTG-CAATGCCCCCGTCAC---G  
--ACATC---AGTTGTT---CTC---GGTGCA--TGCCCCGGTCAC--GG  
C-ACATC---AATTGTT---CTCT--CGTACCA--TACCCNCTNCGC-G-  
C-ACATC---AGATGTT---CTC---TGTGACAC-TGACCCCTCAC--GG  
AGGCATCA---CGTTGTT---CTCG--ACTGACAAAGCCCCGGGAC---G  
CCACATCA---AGATGTT---CTC---TGTGACAGACGACGCGCTCGCGG  
CCACATC---AGTTGTT---CTC---TCTGACACGACGCGGCTCGCGG  
CCA-ATGC---AGATGTT---CTC---GGTG-CACACGACGCGGTCA--GG  
CCA-AGG---CAG-ACGTTCTCTC---TGACACACGACGCC-TCA---GG  
CCA-ATG---AAG-ATGTTCTCTC---TG-CACAAAAGCGG-TCAT--GG  
CCACAGG---CAG-CGGTCCCTC---AG---CCAGAAGGCC-TCG---TG  
CCA-AGA---AAA-ACTTCCCTC---AA---CAAAAAGCCT-TCG---GG  
CCCACGT---CAA-AC-TTCTCTC---GG---CGACATTGCC-TCT---GC  
CCC-ACG---TAA-ACTTTCTCTC---AA---CAAAAACCTCC-TCG---TG  
CCACAG---CAG-ACTTTTCTTA---AT---CCAAAACCC-TCG---GC  
CCA-AGT---CAG-ATGTTCTCTC---TG---CACAGAAGCC-TCAC--GG  
CCA-AGG---CAGTACGTTCTCTT---TGACACATGAAGGNGTCAC--GG  
--ACATC---AGTTGTT---CTC---GGTG-CAG-TGCCCCGGCACA--G  
--ACACTC---AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCACA--G  
--ACATC---AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCGC---G  
--ACATC---AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCACA--G  
--ACATC---AGTTGTT---CTC---GGTG-CAG-CACCCCGGCACAC--G  
--ACAACG-ATGTTTTT---GGTG-CAG-CACCCCGGCAC-GCG  
--ACAAN---AGTTGTTNTT---GGTN-NNT-TACTCCGGCGG-GCG  
--ACATC---AGTTGTTTTT---GGTG-CAG-TACCCCGGCACA-CG  
--ACA---AAAATTTTTTC---GGTT-CAN-TTCCCGGCAC-GCG  
--AAATC---AATTTTT---CTC---GGTN-CAG-NGCCCGGCAC-GCG  
--ACATC---AATTGTTTTC---GGTT-CAA-TTCCCGGCAC-GCG  
--ACATT---AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCAC-NC-

*B. radicans* 4831  
*H. quarrei* 6586  
*H. dissoluta* 5154  
*H. rudis* 6576  
*H. anamesa* 4841  
*E. muticus* 6954  
*CoixBuckler*  
*Arundinellaneपालensis*  
*Pennisetumsetaceum*  
*Panicumbisulcatum*  
*Chimonobambusabrevinoda*  
*Danthoniacalifornica*  
*Eragrostisdielsii*  
*Aristidaharvardii*  
*Zeadipto*  
*ZmaysGfl*  
*Cleistachnesorghoides*  
*I. cylindrica* 6632  
*D. filliformis* 5874  
*T. spicatus* 4846  
*P. franksae* 4705  
*C. plurinoides* 6404  
*A. eucomus* 5869

CGCCTAT--TGTATTCAACCTTACG-ATCCATTACGAAACT---CAGCAC  
CGGCTAT--TGTATTGGA-----  
-----  
ANTTCNAANNTAATGGACCCATCAGCACATTAAGCGCATCACC-----  
-----  
TTATCTGCCCTAATGACCCATTGTGCGACCTTCAGCCATTGCCCTNNG  
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CGGCCAG--CGTCTGGGCC--TAAGACCCATGGA-GCACC--GCAGCGC  
CAGCTGG--CTTATTGCC--TAAGGACCCATGAC-G-ACC--GAAGCGC  
TAGCTAG--CTTGTGGCCCTTAAGGACCCAGGA----ACC--GTAGCGC  
TAGCCGG--CGCAGCGGCCAG--GGACCCATCG----ACC--GGAGCGC  
CAGCCGG--CGAGACGGCCCTCA--GGACCCCTTCG-ATACC--GCAGCGC  
TATCAGGTACGGGAATGGCCCTCA-TGACCCATGA----ACC--GAAGCGC  
CAGCCGG--CGAGAGGGCCCTGTACGACCCATCG----ACC--GCAGCGC  
TGGCCGG--CCATTCGGGCC--TACGGACCCATCG----AGC--G-ACCGA  
CGGCCGG--CCATTCGGGCC--TAAGGACCCATCG----AGC--G-ACCGA  
CGGCCGG--CGATTCGGGCC--TAAGGACCCATGGA-GCACC--GCAGCGC  
C-GCCAG--TGCCTGGCC--TAA-----  
CGNCCAGGTTGCGTANGCC-----  
CGGCCGGC--CGTAGGCC--TAAGGACCCATCGA-GCACC--GAAGCGC  
CTGCCGA-AGCGATTGACCCTAAGGGACCCATCTA-CCACC--GCAGCGC  
CTTCCG---GCGCTCCGCCCTAATG-ACCCATCTA-GCACC--TAAGCGC  
AGGCCG---GCGCATCGCCCTAAGG-ACCCATCGA-GCACC--GGAGCGC

*D. filliformis* 5872  
*H. contortus* 4271  
*H. contortus* 4741  
*B. radicans* 4831  
*A. appendiculatus* 6638  
*A. appendiculatus* 6677  
*A. appendiculatus* 6543  
*H. hirta* 6656  
*H. hirta* 6550  
*H. quarrei* 5088  
*B. insculpta* 5681  
*T. triandra* 6555  
*Spies* 6620  
*H. pilgeriana* 4738  
*A. appendiculatus* 6625  
*Bothriochloa* 7489  
*H. longofolium* 6560  
*H. longofolium*  
*Panicum schinzii* 6652  
*B. serata*  
*S. halepense* 9318  
*Papalum dilatatum* 6617  
*Paspalum*  
*Brachiaria*  
*Aristida adscencianis* 6713  
*H. contortus* 6628  
*T. leucothrix* 6627  
*S. halepense* 9319  
*S. bicolor* 6588  
*S. bicolor* 6589  
*E. muticus* 5074  
*E. muticus* 6715  
*M. capensis* 6655  
*C. prolixus* 6682  
*C. excavatus* 5877  
*C. plurinodum* 6406  
*H. anamesa* 4807  
*A. amethystinus* 6685  
*I. cylindrica* 5114  
*I. cylindrica* 6582  
*Monocymbium*  
*A. distachyos* 6684  
*B. bladhii* 5173  
*S. nigrostis* 6567  
*H. fillipendula* 5144  
*H. tamba* 6585  
*A. schirensis* 6581  
*E. villosa* 5085  
*C. excavatus* 6577  
*C. marginatus* 4397  
*Evillosa*  
*Saccharum officinarum*  
*Imperata cylindrical*  
*Bothriochloa insculpta*  
*Dichanthium annulatum*  
*Cytopogon prolixus*  
*Cplurnodum*  
*Cmarginatus*  
*Cexcavatus*  
*Hyparrhnia tamba*  
*Hanamesa*  
*Heteropogon contortus*

CGGCGC-----ATCTGCCCTAAGG-ACCCATCGA-GCACC--TGAGCGC  
CGGACGG-CGA-ATCGGCCCTCAGG-ACCCATCGA-GCACC--GCAGCGC  
CGGACGG-CAA-ATCGGCCCTTAGG-ACCCATCGA-GCACC--GCAGCGC  
CGGCTAG-CGT-ATTGGCCCTTACG-ACCCATTGA-GCACC--GAAGCGC  
CGGCTAT-CGT-ATTGGCCCTTA-G-ACCCATTGA-GCACC--TAAGCGC  
CGGCTAT-CGT-ATTGGCCCTTACG-ACCCATTGA-GCACC--TAAGCGC  
CG-CTAT-CGT-ATTGGCCCTTACG-ACCCATTGA-GCACC--TAAGCGC  
TGGTCCG-TGT-GTCTGCCCTAAGG-ACCCATCGA-GCACC--TAAGCGC  
TGGTCCG-CTT-GTCCGCCCTAAGG-ACCCATCAA-GCACC--NAAGCNC  
CGCTCCG--GT-ATTNGCCCTAAG--ACCCATCGA-GCACC--TAAGCGC  
CGGCCGG-CGT-ATCGGCCCTGAGG-ACCCATCAA-GCACC--TAAGCGC  
CGGCCGG-CGT-ATNTGCCCTTATG-ANCCATCTA-GCACC--GNAGCGC  
CGGCCGG-CGC-ATTGGCCCTAAGG-ACCCATCTA-CCACC--GCAGCGC  
CGGCCGG-CGC-GACGCCCTAAGG-ACCCATCGA-GCACC--GAAGCGC  
CGGCCGG-CGC-GACGCCCTAAGG-ACCCATCGA-GCACC--TAAGCGC  
CGGCCGG-CGT-ATCTGCCCTAAGG-ACCCATCGA-GCACC--GAAGCGC  
GTATTCC-ACTCAATGCCCTTCAATG-GACCCT--A-GAACC--G-AGCGC  
GTATCCG-ACGCAATGCCCTCAATG-GACCCT--A-TTACC--G-AGCGC  
CAGCTAT-CTCCATAACCTCAAGGG-ACCCATTTA-CGACC--GTAGCGC  
TACCTGG--CTTG-TCCGCTCTAAG-ACCCATAAA-GGACC--TAAGCCA  
TACCTA--TCTTATTGGCCTTAA--ACCCATATG-TGAAC--TAAGCGT  
TACCTGG-TACTATTGGCCTTACG--ACCCATCAT-CGACC--TCAGCGC  
CAGCTG--GCTTATTGGCCTTAA--ACCCATAGA-CGACC--GAAGCGC  
GAACGT--GCTGTTGGCCTTAAA--ACCCAT-GA-GCACC--TATAGCAC  
-AGCTA--TCTTGTGGCCTTAA--ACCCATACT-TGACC--GCAGCGC  
CANCCG--GCGCAGCGGCCCTAAGG-ACCCACACG-CGACC--GCAGCGC  
CAGCCG--GCGCAGCGGCCCTAAGG-ACCCACACG-CGACC--GCAGCGC  
TGGCCG--GTGCATCTGCCCTAAG-ACCCCTTCNG-GCACT--TAAGCCC  
TGGCCG--GTGCATCTGCCCTAAG-ACCCCTTCGG-GCA-T--GC-GCCC  
TGGCCG--GTGCATCTGCCCTAAG-ACCCCTTCGG-GCACT--GC--GCA  
CAACCG--CTTATCGGCCCTAAGG-ACCCATTNT-GCACA--CCANNC  
CAACNG--CTTATCTGCCCTAAGG-ACCCATTGT-GCACA--TCAGCGC  
CGGCNGT--TGCATCTGCCCTAAGG-ATCCATGGAAGCNC--TCTTAC  
C-GCCGG--TCCGACTGCCCTAAGG-ACCCATCGA-GCAAT--TAAGCAC  
CGGCCGTG--CGTCTGCCCTAAG-G-ACCCATCGA-GCACC--AAGCAC  
CGGCTT--CGCTTGACCTAAG-ATCCATCGA-GCACC--TAAGCCC  
CAGCCC--TTGCGACTGCCNTAAGG-ATCCATCA--GCAAT--TAAGCAA  
CAACTTGGCGTTATCGAAACCTAATGACTCATCGAATTA-----T-GCAC  
CGGCCGTTGGCGTTTGGCCCTATCGGATCCATCGA-GCAAT--TCAGCGC  
CGGCCAGTTGGCGTTGAACTATCG-ATCCATCGA-GCAAT--TCAGCAC  
CGCAAGGTTGATTTGAAATAATGGATGGATCGA-GCACAT-TAAGCGC  
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CG-CAACTTGTATCTGAAA-CTATGAATCCATCGAGCA--ATTATGCAC  
CG-CAACTTTCGT-TCAA-AACTCGATGGTTTGGCGG--ATTCTGCACA  
CC-CAGTTT-----  
TG-CAGCTTGTCTTCTGCGCAATCGATGGATCTAGAGC--ATTATGCAC  
CG-CAACTTGCAT-TCAANN-----  
AC-CTACTTACGT-TCNNA-AACTCGATGGTTGCGGN--ATTGAGCAA  
CG-CAACTTGCAT-TCAAC-CTCTCGATCGATCGAGCAC--ATTGAGCAC  
CGGCAACTTGGANTGAAA-----  
CATACGGCGC-ATCGG-----  
CGGCCGGCG-C-ATCGG-----  
CGGCCGGCG-TG-TCGGCCCTAAGG-ACCCATCGA-GCACC--GCAGCGC  
CGGCCGGCG--CATCGGCCCTAAGG-ACCCATCGA-GCACC--GAAGCGC  
--ACCGCG--CATCGGCCCTAAGG-ACCCATCGA-GCACC--GAAGCG--  
--ACAGCG--TATTGTGCTAAGG-AGGCCATGGAGCGAAC-GAAGCGC  
-AACCGAG--ACCTGTCTCCAGG-AGGGATCGA-GCAGC--GAAGGGC  
--ACCGCG--CATGTGCTAAGG-AGGCCATCGAGCACC--GAAGCGC  
--GCCGGCG--TATTGTCTAAGG-AGGCCATCGAGCACC--GAAGCGC  
--GCCGGCG--CATCGGCCCTAAGG-ACCCATCGA-GCACC--GAAGCGC  
--GCCGGCG--TATTGGCCCTAAGG-ACCCATCGA-GCACC--GAAGCGC  
-AGAGCGCA--AATTGGCCCTTAGG-ACCCATTGA-GCACC--GNAGCGC

*B. radicans* 4831  
*H. quarrei* 6586  
*H. dissoluta* 5154  
*H. rudis* 6576  
*H. anamesa* 4841  
*E. muticus* 6954

ATCACC-CCTCGGAT--CGCAACCCCT--NCCNTNTTCG-ACTACCCGA  
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-----  
ACCCAATCCCGACA-----

CoixBuckler  
 Arundinellaneapalensis  
 Pennisetumsetaceum  
 Panicumbisulcatum  
 Chimonobambusabrevinoda  
 Danthoniacalifornica  
 Eragrostisdielsii  
 Aristidaharvardii  
 Zeadiplo  
 ZmaysGfl  
 Cleistachnesorghoides  
 I. cylindrica 6632  
 D. filliformis 5874  
 T. spicatus 4846  
 P. franksae 4705  
 C. plurinoides 6404  
 A. eucomus5869  
 D. filliformis 5872  
 H. contortus 4271  
 H. contortus 4741  
 B. radicans 4831  
 A. appendiculatus 6638  
 A. appendiculatus 6677  
 A. appendiculatus 6543  
 H. hirta 6656  
 H. hirta 6550  
 H. quarrei 5088  
 B. insculpta 5681  
 T. triandra 6555  
 Spies 6620  
 H. pilgeriana 4738  
 A. appendiculatus 6625  
 Bothriochloa 7489  
 H. longofolium 6560  
 H. longofolium  
 Pannicum schinzii 6652  
 B. serata  
 S. halepense 9318  
 Papalum dilatatum 6617  
 Paspalum  
 Brachiaria  
 Aristida adscencianis 6713  
 H. contortus 6628  
 T. leucothrix 6627  
 S. halepense 9319  
 S. bicolour 6588  
 S. bicolour 6589  
 E. muticus 5074  
 E. muticus 6715  
 M. capensis 6655  
 C. prolixus 6682  
 C. excavatus 5877  
 C. plurinodus 6406  
 H. anamesa 4807  
 A. amethystinus 6685  
 I. cylindrica 5114  
 I. cylindrica 6582  
 Monocymbium  
 A. distachyos 6684  
 B. bladhii 5173  
 S. nigrostis 6567  
 H. fillipendula 5144  
 H. tamba 6585  
 A. schirensis 6581  
 E. villosa 5085  
 C. excavatus 6577  
 C. marginatus 4397  
 Evillosa  
 Saccharumofficinarum  
 Imperatacylindrical  
 Bothriochloainsculpta

ACCGTC-GCTC-GGA-CCGCGA-----  
 ATCGCC-GCTC-GGA-CC-----  
 TTTGTC-GCTC-GGA-CC-----  
 TTCGTC-GCTC-GGA-CC-----  
 A-CGTC-GCTC-GGA-CC-----  
 T-CGTC-GCTC-GG-----  
 A-TCTC-GCTC-GGA-CC-----  
 A-AGCC-GCTC-GGA-CC-----  
 GCTTGT-CCTC-GGA-CCGCGA-----  
 GCTTGC-CCTC-GGA-CCGCGA-----  
 ACCGCC-GCTC-GGA-CCGCGACCCAG--GTCAGTCGGG-ACT-----  
 -----  
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 ATCACC-CCTC-GGA-CCGC-ACCCCA--GTCATTCTGG-ACTACCCGC  
 ACCGCC-GCTC-GGA-CCGCAACCCAG--GTCATTCTGG-ACTACCCGC  
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 ATCCC--GCTC-GGA-CCACGACCCAG--GTCAGTCTGG-ACTACCCGC  
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 CTCGCC-GCTC-GGA-CCGCGACCCAG--GTCAGTCTGG-ACTACCCGC  
 TTTTCC-GCTTTGGA-ACGCGACCC-AN--GTCA-TCTGG-ACTACCCGC  
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 ATCGCC-GCTC-GGA-CCGCGACCCAG--GTCAGTCGGG-ACTACCCGC  
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 TTGGCC-GCTC-GGA-CCGCGACCCAG--GTCATTCTGG-ACTACCCGC  
 ACAAC--CCTT-GTA-TCGCGACTCCCT--GTCATGCTGG-ACTACCCGC  
 ACA---CCTT-CTA-CCGC-ACCCCAT--GTCA-GCTGG-ACTACCCGC  
 CTTGG-TGCTC-GGA-CCGCGACCCAG--GTCAGTCGGG-ACTACCCGC  
 CTTAACTGCTC-GGA-CCGCGACCCCAT--GTCATGCTGG-ACTACCCGC  
 TTCCT-CGCTC-GGA-CCGC-ACCCCAT--GTCATACTGG-ACTACCCGC  
 TTGTA-CACTC-GGA-CCGC-ACCCCAT--GTCATGCTGG-ATTACCCGC  
 TTTGT-CGCTC-GGA-CCGCGACCCAG--GTCAGACGGG-ACTACCCGC  
 TTTGT-TGCTC-GGA-CCGCGACCCCAT--GTCATACTGG-ACTACCCGC  
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 CAGCC--GCTC-GGA-CCGCGACCCAG--GTCAAACGGG-ACTACCCGC  
 CAGCC--GCTC-GGA-CCGCGACCCAG--GTCAGACGGG-ACTACCCGC  
 CATCCATGCTC-GGA-CCGCGACCCAN--GTCAGTCNGG-ACCACCCGC  
 ANTCCATGCTC-GGA-CCGCGACCCAN--GTCANNNGG-ACCACCCGC  
 CATCGATGCTC-GGA-CCGCGACCCAG--GTCAGTCTGG-ACCACCCGC  
 ATTGCC-NCTC-GGA-CCNACCCAG--GNNAGCCC-----  
 ATTTCC-CCTC-GGA-TCGCGATCCCCT--GTGATTTTGG-ACTATACGC  
 ATCACC-GCTC-GGATCCGA--CCCAT--GTCATTCTGG-ACTACCCGC  
 ATCGCC-CCTC-GGA-CCGCAACCCCAT--GCCATTCTGGTACTACCCGC  
 ATCACC-GCTC-GGA-CCGC-ACCCCAT--GTCATTCTGG-ACTACCCGC  
 ATCNCC-NCTC-GNA-TCNCCACCNAT--GTCAACTCTGG-ACTACC-GN  
 NATCNCCCCTC-GGA--CGCAANCCCT--GNCATNTTGN-ACTANNCGN  
 TTCATC-CCTC-GGA-TCGCAAT-----  
 ATCGCC-GCTC-GGA-TCAGATCCCAT--GTCATCCTGG-ACTATCCGC  
 ATCNCC-GCTC-GTATCACAATCTCAT--GTCATTCTGG-ACTACCCGC  
 TTCGCC-GCTC-GGA-TCGCAATTCAT--GTCATTTTGGTACTATCCGC  
 TTCACACC-TC-GTA-TCGCAATTCGCT--GCCATCTTCAT-CTATCCGA  
 TTCNCCCCTC-GGA-TCGCAATTCAT--GACATTTTGG-ACTATCCGA  
 TTCACACCAT--GTA-TCGCAATTTGCT--ACGTTCTTCAT-ACATGCCA  
 -----  
 TTCACCGCTCG-GTA-TCGCAATTTCTCT--GTCATCTTCGT-CGACCCGA  
 -----  
 TTCACTGCAGG-GGA-TCGCAATTTCCATANCGTTCTTCATACTATNCCA  
 TTCACCCC-TC-GTA-TCGCAATTCGCT--GCCATCTTCAA-CTATGCCA  
 -----  
 -----  
 ATCGCC-GCTC-GGA-CC-----  
 TTACGCCGCTC-GGA-CCGCGA-----

<i>Dichanthiumannulatum</i>	ATCACC-GCTC-GGA-CCGCGA-----
<i>Cytopogonprolixus</i>	ATCCCC-GCTC-GGA-CCGCGA-----
<i>Cplurnodus</i>	ATCCCC-GCTC-GGA-CCGCGA-----
<i>Cmarginatus</i>	ATCCCC-CCTC-GGA-CCGCGA-----
<i>Cexcavatus</i>	ATCGCC-GCTC-GGA-CCGCGA-----
<i>Hyparrhnetamba</i>	ATCNCC-NCTC-GGA-CCGCGA-----
<i>Hanamesa</i>	ATCGCC-GCTC-GGA-CCGCGA-----
<i>Heteropogoncontortus</i>	ACAGCC-GCTC-GGA-CCGCGA-----

**Appendix G: CLUSTAL W (1.74) multiple sequence alignment data of the Andropogoneae specimens for *ITS1* + *5.8S* gene + *ITS2* region.**

Aameth6685  
Aappen6543  
Aappen6946  
Adist6684  
Aeuco5869  
Aristidaharvardii  
Arundinellaneapalensis  
Aschiren6581  
Bbladh5173  
Binsc5681  
Bradi4784  
Cdieter6642  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Dfili5874  
Emuti5074  
Emuti6715  
Eragrostisdielsii  
Eviol5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdis5145  
Sbicolour9319  
Hfili5144  
Hhirt6656  
Hlongi6560  
Hpilg4603  
Hpilg4738  
Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Panicumbisulcatum  
PentasthisaristifoliaSpies6  
Pfrank4705  
Sbic6588  
Sbic6589  
Shalep9318  
Tspic4846  
Ttria6555  
Zeadiplo  
ZmaysGfl  
Eviollosa  
Saccharumofficinatum  
Imperatocylindrica  
Bothriochloainsculpta  
Dichanthiumannulatum  
Cybopogonprolixus  
Cplurnodus  
Cmarginatus  
Cexcavatus  
Hyparrhrietamba  
Hanamesa  
Heteropogoncontortus

TCGTGACC-GCGTCGATGACC--GT-AACAAA-GTTTCCCTCAGTT--G  
TCGTGACC-----GAAGGAATTC-GT-AACAAA-GTTTCCCTCAGTT--C  
TCGTGACC-----GAGGAATTC-GT-AACAAA-GTTTCCCTCAGTT--C  
TCGTGACAC---GAAAGAAA-C-GT-AACAAACGCTTCCCTCAGTT--C  
TCGTGACCCA--GAAA-AATCC-GT-AACAAG-GCTTCCCTCATGCATG  
TCGTGACCC---TGACCAAAACAGACCGCAACGTGTCCCTG---C  
TCGTGACCC-TTAAACAAAACAGACCGCAACATGTTCCCATGT---  
??  
TCGTGACCC--TGAAGGAATTCAG--ACCGAAGCTTCCCTTAAT--G  
TCGTGAGC---GGAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT---  
TCGTGAACCT--GAAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G  
TCGTGACCCCG--ACGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G  
TCGTGACCC--CTTAAACAAAACAGACCGCAACGTGTCTCTCT--GT--G  
TCGTGACAATTCGAAGGAATTCG-GT-AACAAGAGCTTCCCTCAGTT--C  
TCGTGACCC--ATGTAGATATTC-GT-AACAAA-GCTTCCCTCAGTT--C  
TCGTGACCC---GGTTAAACCG-GT-AACAAA-GCTTCCCTCCCTT--G  
TCGTGACCC---GAAACAAAACCGACCGCAACGTGTCCCTCAT---C  
TCGTGACCGA--GGA---ATTC-GT-AACAA--GCTTCCCTCAGTT--G  
TCGTGACCCG-TGAAAAAACC-CA-AACAAGAGCTTCCCTCAGTT--C  
TCGTGAAC---GGAGGAATTC-GT-AACAAA-GTTTCCCTCAGT---G  
TCGTGACCC---GAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G  
TCGTGACCC---TGACCAAAATAGACCGCAACATGTCTCATCCATG---C  
TCGTGACCC---TGAAACAAATC-GT-AACAAA-GCTTCCCTCCTTTG--T  
TCGTGAGCC---GGAGGAATTC-GT-AACAAA-GTTTCCCTCCTCAT--G  
TCGTGACCC---GAGGAATTC-GT-AACAAA-GCTTCCCTCCTCAT--G  
TCGTGACCC--TGAAAGAAATC-GA-AACAAA-GCTTCCCTCTGT--G  
TCGGTACT---TGGAGGAATTC-GT-AACAAA-GCTTCCCTCTGAT--G  
TCGTGACCC---TTAAACAAAACAGACCGTGAACATGTCTCTCA----G  
TCGTGATAGC---GGAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G  
TCGTGACG---AGGAAAATTC-GT-AACAAA-GCTTCCCTCAGTT--G  
??  
TCGTGACCC---GGAGGAATTC-GT-AACAAA-GTTTCCCTCCGTT--G  
TCGTGACCC---GGAAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--C  
TCGTACCC---TGGAAGAAAACAGA-AACAAA-GCGTCCCTCTCAT--G  
TCGTGACCC--TTAAACAAAACAGACCGCAACGTGTCTCTCATGT---  
TCGTGACCC---GGAGGAATTC-GT-AACAAA-GCTTCCCTCTGAT--G  
TCGTGACCC---TGAAAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G  
TCGTGACC---GGAAGA-ATTC-GT-AACAAA-GCTTCCCTCAGTT--C  
TCGTGAGGC---GAAGGATT-C-GT-AACAAG-GTTTCCCTCAGTT--G  
TCGTGACCC--TTAACCAAAACAGACCGTGAACATGTCTCATCCATG---  
TCGTGACCC---GAAACCAATGCAGACCGTGAACATGCACATGTT---C  
TCGTGAC---GAGGAATTC-GT-AACAAA-GTTTCCCTCAGTT--G  
TCGTGACC---TTAAACAAAACAGACCGTGAACATGTCTCTCATGT---  
TCGTGACCC---AGGAGGAATTC-GT-AACACA-GTTTCCCTCAGTT--C  
TCGTGACAC---GAAGGAATTC-GT-AACAAA-GCTTCCCTACGTT--C  
TCGTGAT---GGAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G  
TCGTGACC--GTGAAGGTAACG-CC-AACAAGAGTTTCCCTCAGTT--G  
CTGTGACCC--TTAAACGAAAACAGACCGCAACGAGTACCCCGT---C  
CCGTGACCC--TTAAACAAAACAGACCGCAACGAGTACCCCGT---C  
-----CGTGAACAGTCTCTCTGT---C  
TCGTGACC---TTAAACAAACAGACCGCAACGCGTCCCTCGT---CC  
TCGTGACCC--TTAAACAAAACAGACCGCAACGAGTCTCTCTGT---CC  
TCGTGACCC--TTAAACAAAACAGACCGCAACGTGTCTCTCATGT---C  
TCGTGACCC--TTAAACAAAACAGACCGCAACGTGTCTCTCATGT---C  
TCGTGACCC--TTAAACAAAGCAGACCGTGTCTCTCTGT---C  
TCGTGACCC--TTAAACAAAACAGACCGTGTCTCTCTGT---C  
TCGTGACCC--TTAAACAAAACAGACCGTGTCTCTCTGT---C  
TCGTGACCC--TTAAACAAAACAGACCGTGTCTCTCTCATGT---C  
TCGTGACCC--TTAAACAAAACAGACCGTGTCTCTCTCATGT---C  
TCGTGACCC--TTAAACAAAACAGACCGTGTCTCTCATGT---C  
TCGTGACCC--TTAAACAAAACAGACCGTGTCTCTCATGT---C

Aameth6685

CCCCCGGCTTC-----GGCTCGGC-CAAGGCACCCAT

*Aappen6543* CNCCCNGGCTAC-----GGTTCGGC-CCTGGCTCCCNAG  
*Aappen6946* ACCGCGGGCTAG-----GGCCCCGCACTCGGCCCCCTAG  
*Adist6684* CCCCCTGGCCCTT-----GGTTCAC-CTTGGCCCCCTTT  
*Aeuco5869* CCACCGGGCTCC-----GGCCCCGC-AAAGGCTCCCGAG  
*Aristidaharvardii* CGTCCG-GCGTC-----TCCCACGACGCCCGGC-CAAGGCCCCCGAC  
*Arundinellanepalensis* -CGCCGGACTTC-----GGTCCGGT-TTAGGCCCCCGAG  
*Aschiren6581* ???  
*Bbladh5173* CCGCCGGGCTCC-----GGCCCCGC-CTAGGCCCCCGAG  
*Binsc5681* CCACCGGGCCTC-----GGCC-GGC-ACAGGCCCCCGAG  
*Bradi4784* CCGCCGGGCTCC-----GGCCCCGCCAAAGGCTCCCGAG  
*Cdieter6642* CCCCCGGGCTTC-----GGTCCGGC-CAAGGCCCTCTAG  
*Cleistachnesorghoides* CCGCCGGGCTCC-----GGCCCGGCANNNGNCCCGAG  
*Cmarg4397* CCGCCGGGCTTC-----GGCCCCGC-CAAGGCCCCCGAG  
*Cplur6406* CTCCCNCGCTAG-----GG--CCGG-CTTGGCCCCCTAT  
*Cprolix6682* CCGCCGGGCTTC-----GGTCCGGC-CAAGGCCCCCGAG  
*Danthoniacalifornica* CGGCCACGCGCGGGACTCGTCCCGCCGCGGCCCTAGGCCCGGAC  
*Dfili5872* CCACCGGGCTCC-----GGCCCCGC-AAAGGCTCCCGAG  
*Dfili5874* CCACCGGGCTCG-----GGCCCCGC-AAAGGCTCCCGAG  
*Emuti5074* CCACCGGGCTC-----GGCCCCGA-CAAGGCCCCCGAG  
*Emuti6715* CCACCGGGCTC-----GGCCCCGA-CAAGGCCCCCGAG  
*Eragrostisdielsii* CGTCCG-GTGATGGGGCTTG-CCTCGCTCCCGAC-TTAGGCCCCAGAC  
*Evill5085* CTGCCGGGCTCC-----GGCTCGGC-CGAGGTCCCGAG  
*Hanam4807* TCGCCGGGCTCC-----GGCCCCGC-CAAGGTTCGGAG  
*Hanam4841* TCGTGGGCTTC-----GGCCCCGC-CAAGGTTCGGAG  
*Hcon4741* TCGTCCGAGCTC-----GGCTCGGC--AAGGCCCCCGAG  
*Hdiss5145* TCGCCGGGCTCC-----GGCCCCGC-CAAGGTTCGGAG  
*Sbicolour9319* TCGTCCGAGCTTT-----GGTCCGGCACAAGGTCCCTTG  
*Hfilli5144* TCGCCGGGCTCC-----GGCCCCGC-CAAGGTTCGGAG  
*Hhirt6656* ACCCTGGGCTTC-----GGCCCCGC-CAAGGTTCGGAG  
*Hlongi6560* ???  
*Hpilg4603* CCGCCGGGCTTC-----GGCCCCGC-CAAGGCCCCCGAG  
*Hpilg4738* CCCCCTGGCAAG-----GGCCCCGCACAAGGCCCCCGAG  
*Hquar5088* TCGCCGGGCTCC-----GGCCCCGC-CAAGGTTCGGAG  
*Hrud6576* -CGCCGGGCTCC-----GGCCCCGC-CAAGGTTCGGAG  
*Htamb6585* TCGCCGGGCTCC-----GGCCCCGC-CAAGGTTCGGAG  
*Icylin5114* CCGCCGGGCTCC-----GGCCCCGC-CAAGGTTCGGAG  
*Icylin6632* CCCCCTGGCCAG-----GGTTCGGC-CCCGGCCCTAT  
*Mcape6655* CCCCCTGGCTAC-----GGCTCGGCACAG--CCCCTAG  
*Panicumbisulcatum* CGT-CGGCTAT-----GGCCCCGC-AAAGGCACCCACC  
*PentastichisaristifoliaSpies6* CGGCCACACGA-----GCTCACGCGTGTGGC-CTAGGCCCCCGAC  
*Pfrank4705* CCGCCTGGCTTC-----GGCTGGT-CAAGGCCCCCTGAG  
*Sbic6588* -CGTCCGAGTTC-----GGCTCGGCACAAGGTCCCTTG  
*Sbic6589* ACCNCGGGCTGG-----GGCTCGGC-CTAGGCCCCCTTG  
*Shalep9318* CTCTGGGCTCC-----TGCCCGTG-CAAGGCCCCCGAG  
*Tspic4846* CCACCGGGCTTC-----GGCCCCGC-AAAGGCTCCCGAG  
*Ttria6555* CCCCCTGGCCTC-----GGCCCCGC-CACGGCCCGAG  
*Zeadiplo* CGCCGG-GCTCC-----GGCCCCGCACGCTGCCCCCCC  
*ZmaysGfl* CGCCGG-GCTCC-----GGCCCCGCACGCTGCCCCCCC  
*Evillosa* -GCCGG-GCTTC-----GGCTCGGC-CAAGGCACCYGAG  
*Saccharumofficinatum* -GCCGG-GCTTC-----GGCCCCGCACGAGGTCCCGAG  
*Imperatacylindrical* -GCCGG-GCTTC-----GGCCCCGC-CAAGGCCCCCGAG  
*Bothriochloainsculpta* -GACGG-GCTCC-----GGATCGGA-CAAGGCCCCCGAG  
*Dichanthiumannulatum* -GCCGG-GCTTC-----GGCTCGGA-CAAGGCCTCGAG  
*Cytopogonprolixus* -GCCGG-GCTTC-----GGCTCGGN-CAAGGTTCGGAA  
*Cplurnodus* -GCCGG-GCTTC-----GGCTCGGA-CAAGGTTCGGAA  
*Cmarginatus* -GCCGG-GCTTC-----GGCTCGGC-CAAGGTTCGGAA  
*Cexcavatus* -GCCGG-GCCTC-----GGCCTGAGCAAGGTTCGGAA  
*Hyparrhrietamba* -GCCGG-GCTCC-----GGCTCGGC-CAAGGTTCGGAA  
*Hanamesa* -GTCGG-GCTCC-----GGTCCGGC-CAAGGTTCGGAA  
*Heteropogoncontortus* -GTCGA-GCCTC-----GGTCCGGC--AAGGCCCCCGAT

*Aameth6685* CCTCGTCCC-----GGGGCGGAGGTGCCGC--AAAAGAACCACGGC  
*Aappen6543* CCCCCTCCA-----GGGGCGAAGGGCCGC--AAAACAACCACGGC  
*Aappen6946* CCCCCTCCC-----GGGGCGGAGGGCCGC--AAAACAACCACGGT  
*Adist6684* TCCCCTCCC-----TGGGTAATCGCCCGAA--AA-ACAACCACGGC  
*Aeuco5869* CCCCCTCCC-----GGGGCGGAGGGCCGC--AAAAGAACCACGGC  
*Aristidaharvardii* CTCCTGT-----GGAGGGGAGGGCCGC--AACAGAACCACGGC  
*Arundinellanepalensis* CTCCTCCC-----GGGGCAGAGGGCCCTT--AACAGAACCACGGC  
*Aschiren6581* ???  
*Bbladh5173* CCCCCTCCC-----GGGGCGGAGGAGCCGC--CAAAGAACCACGGC



Binsc5681  
Bradi4784  
Cdieter6642  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Dfili5874  
Emuti5074  
Emuti6715  
Eragrostisdielsii  
Eviol5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdiss5145  
Sbicour9319  
Hfilli5144  
Hhirt6656  
Hlongi6560  
Hpilg4603  
Hpilg4738  
Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Panicumbisulcatum  
PentastichistisaristifoliaSpies6  
Pfrank4705  
Sbic6588  
Sbic6589  
Shalep9318  
Tspic4846  
Ttria6555  
Zeadiplo  
ZmaysGfl  
Evillosa  
Saccharumofficinatum  
Imperatocylindrical  
Bothriochloainsculpta  
Dichanthiumannulatum  
Cybopogonprolixus  
Cplurnodus  
Cmarginatus  
Cexcavatus  
Hyparrhnetamba  
Hanamesa  
Heteropogoncontortus

CTCCGTCTCTCATGTCTCGGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CCCCGTCCA-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CCCTGTAC-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CCNGTNT-----GGGGCGGAGGGGCCAC--AAAAGAACCACCGC  
CCCTGTAC-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CCCCNTCTT-----GGGGTGGAGGGGCCGC--AAAATAACCACCGC  
CCCTGTAC-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CTCCCTCTCC-----GGGAGGGAGGGGCCGT--AAAAGAACCACCGC  
CCCCGTCCC-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CCCCGTCCC-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CTCCGGCA-----GAGGGGCCGC--AACAGAACCACCGC  
CTCCGTAC-----GAGGGGCCGC--AACAGAACCACCGC  
TTCTTAC-----GGAGGGCTGGGGGCCGC--AACAGAATCCACGGC  
CCCCGTCTT-----GGGGCGGAGGGGCCGC--AAAAGATCCACGGC  
CCCCGTCCC-----GGGGCGGAGGTGCCGC--AAAAGAACCACCGC  
CCCCGTCCG-----GGGGCGGAGGTGCCGC--AAAAGAACCACCGC  
CTTC-----GGCAGAGGGGCCGC--CACAGAACCACCGC  
CCCCGTCTCTCATGTCTCGGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CTCCAACCT-----GGGGCAGAGGGGCCAC--AAAAGAACCACCGC  
CCCCGTCCC-----GGGGCGGAGGTGCCGC--AAAAGAACCACCGC  
CCCCGTCCG-----GGGGCGGAGGTGCCGC--AAAAGAACCACCGC  
??  
CCCTGTAC-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CCCCGTAC-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CCCCGTCCC-----GGGGCGGAGGTGCCGC--AAAAGAACCACCGC  
CCCCGTCTCTCATGTCTCGGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CCCCGTCTCTCATGTCTCGGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CTCCGCA-----GGGGCGGAGGGGCCGC--AACAGAACCACCGC  
CTCCGTCCC-----GGGGCGGAGGGGCCGC--AACAGAACCACCGC  
CCCCGTCTT-----GGGGCGGAGGGGCCAC--AACAGAACCACCGC  
CACC-TATT-----AGGAGGATGGGGGCCGC--AAAAGAACCACCGC  
CTCCGCA-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CACCCTCAA-----GGGGC-GAGGGGCCGC--AACAGAACCACCGC  
CTCCAACCT-----GGGGCAGAGGGGCCAC--AAAAGAACCACCGC  
CTCCCTCT-----GGGGCAGAGGGGCCAC--AAAAGAACCACCGC  
CTCC-TCAT-----GGAGGAGAGGGGCCAC--AAAAGAACCACCGC  
CCCCGTCCC-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
CTCCGTCT-----GGGGCGGAGGGGCCAC--AAAAGAACCACCGC  
GAACCTCCC-----GGGGAAGGGGGCCGCGAAAAAGAACCACCGC  
TCCAGTCCC-----GGGGCGGAGGGTGCAGCGAAAAAGAACCAYGGC  
CTCCGTCCC-----GGAGCGGAGGGGCCAC--AACAGAACCACCGC  
CTCCGTCT-----GGGGCGGAGGGGCCGC--AACAGAACCACCGC  
CTCCGTCT-----GGGGGGGGGGGGCC--GTCCGAAAAAGAACCACGGC  
CCCTGTCT-----GGGGGGGGGGGGCC--AAAAGAACCACCGC  
CCCTTCCC-----GGGGGGGAGGGTCTCCC--AAAAGAACCACCGC  
CCC-TTCCC-----GGGGGGGAGGGCNC--AAAAGAACCACCGC  
CCCTTCCC-----GGGGGGGAGGGTCTCCC--AAAAGAACCACCGC  
CCCCTTCCC-----GGGGGGGAGG-TTCCC--AAAAGAACCACCGC  
CCCCCTCCC-----GGGGGGGAGG-TTCCCCAAAAGAACCACCGC  
CCCCCTCCC-----GGGGGGGAGG-TTCCCCAAAAGNAACC-ACGGC  
CTTCGA-CA-----GAGGGG-----CCACCACAGAACCACCGC

Aameth6685  
Aappen6543  
Aappen6946  
Adist6684  
Aeuco5869  
Aristidahvardii  
Arundinellapalensis  
Aschiren6581  
Bblad5173  
Binsc5681  
Bradi4784  
Cdieter6642  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica

GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AAGCGTCAAGGAACACTTA-TATG-CCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATG-CCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCGAA---CGGCGTCAAGGAACACTGTCTATTGCCTG--GC  
GCCAT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
??  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCTT---AGGCGTCAAGGAACACTTA-TATTGCCTT--GC  
GCCGAC---CGGCGCAAGGAACACTGTTTATTGCCTT--GC



*Hcon4741* TCGGTGGAGCGGTTCGGC-CTGCCTTCTGCTCCCCGCGCGGC---GATGAT  
*Hdiss5145* AAGGTGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Sbicolour9319* ACAGTGGAGTGGTTCGGC-NTGCGTTGGGGTCCCGATGAAGAACAAAAGAT  
*Hfilli5144* CCGTGGG--CGGTTCGGC-CTGCCTTTCGCTCCCCGTGCAGC---GATGAT  
*Hhirt6656* CCGGTGGAGCGGTTCGC-TTGCTTCCGCTCCCCGTGCAGC---GATGAT  
*Hlongi6560* ???  
*Hpilg4603* CCGGTGGAGCGGTTCGGC--TGCTTCCGCTCCCCGCGCAGC---GATGAT  
*Hpilg4738* CCGGCGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Hquar5088* CCGGTGGAGCGGTTCGGC-CTGCCTTTCGCTCCCCGTGCAGC---GATGAT  
*Hrud6576* AAGGTTGGAACCAAGGC-TTCCGTGGGGGCACCGATGAAGA-CGGATGAT  
*Htamb6585* AAGTGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Icylin5114* TCGGCGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Icylin6632* TCGGCGGAGCGGCCG-C-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Mcape6655* TCGGCGG-ACGCTTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Panicumbisulcatum* TCGGGTGTGGTTCGGC-TTGCCGGCCACGCTCGTGCAGC---GATGAT  
*Pentaschistis aristifolia Spies6* CGGTGGCGGTTCGGC-CTGCCAGCCGCGCCGCGCAGC---GATGAT  
*Pfrank4705* TCGACGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Sbic6588* GCAGTGGAG-GCTTCGGC-CTCCGTTGGTGCCGCGATGAAGAACGCAAGAT  
*Sbic6589* ACA-CGAGTGGTTCGGC-CTGCCTTCCGCTCCCTGAGCAGC---GATGAT  
*Shalep9318* TCGGGGCTTTGGTTCGGC-TTGCCGATCATGCCCTCGTGCAGC---GATGAT  
*Tspic4846* CCGGCGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Ttria6555* TCGGCGGAGCGGTTCGGC-TTGCTTCCGCTCCCCGCGCAGC---GATGAT  
*Zeadiplo* CCGGCGGAGCGGTTCGGC-CCGCTTCCGCTCCCCGGGCGAGC---GGTTA-  
*ZmaysGfl* CCGGCGGAGCGGTTCGGC-CCGCTTCCGTTCCCGAGGCGAGC---GGTTA-  
*Evillosa* --GGCGGAGCGGTTCGGC-TTGCTTCCGCTCCCCACGCGGT---GATCAT  
*Saccharum officinarum* --GGTGGAGCGGTTCGGC-CCGCTTCCGCTCCCCGCGCAGC---GATGAT  
*Imperata cylindrical* --GGCGGA-CGGTTCGGC-CTGCCTTCCGCTCCCCGCGCAGA---GTGCT  
*Bothriochloa insculpta* --GGCGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Dichanthium annulatum* --GGTGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Cybopogon prolixus* --GGTGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Cplurnodus* --GGTGGAGNGGCTTCGGC-CTGCCTTCCGCTCACCAGCAGC---GATGAT  
*Cmarginatus* --GGTGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Cexcavatus* --GGTGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Hyparrhrietamba* --GGTGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Hanamesa* --GGTGGAGCGGTTCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Heteropogon contortus* --GGTGGAGCGGTTCGGC-CTGCCTTCTGCTCCCCGCGCGGC---GATGAT

*Aameth6685* ATAC-TAATC  
*Aappen6543* ATC-TTAATC  
*Aappen6946* ATC-TTAATC  
*Adist6684* ATCTCTAATC  
*Aeuco5869* ATC-TTAATC  
*Aristidaharvardii* ATCA-TAAC-  
*Arundinella nepalensis* ATG--TAAC-  
*Aschiren6581* ??????????  
*Bbladh5173* ATCTTTAATC  
*Binsc5681* ATC-TTAATC  
*Bradi4784* ATC-TTAATC  
*Cdieter6642* ATC-TTAATC  
*Cleistachnesorghoides* ATC-TTAATC  
*Cmarg4397* ATC-TTAATC  
*Cplur6406* ATC-TTAATC  
*Cprolix6682* ATC-TTAATC  
*Danthoniocalifornica* ??????????  
*Dfili5872* ATC-TTAATC  
*Dfili5874* ATC-TTAATC  
*Emuti5074* ATC-TTAATC  
*Emuti6715* ATC-TTAATC  
*Eragrostis dielsii* TTTTGTAAATC  
*Eviol5085* ATC-TTAATC  
*Hanam4807* ATC-TTAATC  
*Hanam4841* ATC-TTAATC  
*Hcon4741* ATC-TTAATC  
*Hdiss5145* ATC-TTAATC  
*Sbicolour9319* ATCA-TCC--  
*Hfilli5144* ATC-TTAATC  
*Hhirt6656* ATC-TTAATC  
*Hlongi6560* ??????????  
*Hpilg4603* ATC-TTAATC  
*Hpilg4738* ATC-TTAATC

<i>Hquar5088</i>	ATC-TTAATC
<i>Hrud6576</i>	ATC-TTAATC
<i>Htamb6585</i>	ATCTTTAATC
<i>Icylin5114</i>	ATC-TTAATC
<i>Icylin6632</i>	ATC-TTAATC
<i>Mcape6655</i>	ACC-TTAAAC
<i>Panicumbisulcatum</i>	ATC-TTAATC
<i>PentaschistisaristifoliaSpies6</i>	ATAC-TAATC
<i>Pfrank4705</i>	ATC-TTAATC
<i>Sbic6588</i>	ATC-TTAATC
<i>Sbic6589</i>	ATC-TTAATC
<i>Shalep9318</i>	ATC-TTAATC
<i>Tspic4846</i>	ATC-TTAATC
<i>Ttria6555</i>	TTA-TTAATC
<i>Zeadiplo</i>	CACCTTAGTC
<i>ZmaysGfl</i>	CACCTTAATC
<i>Evillosa</i>	ATAC-TAATC
<i>Saccharumofficinatum</i>	ATCT-TAATC
<i>Imperatacylindrical</i>	ATC-TTAATC
<i>Bothriochloainsculpta</i>	ATAC-TAATC
<i>Dichanthiumannulatum</i>	ATAC-TAATC
<i>Cytopogonprolixus</i>	ATCC-TAATC
<i>Cplurnodus</i>	ATCC-TAACC
<i>Cmarginatus</i>	ATCC-TAACC
<i>Cexcavatus</i>	ATC-TTAATC
<i>Hyparrhrietamba</i>	ATCC-TAATC
<i>Hanamesa</i>	ATC-TTAATC
<i>Heteropogoncontortus</i>	ATC-TTAATC
<i>Ameth6685</i>	AACACG-ACTCTCGG-CAACGGATATCT
<i>Aappen6543</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Aappen6946</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Adist6684</i>	CACACG-ACTCTCGGTCAACGGATATCT
<i>Aeuco5869</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Aristidaharvardii</i>	AACACG-ACTCTCGG-CAACGGATATCT
<i>Arundinellaneapalensis</i>	AACACG-ACTCTCGG-CAACGGATATCT
<i>Aschiren6581</i>	????????????????????????????????
<i>Bbladh5173</i>	CACACGAAGTCT--GG-GAAGGTATATCT
<i>Binsc5681</i>	CACAC--ACTCTCG--CAACGGATATCT
<i>Bradi4784</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Cdieter6642</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Cleistachnesorghoides</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Cmarg4397</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Cplur6406</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Cprolix6682</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Danthoniacalifornica</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Dfili5872</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Dfili5874</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Emuti5074</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Emuti6715</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Eragrostisdielsii</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Evill5085</i>	TACATG-ACTATCCG-CAACGGATATCT
<i>Hanam4807</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Hanam4841</i>	CACGCG-ACTCTCGG-CAACGGATATCT
<i>Hcon4741</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Hdiss5145</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Sbicolour9319</i>	-----
<i>Hfilli5144</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Hhirt6656</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Hlongi6560</i>	????????????????????????????????
<i>Hpilg4603</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Hpilg4738</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Hquar5088</i>	CACACC-ACTCTCAG-TAACGGATATCT
<i>Hrud6576</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Htamb6585</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Icylin5114</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Icylin6632</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Mcape6655</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>Panicumbisulcatum</i>	CACACG-ACTCTCGG-CAACGGATATCT
<i>PentaschistisaristifoliaSpies6</i>	????????????????????????????????
<i>Pfrank4705</i>	GACACG-ACTCTCGG-CAACGGATATCT

*Sbic6588*  
*Sbic6589*  
*Shalep9318*  
*Tspic4846*  
*Ttria6555*  
*Zeadiplo*  
*ZmaysGf1*  
*Evillosa*  
*Saccharumofficinatum*  
*Imperatacylindrical*  
*Bothriochloainsculpta*  
*Dichanthiumannulatum*  
*Cytopogonprolixus*  
*Cplurnodus*  
*Cmarginatus*  
*Cexcavatus*  
*Hyparrhrietamba*  
*Hanamesa*  
*Heteropogoncontortus*

????????????????????  
CACACG-ACTCTCGG-CAACGGATATCT  
CACACG-ACTCTCGG-CAACGGATATCT  
CACACG-ACTCTCGG-CAACGGATATCT  
CACACG-ACTCTCGG-CAACGGATATCT  
GACACG-ACTCTCGG-CAACGGATATCT  
GACACG-ACTCTCGG-CAACGGATATCT  
CACACG-ACTCTCGG-CAACGGATATCT  
GACACG-ACTCTCGG-CAACGGATATCT  
CACACG-ACTCTCGG-CAACGGATATCT  
CACACG-ACTCTCGG-CAACGGATATCN  
CACACG-ACTCTCGG-CAACGGATATCN  
CACACG-ACTCTCGG-CAACGGATATCN  
CACACG-ACTCTCGG-CAACGGATATCN  
CACACG-ACTCTCGG-CAACGGATATCN  
CACACG-ACTCTCGG-CAACGGATATCN  
CACACG-ACTCTCGG-CAACGGATATCN  
CACACG-ACTCTCGG-CAACGGATATCN  
CACACG-ACTCTCGG-CAACGGATATCN

*Ameth6685*  
*Aappen6543*  
*Aappen6946*  
*Adist6684*  
*Aeuco5869*  
*Aristidaharvardii*  
*Arundinellaneapalensis*  
*Aschiren6581*  
*Bbladh5173*  
*Binsc5681*  
*Bradi4784*  
*Cdieter6642*  
*Cleistachnesorghoides*  
*Cmarg4397*  
*Cplur6406*  
*Cprolix6682*  
*Danthoniacalifornica*  
*Dfili5872*  
*Dfili5874*  
*Emuti5074*  
*Emuti6715*  
*Eragrostisdielsii*  
*Evill5085*  
*Hanam4807*  
*Hanam4841*  
*Hcon4741*  
*Hdiss5145*  
*Sbicolour9319*  
*Hfilli5144*  
*Hhirt6656*  
*Hlongi6560*  
*Hpilg4603*  
*Hpilg4738*  
*Hquar5088*  
*Hrud6576*  
*Htamb6585*  
*Icylin5114*  
*Icylin6632*  
*Mcape6655*  
*Panicumbisulcatum*  
*PentaschistisaristifoliaSpies6*  
*Pfrank4705*  
*Sbic6588*  
*Sbic6589*  
*Shalep9318*  
*Tspic4846*  
*Ttria6555*  
*Zeadiplo*  
*ZmaysGf1*  
*Evillosa*

CGGCTCTCGCATCGATGAAGAA-----  
CGGCTCTCGCATCGATGAAGAA-----  
CGGCTCTCGCATCGATGAAGAA-----  
CGGCTCTCGCATCGATGAAGAA-----  
CGGCTCTC-----  
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CGGCTCTCGCATCGATGAAGAACGTAGCAAAA  
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CGGCTCTCGCATCGATGAAGAACGTAGCAAAA

*Saccharumofficinatum*  
*Imperatocylindrica*  
*Bothriochloainsculpta*  
*Dichanthiumannulatum*  
*Cytopogonprolixus*  
*Cplurnodus*  
*Cmarginatus*  
*Cexcavatus*  
*Hyparrhietamba*  
*Hanamesa*  
*Heteropogoncontortus*

CGGCTCTCGCATCGATGAAGAACGTAGCAAAA  
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*Aameth6685*  
*Aappen6543*  
*Aappen6946*  
*Adist6684*  
*Aeuco5869*  
*Aristidaharvardii*  
*Arundinellaneopalensis*  
*Aschiren6581*  
*Bbladh5173*  
*Binsc5681*  
*Bradi4784*  
*Cdieter6642*  
*Cleistachnesorghoides*  
*Cmarg4397*  
*Cplur6406*  
*Cprolix6682*  
*Danthoniacalifornica*  
*Dfili5872*  
*Dfili5874*  
*Emuti5074*  
*Emuti6715*  
*Eragrostisdielsii*  
*Evill5085*  
*Hanam4807*  
*Hanam4841*  
*Hcon4741*  
*Hdiss5145*  
*Sbicolour9319*  
*Hfilli5144*  
*Hhirt6656*  
*Hlongi6560*  
*Hpilg4603*  
*Hplig4738*  
*Hquar5088*  
*Hrud6576*  
*Htamb6585*  
*Icylin5114*  
*Icylin6632*  
*Mcape6655*  
*Panicumbisulcatum*  
*PentaschistisaristifoliaSpies6*  
*Pfrank4705*  
*Sbic6588*  
*Sbic6589*  
*Shalep9318*  
*Tspic4846*  
*Ttria6555*  
*Zeadiplo*  
*ZmaysGfl*  
*Evillosa*  
*Saccharumofficinatum*  
*Imperatocylindrica*  
*Bothriochloainsculpta*  
*Dichanthiumannulatum*  
*Cytopogonprolixus*  
*Cplurnodus*  
*Cmarginatus*  
*Cexcavatus*

TGCG-ATACCTAGTGT-GAATTGC  
TGCGGATACGTGGCGT-GAATTGC  
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TGCG-ATACCTGGTGT-GAATTGC  
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ATGCGATACCTAGCGT-CAATTCC  
TGCCGT-CCTAGCGT--NATTCC  
TGCGGATACCTGGTGT-GAATTGC  
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TGCG-ATACCTGGTGT-GAATTGC  
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TGCCGATACCTAGCGT-GAATTCC  
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TGCG-ATACCTGGTGT-GAATTGC  
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TGCGGATACCTGGCT-AAATTGC  
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TGCT-ATACCTCGAGT-GAATTGC  
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TACGGATACCTAGGCT-CAATTGC  
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TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC

*Hyparrhnetamba*  
*Hanamesa*  
*Heteropogoncontortus*

T-CG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC  
TGCG-ATACCTGGTGT-GAATTGC

*Aameth6685*  
*Aappen6543*  
*Aappen6946*  
*Adist6684*  
*Aeuco5869*  
*Aristidaharvardii*  
*Arundinellanepalensis*  
*Aschiren6581*  
*Bbladh5173*  
*Binsc5681*  
*Bradi4784*  
*Cdieter6642*  
*Cleistachnesorghoides*  
*Cmarg4397*  
*Cplur6406*  
*Cprolix6682*  
*Danthoniacalifornica*  
*Dfili5872*  
*Dfili5874*  
*Emuti5074*  
*Emuti6715*  
*Eragrostisdielsii*  
*Eviol5085*  
*Hanam4807*  
*Hanam4841*  
*Hcon4741*  
*Hdiss5145*  
*Sbicolour9319*  
*Hfilli5144*  
*Hhirt6656*  
*Hlongi6560*  
*Hpilg4603*  
*Hplig4738*  
*Hquar5088*  
*Hrud6576*  
*Htamb6585*  
*Icylin5114*  
*Icylin6632*  
*Mcape6655*  
*Panicumbisulcatum*  
*PentaschistisaristifoliaSpies6*  
*Pfrank4705*  
*Sbic6588*  
*Sbic6589*  
*Shalep9318*  
*Tspic4846*  
*Ttria6555*  
*Zeadiplo*  
*ZmaysGfl*  
*Evillosa*  
*Saccharumofficinatum*  
*Imperatacylindrical*  
*Bothriochloainsculpta*  
*Dichanthiumannulatum*  
*Cyobopogonprolixus*  
*Cplurnodus*  
*Cmarginatus*  
*Cexcavatus*  
*Hyparrhnetamba*  
*Hanamesa*  
*Heteropogoncontortus*

ATACATCCCG--CTGAACCATCT--AGTTTTTTAAACGCAAGTTGCGCCC  
ATAAATCCCG--CG-AACCATCG--AGTTTTTTGAA-CGCAAGTTGCGCCC  
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--TAATGTGC--TC-AATGGGTCC--TTATGGCCGAA--TACACCG--GCCGCG  
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AG-AATCCCG--CG-AATCATCG--AGTTTTTTGAA-CGCACATTCGSCCC

*Aameth6685*  
*Aappen6543*  
*Aappen6946*

C-AGGGC-TTCTGGTCTGA-GGACACCTCTTCCTGGCCGC-TCACGCC--  
-GAGGCC-TTCTGGTCTGA-GGGCACCTCTGCCTGGGCG--TCACGCCG--  
???

Adist6684  
Aeuco5869  
Aristidaharvardii  
Arundinellapalensis  
Aschiren6581  
Bbladh5173  
Binsc5681  
Bradi4784  
Cdieter6642  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Dfili5874  
Emuti5074  
Emuti6715  
Eragrostisdielsii  
Eviol5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdiss5145  
Sbicolour9319  
Hfilli5144  
Hhirt6656  
Hlongi6560  
Hpilg4603  
Hpilig4738  
Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Panicumbisulcatum  
PentasthisaristifoliaSpies6  
Pfrank4705  
Sbic6588  
Sbic6589  
Shalep9318  
Tspic4846  
Ttria6555  
Zeadiplo  
ZmaysGfl  
Evillosa  
Saccharumofficinarum  
Imperatocylindrical  
Bothriochloainsculpta  
Dichanthiumannulatum  
Cybopogonprolixus  
Cplurnodus  
Cmarginatus  
Cexcavatus  
Hyparrhnetamba  
Hanamesa  
Heteropogoncontortus

TGACCCCGTTCTTTCACACGACACCAATGTCATGTGCGGTACGCTGAT  
-GAAGCC-TTTTGGTTCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
-GAGGCC-TTCTGGCCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
-GAAGCC-TCCAGGCCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
C-ACGCCTTTTCTTTGAC--GACACAAGTGTGCGCGTACACCAA--  
T-ACCGGGCTCTTTCAC-CGACAACAATGGGA-GTCGCC-CACCACG--  
-GAGGCC-TTCTGGTTCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
T-ATGCC-TTCTGNTCAA-GGGCACCTTGCCTGGGCG--TCACGCCG--  
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-GAGGCC-TTCTGGCCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
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-GACGCC-TTCTGTTCGA-GGGCAGCTTGCCTGGGCG--TAACACC---  
G--AGCC-TTCTGGTC-A-GGGCACCTTGCCTGGGCG--TCACGCCA--  
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-GAGGCC-TTCCGGCCGA-GGGCAGCTTGCCTGGGCG--TCACGCCA--  
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-GA-GCC-TTCTGGTTCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
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-GAGGCC-TTCTGGTTCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
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-GATGCC-TTCTGGTTCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
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C-ACGCC-TTCTGGTTCGA-GGGCACCTTGCCTGGGCG--TCACGCCA--  
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-GAAGCC-TTCTGGCCGA-AGGCACNTTGCCTGGGCG--TCACGCC---  
-GAGGCC-TTCTGGTTCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
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-GAGGCC-TTCCGGCCGA-GGGCAGCTTGCCTGGGCG--TCACGCCA--  
-GACGCC-TTCTGGTTCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
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-GAGGCC-TTCTGGTTCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
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-GAGGCC-TTCTGGTTCGA-GGGCAGCTTGCCTGGGCG--TCACGCC---  
-GAGGCC-TTCTGGTTCGA-GGGTACGCTTGCCTGGGCG--NCACGNC---  
-GAGGCC-TTCTGGTTCGA-GGGTACGCTTGCCTGGGCG--NCACGAC---  
-GAGGCC-TTCTGGTTCGA-GGGTACGCTTGCCTGGGCG--NCACGAC---  
-GAGGCC-TTCTGGTTCGA-GGGTACGCTTGCCTGGGCG--TCACGAC---  
-GAGGCC-TTCTGGTTCGA-GGGTACGCTTGCCTGGGCG--TCACGCC---  
-GAGGCC-TTCTGGTTCGA-GGGTACGCTTGCCTGGGCG--TCACGCC---  
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Aameth6685  
Aappen6543  
Aappen6946  
Adist6684  
Aeuco5869  
Aristidaharvardii  
Arundinellapalensis  
Aschiren6581  
Bbladh5173  
Binsc5681  
Bradi4784

ATAGCAC-ACTCACAAGATGAC--CCTCGGCC--A-CAGATTTGGT-  
AAAACAC-ACTCCCAACCCAA----AATCGGGG--AGGGCATGTG-T-  
??  
GAATGAG-ACGTCGACACGATTTTC-GGCAGACGG--CACACTTTGG-  
AAAAGAC-ACGTCACAGCCAC--AATCAAG----GAGTACTGGC-  
AAAAGAC-ACTCCCACCCACCC--AGGTTG-----GGG-ACGTGGC-  
AAAAGAC-ACTCCCAACCCAAC--CAGGGG----AGGG-ATGTGGC-  
-T--GAC-ACTCACAAAACACCC-T-TGCAGAC-----CCACCTTCTG-  
GC-TGTG-CCCGGCACGGCTATCC--GGCAAC-----CAAACCTGGG-  
AAAAGAC-ACTCCCAACCCAC----CCTCGGGG--AGGGCAGCTGGT-  
AAAATAC-ACTCCCAACCC-AT---ACTCGGGG--AGGG-ATGTTGTG



Cdieter6642  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Dfili5874  
Emuti5074  
Emuti6715  
Eragrostisdielsii  
Eviol5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdiss5145  
Sbicolour9319  
Hfilli5144  
Hhirt6656  
Hlongi6560  
Hpilg4603  
Hplig4738  
Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Panicumbisulcatum  
PentaschistisaristifoliaSpies6  
Pfrank4705  
Sbic6588  
Sbic6589  
Shalep9318  
Tspic4846  
Ttria6555  
Zeadiplo  
ZmaysGfl  
Eviolosa  
Saccharumofficinarum  
Imperatacylindrical  
Bothriochloainsculpta  
Dichanthiumannulatum  
Cybopogonprolixus  
Cplurnodus  
Cmarginatus  
Cexcavatus  
Hyparrhnietaмба  
Hanamesa  
Heteropogoncontortus

??  
AAAAGAC-ACTCCCAACCCACCC--AAAGGGGAG-GAGGG-ACGTGGT-  
-AAATAC-ACTCCCAACC-AATCCTCGGCAGAAG---GACGCTTTGG-  
AAAAGAC-ACTCCCAACAGCACCCCTCGAGCGGCATAATGG-ACCTGAC-  
AAATGAC-ACTCCCAACCCAT----CCTCGGGT---ACGG-ACATGGT-  
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TAATAAC-ACTCACAACATAC----AATCGCGGC---AGC--ACCTTCT-  
ACATTAC-ACTCCGAACCCA----CCCTCGGGG---AGGG-ATGTGGT-  
TC-TGTC---CCCAATCCACTTGT-TGCCAGGC---CCC-ACCTTTG-  
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-AAACAC-GCTCCCAACCCACTAACCCCTGTGGT---GGG-ATGCGGC-  
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GCTGTAC-ACTGCCACAAC-CACTAT-GGCAGTAG--CT--ACCTTCTG  
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AAAACAC-ACTCCCAACCCAC---ACAAGG---AAAGG-ACCTTCT-  
AAAAGAC-ACTCCCAACCCATCC---ATGGGGT---GGG-ACGTGGT-  
??  
--GTACC-CGCTCAGATACTAG---CAGCAA---ATAGG-ACTTCT-  
ACAGTAC-ACTCCCAACCCA---GCCTCGGGG---AGGG-ATGTGGT-  
ACAG-AC-ACTCCCAACCCA---GGCTCGGGG---AGGG-ATGTGGT-  
AAAAGAC-ACTCCCAACCCA---TCCCTGGGT---AGGG-ACCTGGT-  
AAAAGAC-ACTCCCAACCCACCC---TCGGGGG---AGGG-ACGTGGT-  
AAAAGAC-ACTCCCAACCCAC---CCTAGGG---AGGG-ACGTGGT-  
AAAAGAC-ACTCCCAACCCCC---CCTGCGGGGCGAGGG-ACGTGGC-  
AAAAGAC-ACTCCCAACCCCC---CC-GCGGGGCGAGGG-ACGTGGC-  
AAAAGAC-ACTCCCAACCCGA---CCTCGGGG---AGGG-ACGTGGT-  
AAAAGAC-ACTCCCAACCCAC---CCGAGGGG---AGGG-ATGTGGT-  
AAAAGAC-ACTCCCAACCCAC---CCTCGGGG---AGGG-ACGTGGT-  
AAAAGAC-ACTCCCATCCCAA---CCTCGGGGGGGGG---ACGTGGT-  
AAAAGAC-ACTCCCAACCCAA---CCTCGGGGGGGG---ACGTGGT-  
AAAAGAC-ACTCCCATCCCAA---TCTCGGGGGGGG---ATGTGGT-  
AAAAGAC-ACTCCCATCCAG---TCGCGGGGGGGG---ATGTGGT-  
AAAAGAC-ACTCCCATCCAG---TCTCGGAGAGGG---ACGTGGT-  
AAAAGAC-ACTCCCATCCAC---CCTCGGGGAGGG---ACGTGGT-  
AAAAGAC-ACTCCCAACCCAC---CCTCGGGGGGG---AGTGN-  
AAAAGAC-ACTCCCATCCAN---CCTCGGGGGGG---ACGTGGT-  
AAAANAA-NCTCCCAACCCAN----CTGGGACGGG-----ACGTGGT-

Aameth6685  
Aappen6543  
Aappen6946  
Adist6684  
Aeuco5869  
Aristidaharvardii  
Arundinellaneपालensis  
Aschiren6581  
Bbladh5173  
Binsc5681  
Bradi4784  
Cdieter6642  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Dfili5874

GTTTCGCC-CCTCGTGG-CACGAGGGGCCGAGC--ACCACAAGTT-  
GTTTGGCC-TCCCGTGC-CGCAGGG-CGCG--TGGGCCG-AACTT-  
??  
-TTTACCCTCCTTGC-GGCAAGGA-GGGGCCGAACACAGTGT-  
GTATGGCC-CCCGTGC-CGCACGG-CGCGG-TGGGCCG-AAATT-  
GTATGGCC-CCCGTGC-CGCATGG-CGCGG-TGGGCCG-AAATT-  
-TT-TACCGCCCCCTGC-CGCACGGG-GG-GCCGAACACCAAGTC-  
-CTCACCT-CGCCCTGC-GGCA-GGG-GGAGCCAAACACAC-TG-  
GTTTGGCT-CCCGTGC-CGCAGGG-CGCAG-TGGGCCG-AAATT-  
-TTTCGCTCCCCCTTCC-GGCAAGGGCGAGGCTGAACACAACCTTC  
??  
GTTTGGCC-TCCCGTGC-CTCACGG-CGCGG-TGGGCCG-AAATT-  
-TT-GACCGCGCCCTGC-GGCAAGGG-GGGGCCAGACACACTG-  
GTCTGGCC-CCCGCCCCGGCAGGG-CGAAG-TGGGCCCAATAT-  
GTCTGGCC-CCCGTGC-CGCAAGG-CGGGG-TGGGCCCAAGTT-  
GTTTGGCT-CCCGTGC-CGCAGGG-CGCGG-TGGGCCG-AAATT-  
GTATGGCC-CCCGTGC-CGCACGGGGGGG-TGGGCC--AAATT-  
GTTTGGCC-CCCGTGC-CGCAGGG-CGCGA-TGGGCTGAAGTT-

Emuti5074  
Emuti6715  
Eragrostisdielsii  
Eviol5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdiss5145  
Sbicolour9319  
Hfilli5144  
Hhirt6656  
Hlongi6560  
Hpilg4603  
Hplig4738  
Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Panicumbisulcatum  
PentastichisaristifoliaSpies6  
Pfrank4705  
Sbic6588  
Sbic6589  
Shalep9318  
Tspic4846  
Ttria6555  
Zeadiplo  
ZmaysGfl  
Evillosa  
Saccharumofficinarum  
Imperatocylindrical  
Bothriochloainsculpta  
Dichanthiumannulatum  
Cybopogonprolixus  
Cplurnodus  
Cmarginatus  
Cexcavatus  
Hyparrhnietaмба  
Hanamesa  
Heteropogoncontortus

GTTTGGCC-CCTCGCGC-CTCAGG--CGCGG-TGGGCC--AAATTT  
GTTTGGCC-CCTC-CGC-CTCANGG-CACGG-GGGGCC--AAAATT  
ATTTGGCT-CCCCGTGC-AACACGA-CACGG-TGGGCC-AAGTT-  
-TCCTTCTCCCTGCC-AGGA-CGA-GGAACCAAACACCATGAT-  
GTTTGGCC-CCCCGTGC-CGCAAGG-CGGGG-TCGACCACAACCTT-  
NTT-----  
GTTTGGCC-TCCCGTGC-CGCAAGG-CGCGG-TAGGCCG-AAGTT-  
CTTTGCC-CCCCGTGC-CATACGA-CGCGGGTGAACACAACTT-  
TTTTGGCT-CCCCGTGC-CGCAAGG-CGCGG-TGGGCC-AAGTT-  
-GCCAACCCACCTTAT-TGAACGTT--GGG-CAAACACCCCTT-  
GTTTGGCC-CCCCGTGC-CGTATGG-CGCGG-TGGGCC-AACTT-  
ATTTGGCT-CCCCGT-CTCGCAAGGGCGGTG-GGCCTC-AAGAT-  
GTTTGGCC-CCCCGTGC-CGCAAGG-CGCGG-TGGGCC-AAGTT-  
??  
GTTTGGCC-CCCCGTGC-CGTACGG-CGCGG-TGGGCC-AAGTT-  
GTTTGGCC-CCCCGTGC-CATACGG-CGCGG-GGACACGAAATT-  
-CTCTACCACCCGTGC-CGCACGA-CGGGG-CATACACCAACAT-  
GTTTGGACCCCCCTGCCGGCAGGG-GGGCGGTGGGACACAAGTTG  
GTTTGGAC-CCCCGTGC-CGCAGGG-CGCGG-TGGGCC-AACTT-  
GTTTGGCT-CCCCCTTC-CGCAAGG-CGCGA-TGGGCC-AAAATT-  
GTTTGGCT-CCCCGTGC-CTCAGG-TGCGG-TGGGCC-AAGTT-  
??  
GTTTGGAT-CCCCGTGC-CGCATGGGCACAGTGGGCCGAAAGTT-  
TTTTGGCT-CCCCGTGC-CGCATGGGCATGG-CGCGG-TGGGCC-AAGTT-  
TTTTGGCT-CCCCGA-GCCGCATGG-CGCGG-TGGGCC-AAGTT-  
GTTTGGCT-CCCTGTGC-TGCAAG-CGCGG-GGGGCTG-AAATT-  
GTTTGGCC-CCCCGTGC-CGCAGGG-CGCGG-TGGGCC-AATTT-  
GTTTGGCT-CCCCGTGC-CGCAGGG-CGCGG-AGGGCCA-AAATT-  
GTCTGGCC-CCCCACGC-CGCAGGG-CGAGG-CGGGCC-AAGCA-  
GTCTGGCC-CCACGCGC-CGCAGGG-CGAGG-TGGGCC-AAGCA-  
GTTTGGCC-CCTCGTGC-CGCAAGG-CGCGG-TGGGCC-AAGTT-  
GTTTGGCT-CCCCGTGC-CGCAGGG-CGCGG-TGGGCC-AAGTT-  
GTTTGGAC-CCCCCGC-CGCAGGG-CGCGG-TGGGCC-AAGTT-  
GTTTGGCC-CCTCGTNC-CTCAAG-CACGG-TGGGCC-AAGTT-  
GTTTGGCCCCCCTNNTC-CCNAAG-GGGNGGGGGGCC--AAGTT-  
GTTTGGCCCCCCTTCC-CAAGGG--GAGG-TGGGCC-AAGTT-  
GTTGNNCCCCNNTGCC-CAAGGG--GAGG-TGGGCC-AAGTT-  
GTTTGGCCCCCCTTCTACAANAN--NACGG-TGGGCC-AAGTT-  
GTTTGGCCCCCCTGCC-GAATGGG-GAGG--TGGTACCNAAGTT-  
-TTTGGCCCCCTCCGTC-AAAGGGG-GGGG--GGCC--AAGNT-  
TTTTGGCCCCCCTTCC-CCATTGG-CGCGG-TGGGCCGGAATT-  
GTTTGNCCCCCGTNC--GCAGG-CGCGG-TAGGCCG-AAGTT-

Aameth6685  
Aappen6543  
Aappen6946  
Adist6684  
Aeuco5869  
Aristidaharvardii  
Arundinellaneapalensis  
Aschiren6581  
Bblad5173  
Binsc5681  
Bradi4784  
Cdieter6642  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Dfili5874  
Emuti5074  
Emuti6715  
Eragrostisdielsii  
Eviol5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdiss5145

CTAGCTGACGGGTGATACTCGT-GACTGT-CTTTGGACG-TGACTGGCC  
GGGGCTGCCG--CTAATCG-TGCCGG-CACA-TCACG-TGGTGGGCG  
??  
CCCTCTCCCGAG--GTAG-TCGTGTGCGAGTG-ACTTTGATGTTGAG-  
GGGGCTGCCG--T-AATCC-TGCCAGG-CACA-GCACC-TGGTGGGC-  
GGGGCTGCCG--CGTATCGG-TCGGG-CACA-GCACC-TGGTGGGCG  
GGGGCTGCCG--CAAACCGT-TCGGG-CACA-GCATG-TGGTGGGCG  
CC-TCCTGCAAG--GCTAATCTGGG--AGTG-TATTTAGCG-T-GACG  
CCCTC-CCCTTG--GGAG-GCTTGG--GAGTG-ACTTTGGCGGT-GACG  
GGGGCTGCCG--GAATCG-TGTCGAG-CACA-GCACC-TGGTGGGCG  
CGGTCTGCCGAGCGTAATCTTGCCGAGGTCACA-TTACG-TGGTTGACG  
??  
GGGGCTGCCG--CGAATCGT-TCGGG-CACA-GCACC-TGGTGGGCG  
CCCTT-GCCGAG--GGAG-TGTGCG-GAGTG-ACTTTTCGCGTG-GACG  
GGGGCTGCCG--TAATCT-CGCCAGG-CACA-TTACC-TGGTGGGCG  
GGGGCTGCCG--GGAATCGTCCCGGGGCA-TCACG-TGGTGGGCG  
GGGGCTGCCG--CGTACC--TGCCGGGCGCA-GACA-AGGTGGGCG  
GGGGCTGCCG--AAATCT-TGCCAGG-CACA-TCACG-TGGTGGGCG  
GGGGCTGCCATC--TGAATCATA-CCGGTACACA-TCACG-CGGTGGGCG  
GGGGCTGCCG--CATATCG-CGTCCGG-CACA-TCACG-TGGTGGGCG  
GGGGCTGCCG--CAGATCG-CGTCCGA-CACA-TCTT--TGGTGGGCG  
GGGGCTGCCG--CTTACGGTA-CCGGT-CACA-GCACA-AGGTGGGCG  
CCCTC--CCGG--GGAGTGC--TGGC-AGTG-TCTTTTCGCG-T-GACG  
CGGGCTCCCGAC--GGAATCGTTCCGAGTGACA-TTACG-TGGTGGGCG  
??  
GGGGCTGCCG--GAATCG-TGTCGGG-CACA-GCACC-TGGTGGGCG  
GGGGCTCCAGCA-TTATCTATCTTA-GTGAATTTTTT--CGGTATGAC

*Sbicolour9319*  
*Hfilli5144*  
*Hhirt6656*  
*Hlongi6560*  
*Hpilg4603*  
*Hplig4738*  
*Hquar5088*  
*Hrud6576*  
*Htamb6585*  
*Icylin5114*  
*Icylin6632*  
*Mcape6655*  
*Panicumbisulcatum*  
*PentaschistisaristifoliaSpies6*  
*Pfrank4705*  
*Sbic6588*  
*Sbic6589*  
*Shalep9318*  
*Tspic4846*  
*Ttria6555*  
*Zeadiplo*  
*ZmaysGfl*  
*Evillosa*  
*Saccharumofficinatum*  
*Imperatacylindrical*  
*Bothriochloainsculpta*  
*Dichanthiumannulatum*  
*Cytopogonprolixus*  
*Cplurnodus*  
*Cmarginatus*  
*Cexcavatus*  
*Hyparrhrietamba*  
*Hanamesa*  
*Heteropogoncontortus*

GGGGCTGCCGGC---GATTTCGT-GTCGGG-CACA-TCACG-TGGTGGGCG  
CCTTC-CCCAAG--GTACCCT-TGG--AATTT-ACCTTTG-TCATTAAC  
GGGGCTGTCG-A--CGAATCG-TGTCGGG-CACA-TCACG-TGGTGGGCG  
CCGGCTGCCGGT---GTATCGTGACCGGA-CACA-GCGCG-TGGTACGCG  
GGGGCTGCCGGC--GGAATCG-CGCCGGG-CACA-GCACG-TGGTGGGCG  
??  
GGGGCTGCCA-A--CGAATCG-TGTCAGG-CACA-GCACG-TGGTGGGCG  
GCCTCTCCCATGC-CTATCTGTGGCAGTGCACCTTTTAGCTAGTACGGCN  
CTCTCTGCCATG--CATGGCCGTGTCCAGGTGCACCTTTTAGCT-T-AACG  
GGGTCTGCCGAG---GGAGCCTTGGCGAGTGA--CTTTTGGTGGTTGACG  
GGGGCTGCCGGC---CGAA-CG-TGTCGGG-CACA-TGACG-TGGTGGGCG  
GGGGCTGCCGGC---GAATCG-TGTCAGA-CACA-TTATG-TGGTGGACC  
TGGGCTGCTGG---CATACCG-TGCCGGG-CACC-GCACG-TGGTGGGCG  
??  
GGGGCTTCCGGC--ATAACCG-TGTCGGG-CACA-GCATT-CTGTGGGC  
GGGGCTGCCGGC---GATTTCG-TGTCGGG-CACC-TCACG-TGGTGGGCG  
GGGGCTGCCGGC---GATTTCG-TGTCGGG-CACA-TCACG-TGGTGGGCG  
GGGGCTGCCGGCA-TATACAT-TGCCGGG-CACC-GCATG-TGGTGGGCG  
GGGGCTGCCAG---TGAATCACAGCCGGTACACA-TCACG-TGGTGGGCG  
GGGGCTGCCGGC---GAATCG-TGTCGG--CACA--CAC--TGGTGGGCG  
GGTGTCTGCCGGC---GAACCGC-GCCGGG-CGCA-GCACG-TGGTGGGCG  
GGGGCTGCCGGC---GAACCGC-GCCGGG-CGCA-GCACG-TGGTGGGCG  
GGGGCTGCCGGC---ATACCG-TGTCGGG-CACA-GCACG-TGGTGGGCG  
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GGGGCTGCCGGC---GAACCGC-GCCGGG-CACA-GCACG-TGGTGGGCG  
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GGGGCTGCCGGC---GANTCG-TGTTGGG-CACA-GCACA-TGGTGGGCG  
GGGGCTGCTGGC---GATTTCG--TTTCGG-TAGCAAAAGA-TGGTGGGCG  
GGGGCTGCCGGC---GAATCG-TGTTGGG-CACA-GCACG-TGGTGGGCG  
GGGGCTGCCGGC---GATTTCG--TTTCGG-TAACAGAACG-TGGTGGGCG  
GGGTCT-CTNCCGGCGAATCG-TTTCGGG-TACA-ACACA-TGGTGGGCG  
GGGGCTTCCGNC---GAATCG-TATCGGG-CACA-AAAAA-TGGTGGGCG  
GGGNTGCCGNC---GAATCG-TGTCGGG-CACA-GCACG-TGGTGGGCG

*Aameth6685*  
*Aappen6543*  
*Aappen6946*  
*Adist6684*  
*Aeuco5869*  
*Aristidaharvardii*  
*Arundinellanepalensis*  
*Aschiren6581*  
*Bbladh5173*  
*Binsc5681*  
*Bradi4784*  
*Cdieter6642*  
*Cleistachnesorghoides*  
*Cmarg4397*  
*Cplur6406*  
*Cprolix6682*  
*Danthoniacalifornica*  
*Dfili5872*  
*Dfili5874*  
*Emuti5074*  
*Emuti6715*  
*Eragrostisdielsii*  
*Evill5085*  
*Hanam4807*  
*Hanam4841*  
*Hcon4741*  
*Hdiss5145*  
*Sbicolour9319*  
*Hfilli5144*  
*Hhirt6656*  
*Hlongi6560*  
*Hpilg4603*  
*Hplig4738*  
*Hquar5088*  
*Hrud6576*

AGGCATCA--CGTTGTT---CTCG--ACTGACAAAGCCCCGGGGAC---G  
--ACATC---AGTTGTT---CTC---GGTG-CAT-GACCTCGGCAC---G  
??  
CCA-AGG--CAG-ACGTCTCTC---TGACACACGAACGCC--TCA---GG  
G-ACATC---AGTTGT---CTC---CGTG-CAG-TGACCCGTCAA---A  
--ACCT---AGTTGTC---ATC---GGTG-CCG-TGACCCGACGC---G  
--ACAAC---AGTTGTT---CTC---GATG-CAG-TGGCTCGACAC---G  
CC-ACG---TAAACTTTCTCTC---AA---CAAAAACCTC--TCG---TG  
CCA-ATG---AAGATGTTCTCTC---TG-CACAAAAGGGCG--TCAT--GG  
C-ACATC---GTTGTT---CTC---GGTG-CAG-CGCCCGGCAT---G  
CCACATC---AGATGTT---CTC---TCTGACATACA-CTCCTCAC---G  
??  
--ACACCT---AGTTGTT---CTC---GGTG-CAG-CGCCCGGCAC---G  
CCA-AGG--CAGTACGTCTCTT---TGACACATGAAGGCGTCCAC--GG  
C-ACATC---AATTGTT---CTCT--CGTA-CCA-TACCCNTNCG--CG  
--ACATC---AGATGTT---CTC---TGTG-CAATGCCCCGTCAC---G  
--ACCA---ACGTTGT---CAC---GGTG-CAG-CGCCCGGCAC---G  
--ACATC---AATTGT---CTC---GGTG-CAA-TACCCTCAAAA---G  
--ACATC---AATT-T---CTC---GGTG-CAA-TGCCCGGAAA---G  
--ACATC---AAGTTGTT---CTC---GGTG-CAG-CGCTCGGCAC---G  
C-ACATC---AAGTTGTT---CTC---TGTG-CAG-CGACTCGGCAC---G  
--ACAC---ACGGTGT---CTC---GCTG-CTG-CGAACTGGTCCA-AG  
CCACAG---CAG-ACCTTT-CTT---AAT---CAAAAACCC--TCG---GC  
C-ACATC---AGATGTT---CTC---TGTGACAC-TGACCCCTCAC---GG  
??  
G-ACATT---AGTTGTT---CTC---GGTG-CAG-CGCCCGGCAC---G  
C-ACAACA---TTTGTTCCTCGSCGATAAAA-ATCCCTCCCCTCGCT---  
--ACATC---TAATTGTT---CTC---TGTG-CAA-CACCC--ACAC---G  
CCA-AGA---AAA-ACTTCCCTC---AA---CAAAAAGCCT-TCG---GG  
--ACATT---AGTTGTT---CTC---GGTG-CAT-TGCCTCGGCGC---G  
--ATCT---CGCTATA---CTA---AACG-CAA-TGCATCGGGG---C  
--ACATCA---G-TTGT---CTC---GGTG-CAG-TGCCCGGCAC---G  
??  
C-ACATT---AGTTGTT---CTC---GGTG-CAT-TGCTTCGGAGC---C  
C-ACGTCA---AATTCTCNTCCAGAGACTCCATCATGGCTGCCAGT-TA

Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Panicumbisulcatum  
PentaschistisaristifoliaSpies6  
Pfrank4705  
Sbic6588  
Sbic6589  
Shalep9318  
Tspic4846  
Ttria6555  
Zeadiplo  
ZmaysGf1  
Eviollosa  
Saccharumofficinatum  
Imperatacylindrical  
Bothriochloainsculpta  
Dichanthiumannulatum  
Cybopogonprolixus  
Cplurnodus  
Cmarginatus  
Cexcavatus  
Hyparrhnietaamba  
Hanamesa  
Heteropogoncontortus

CCACAGT--CAA-AC-TTCTCTC---GG---CGACATTGCC-TCT---GC  
CCACATCA--AGATGTT---CTC---TGTGACAGACGACGCGCTCGCGG  
--ACATC---AGTTTTT---CTC---GGTG-CA---CACCCGGCGC---G  
--ACACTC---ATTGTT---CTC---GTG-CAG-CGCCCCGGCAGC---G  
--ACTTAC--AGTTGTT---CTC---GGTG-CAG-CGTCCCGGTAC---G  
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G-AGATCT--ACTTGT---GCC---GGTG-CAG-CGCCCCGGCAGC---G  
--ACAC-T-TAAATGTT---CTC---TGTG-CAG-CGCCCC-ACAC---G  
--ACAC-T-TAATTGTT---CTC---TGTG-CAG-CGCCCCGACAC---G  
--ACATCG-TAGGTGTT---CTC---TGTG-CAA-TATCGCCACAT---G  
--ACATC---AGTTGT---CTC---GGTGACAC-TGCCCCGGCAGC---G  
--ACATCT--AATTGTT---CTC---GGTG-CA---CGCCCCGGCAGC---G  
--ACATC--AAGTTGTT---CTC---GGTG-CAG-CGTCCCGGGCGC---G  
--ACATC--AAGTTGTTGTTCTC---GGTG-CAG-CGTCCCGGGCGC---G  
--ACATC---AGTTGTT---CTC---GGTG-CAG-TGCCCCGGCACA---G  
--ACACTC--AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCAGC---G  
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--ACATC---AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCAGC---G  
--ACATC---AGTTGTT---CTC---GGTG-CAG-CACCCCGGCACAC-G  
--CAACTG-TGTTTTT-----GGTG-CAG-CACCCCGGCAGC-GCG  
--ACAAN---AGTTGTTNTT-----GGTN-NNT-TACTCCGGCGG-GCG  
--ACATC---AGTTGTTTTT-----GGTG-CAG-TACCCCGGCACA-CG  
--ACA---AAAATTTTTTTC-----GGTT-CAN-TTCCCGGCAGC-GCG  
--AAATC---AATTTTT---CTC---GGTN-CAG-NGCCCGGCAGC-GCG  
--ACATC---AATTGTTTTC-----GGTT-CAA-TTCCCGGCAGC-GCG  
--ACATT---AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCAGC-NC-

Aameth6685  
Aappen6543  
Aappen6946  
Adist6684  
Aeuco5869  
Aristidaharvardii  
Arundinellaneapalensis  
Aschiren6581  
Bbladh5173  
Binsc5681  
Bradi4784  
Cexc6577  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprollix6682  
Danthoniacalifornica  
Dfili5872  
Dfili5874  
Emuti5074  
Emuti6715  
Eragrostisdielsii  
Evill5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdiss5145  
Sbicolour9319  
Hfilli5144  
Hhirt6656  
Hlongi6560  
Hpilg4603  
Hplig4738  
Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Panicumbisulcatum  
PentaschistisaristifoliaSpies6  
Pfrank4705  
Sbic6588

CAACTTGGCGTTATCGAAACCTAATGACTCATCGAATTA-----T-GCAC  
CG-CTAT-CGT-ATTGGCCCTTACG-ACCCATTGA-GCACAGGTAAGC-C  
??  
CG-CAACTTGCAT-TCAA-ATCTCGATCCATCGAGCG---ATTCTGCGC  
AGGCCG--GCGCATCGCCCTAAGG-ACCCATCGA-GCACC--GGAGCGC  
CAGCCGG--CGAGAGGGCCCTGTACGACCCATCG-----ACC--GCAGCGC  
CGGCCAG-CGTCTTGGCCC--TAAGGACCCATGGA-GCACC--GCAGCGC  
CG-CAACTTGCAT-TCAANN-----G-----G-----G-----  
CG-CAACTTGTATCTGAAA-CTATGAATCCATCGAGCA---ATTATGCAC  
CGGCCGG-CGT-ATCGCCCTGAGG-ACCCATCAA-GCACC--TAAGCGC  
CGCCTAT--TGATTTCAACCTTACG-ATCCATTACGAAACT---CAGCAC  
CG-CAACTTGCAT-TCAAC-CTCTCGATCGATCGAGCAC--ATTGAGCAC  
CGGCCGG--CGCATCGCCCTAA-GGACCCATGGA-GCACC--GCAGCGC  
CGGCAACTTGGGANTGAAA-----G-----G-----G-----  
CGGCCCT---CGCTTGACCTAAAG-ATCCATCGA-GCAANT--TAAGCCC  
C-GCCGG--TCCGACTGCCCTAACG-ACCCATCGA-GCAAT--TAAGCAC  
CAGCCGG--CGAGACGGCCCTCA-GGACCCCTCGC-ATACC--GCAGCGC  
CGCGC-----ATCTGCCCTAAGG-ACCCATCGA-GCACC--TGAGCGC  
CGNCCAGGTTGCGTANGCCC-----G-----G-----G-----  
CAACCGG--CTTATCGGCCCTAAGG-ACCCATTNT-GCACA---CCANNC  
CAACCGG--CTTATCTGCCCTAACG-ACCCATTGT-GCACA---TCAGCGC  
TATCAGGTACGGAATGGCCCTCA-TGACCCATGA---ACC--GAAGCGC  
AC-CTACTTACGT-TCNNA-AACTCGATGGTTCGCGGN---ATTGAGCAA  
CAGCCGG-TTGGGACTGCCNTAACG-ATCCATCA--GCAAT--TAAGCAA  
??  
CGGACGG-CAA-ATCGCCCTTAGG-ACCCATCGA-GCACC--GCAGCGC  
??  
TGGCCG--GTGCATCTGCCCTAAAG-ACCCTTCNG-GCACT--GCAGCCC  
CC-CAGTTT-----G-----G-----G-----G-----  
TGGTCCG-TGT-GTCTGCCCTAAGG-ACCCATCGA-GCACC--TAAGCGC  
GTATCCG-ACGCAATGCCCTCAATG-GACCCT--A-TTACC--G-AGCGC  
CGGCCGG-CGC-GACGCCCTAAGG-ACCCATCGA-GCACC--GAAGCGC  
??  
CGCTCCG--GT-ATTNGCCCTAAG--ACCCATCGA-GCACC--TAAGCGC  
ANTTCNAANNTAATGGACCCATCNAGCACATTAAGCGCATCACC-----  
TG-CAGCTTGCTTTTCTGGCAATCGATGGATCTAGAGC--ATTATGCAC  
CGGCCGGTTGGCTTTCGCCCTATCGGATCCATCGA-GCAAT--TCAGCGC  
C-GCCAG--TGCGTTGGCC--TAA-----G-----G-----G-----  
CGGCNGT--TGATCTGCCCTAAGG-ATCCATGGAAGCACN--TCTTCAC  
TAGCTAG--CTTGTGGCCCTTAAGGACCCAGGA---ACC--GTAGCGC  
??  
CTGCCGA-AGCGATTGACCCTAAGGGACCCATCTA-CCACC--GCAGCGC  
TGGCCG--GTGCATCTGCCCTAAAG-ACCCTTCGG-GCA-T--GC-GCCC

*Sbic6589* TGGCCG--GTGCATCTGCCCTAACG-ACCCTTCGG-GCACT--GC--GCA  
*Shalep9318* TACCTA--TCTTATTGGCCCTAAG--ACCCATATG-TGAAC--TAAGCGT  
*Tspic4846* CGGCCGCG--GGGTAGGCCCTAAGG-ACCCATCGA-GCACC--GAAGCGC  
*Ttria6555* CGGCCGCGCT-ATNTGCCCTTATG--ANCCATCTA-GCACC--GAAGCGC  
*Zeadiplo* TGGCCG--CCATTTCGGCCCTACGG-ACCCATCGAGCGACC--GA-GC--  
*ZmaysGfl* CGGCCG--CCATTTCGGCCCTAAGG-ACCCATCGAGCGACC--GA-GC--  
*Evillosa* CATACGGCGAC-ATCGG-----  
*Saccharumofficinatum* CGGCCGCGC-C-ATCGG-----  
*Imperatacylindrical* CGGCCGCGG-TG-TCGGCCCTAAGG-ACCCATCGA-GCACC--GCAGCGC  
*Bothriochloainsculpta* CGGCCGCGC--CATCGGCCCTAAGG-ACCCATCGA-GCACC--GAAGCGC  
*Dichanthiummannulatum* --ACCGCGC--CATCGGCCCTAAGG-ACCCATCGA-GCACC--GAAGC--  
*Cytopogonprolixus* --ACAGCGC--TATTGTCGCTAAGG-AGGCCATGGAGCGAAC-GAAGCGC  
*Cplurnodus* -AACCAGC--ACCTGTCGTCAGG-AGGGATCGA-GCAGC--GAAGGGC  
*Cmarginatus* --ACCGCGC--CATTGTCGCTAAGG-AGGCCATCGAGCACC--GAAGCGC  
*Cexcavatus* --GCCGCGC--TATTGTCCTAAGG-AGGCCATCGAGCACC--GAAGCGC  
*Hyparrhrietamba* --GCCGCGC--CATCGGCCCTA-GG-ACCCATCGA-GCACC--GAAGCGC  
*Hanamesa* --GCCGCGC--TATTGCCCCTAAGG-ACCCATCGA-GCACC--GAAGCGC  
*Heteropogoncontortus* -AGACGGCA--AATTGGCCCTTAGG-ACCCATCGA-GCACC--GNAGCGC

*Aameth6685* TTCATC-CCTC-GGA-TCGCAAT-----  
*Aappen6543* ATCGCC-GCTC-GGA-TCGCGACCCCAT--GTCATTCTCC-ACTACCCGC  
*Aappen6946* -----  
*Adist6684* TTCACACC-TC-GTA-TCGCAATTCGCT--GCCATCTTCAT-CTATCCGA  
*Aeuco5869* ATCCCC-GCTC-GGA-CCGCGACCCCG--GTCAGTCGGG-ACTACCCGC  
*Aristidaharvardii* A-AGCC-GCTC-GGA-CC-----  
*Arundinellaneapalensis* ATCGCC-GCTC-GGA-CC-----  
*Aschiren6581* ???  
*Bbladh5173* TTCNCCCCCTC-GGA-TCGCAATTCAT--GACATTTTGA-CTATCCGA  
*Binsc5681* CTCGCC-GCTC-GGA-CCGCGACCCCG--GTCAGTCGG-ACTACCCGC  
*Bradi4784* ATCACC-CCTCGGAT--CGCAACCCCT--NCCNTNTTCG-ACTACCCGA  
*Cdieter6642* ???  
*Cleistachnesorghoides* ACCGCC-GCTC-GGA-CCGCGACCCCG--GTCAGTCGGG-ACT-----  
*Cmarg4397* ???  
*Cplur6406* ATCNCC-NCTC-GNA-TCNCCACCNCAT--GTCAATCTGG-ACTACC-GN  
*Cprolix6682* ATCGCC-CCTC-GGA-CCGCAACCCCAT--GCCATTCTGGTACTACCCGC  
*Danthoniacalifornica* T-CGTC-GCTC-GG-----  
*Dfili5872* ATCACC-CCTC-GGA-CCGC-ACCCCA--GTCATTCTGG-ACTACCCGC  
*Dfili5874* ???  
*Emuti5074* ATTGCC-NCTC-GGA-CCNCCACCCAG--GNNAGCCC-----  
*Emuti6715* ATTTCC-CCTC-GGA-TCGCGATCCCT--GTGATTTTGG-ACTATACGC  
*Eragrostisdielsii* A-TCTC-GCTC-GGA-CC-----  
*Evill5085* TTCACTGCAGG-GGA-TCGCATTTCCATANCCTTCTTACTACTATNCCA  
*Hanam4807* NATCNCCCTC-GGA--CGCAANCCCT--GNCATNTTGN-ACTANNCGN  
*Hanam4841* ???  
*Hcon4741* ATCGCC-GCTC-GGA-CCGCGACCCCG--GTCAGTCGGG-ACTACCCGC  
*Hdiss5145* ???  
*Sbicour9319* CATCCATGCTC-GGA-CCGCGACCCAN--GTCAGTCNGG-ACCACCCGC  
*Hfilli5144* ???  
*Hhirt6656* ATCCC--GCTC-GGA-CCACGACCCCG--GTCAGTCGG-ACTACCCGC  
*Hlongi6560* ACA---CCTT-CTA-CCGC-ACCCAT--GTCA-GCTGG-ACTACCCGC  
*Hpilg4603* ATCGCC-GCTC-GGA-CCGCGACCCCG--GTCAGTCGGG-ACTACCCGC  
*Hplig4738* ???  
*Hquar5088* ATCCC--GCTC-GGA-CCGCGACCCCG--GTCATTCTGG-ACTACCCGC  
*Hrud6576* ???  
*Htamb6585* TTCACCCTCG-GTA-TCGCATCTCCT--GTCATCTTCTG-CTATCCGC  
*Icylin5114* ATCGCC-GCTC-GGA-TCACGATCCCAT--GTCATCTTGG-ACTATCCGC  
*Icylin6632* ???  
*Mcape6655* ATCACC-GCTC-GGATCCGA---CCCAT--GTCATTCTGG-ACTACCCGC  
*Panicumbisulcatum* TTCGTC-GCTC-GGA-CC-----  
*PentaschistisaristifoliaSpies6* ???  
*Pfrank4705* ATCGCC-GCTC-GGA-CCGCGACCCCG--GTCAGTCGGG-ACTACCCGC  
*Sbic6588* ANTCCATGCTC-GGA-CCGCGACCCAN--GTCANNNGG-ACCACCCGC  
*Sbic6589* CATCGATGCTC-GGA-CCGCGACCCCG--GTCAGTCGG-ACCACCCGC  
*Shalep9318* TTCCTC-GCTC-GGA-CCGC-ACCCAT--GTCATACTGG-ACTACCCGC  
*Tspic4846* ATCGCC-GCTC-GGA-CCGCGACCCCG--GTCAGTCGGG-ACTACCCGC  
*Ttria6555* TTTTCC-GCTTTTGA-ACGCGACCC-AN--GTCA-TCTGG-ACTACCCGC  
*Zeadiplo* TTGTCC---TC-GGA-CCGCGA-----  
*ZmaysGfl* TTGCC---TC-GGA-CCGCGA-----  
*Evillosa* -----  
*Saccharumofficinatum* -----

<i>Imperatacylindrical</i>	ATCGCC-GCTC-GGA-CC-----
<i>Bothriochloainsculpta</i>	TTACGCCGCTC-GGA-CCGCGA-----
<i>Dichanthiumannulatum</i>	ATCACC-GCTC-GGA-CCGCGA-----
<i>Cybopogonprolixus</i>	ATCCCC-GCTC-GGA-CCGCGA-----
<i>Cplurnodus</i>	ATCCCC-GCTC-GGA-CCGCGA-----
<i>Cmarginatus</i>	ATCCCC-CCTC-GGA-CCGCGA-----
<i>Cexcavatus</i>	ATCGCC-GCTC-GGA-CCGCGA-----
<i>Hyparrhrietamba</i>	ATCNCC-NCTC-GGA-CCGCGA-----
<i>Hanamesa</i>	ATCGCC-GCTC-GGA-CCGCGA-----
<i>Heteropogoncontortus</i>	ACAGCC-GCTC-GGA-CCGCGA-----

**Appendix H:** CLUSTAL W (1.74) multiple sequence alignment data of the Andropogoneae specimens for trnL-F + ITS region.

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Aadscen6713      ?????????????????????????????????????????????????????????????
Aameth6679      TCGTGACCCCGGTGAAAATATG-AC-AACAAAGGCTTCCCTCAGTT--G
Aameth6685      TCGTGACC-GCGTCGATGACCG--T-AACAAAGT-TTCCCTCAGTT--G
Aappen6543      TCGTGAC-----GAAGGAATTC-GT-AACAAA-GTTTCCCTCAGTT--C
Aappen6946      TCGTGACC-----GAGGAATTC-GT-AACAAA-GTTTCCCTCAGTT--C
Adist6684       TCGTGACA---CGAAAGAAA-C-GT-AACAAACGCTTCCCTCAGTT--C
Aeuco5869       TCGTGACCCA--GAAA-AATCC-GT-AACAAG-GCTTCCCTCATGCATG
Arundinellane  
palensis      TCGTGACCC-TTAAACAAAACAGACCGCGAACATGTTTCCCATGT---
Aschiren6581    ?????????????????????????????????????????????????????????????
Bbladh5173      TCGTGACC--CTGAAGGAAATCAG--ACCGAAGGCTTCCCTCAGTT--G
Binsc5681       TCGTGAGC----GGAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT---
BBrachiaria     ?????????????????????????????????????????????????????????????
Bradi4784       TCGTGAACCT--GAAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G
Bserra6554      ?????????????????????????????????????????????????????????????
Cexc6577        ?????????????????????????????????????????????????????????????
Cdieter6642     TCGTGACCCCG--ACGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G
Cleistachnesor  
ghoides      TCGTGACC--CTTAAACAAAACAGACCGCGAACGCGTCTCTCGTG---
Cmarg4397       TCGTGACAATTCGAAGGAATTCG-GT-AACAAGAGCTTCCCTCAGTT--
Cplur6406       TCGTGACC--ATGTAGATATTC-GT-AACAAA-GCTTCCCTCAGTT--C
Cprolix6682     TCGTGACCCGG---TTAAACCG-GT-AACAAA-GCTTCCCTCCCT--G
Danthoniacalif  
ornica      TCGTGACCC--GAAACAAAACCGACCGGAGAACGCGTCCACCTGT---
Dfili5872       TCGTGACCGGA--GGA---ATTG-GT-AACAAG--CTTCCCTCAGTT--G
Emuti5074       TCGTGAAC----GGAGGAATTC-GT-AACAAA-GTTTCCCTCAGT---G
Emuti6715       TCGTGAGC-----GAGGAATTC-GT-AACAAA-GCTTCCCTCAGT---G
Eragrostisdiel  
sii         TCGTGACC---CTGACCAAAATAGACCGGAACATGTCTCATCCATG---C
Evill5085       TCGTGACC--CTGAAACAAAATC-GT-AACAAA-GCTTCCCTCTTTG--T
Hanam4807       TCGTGAGC---CGGAGGAATTC-GT-AACAAG-TTCCCTCTCAT--G
Hanam4841       TCGTGACC---CGAGGAAATC-GT-AACAAG-CTTCCCTCTCAT--G
Hcon4741        TCGTGACC--CTGAAAGAAATC-GA-AACAAA-GCTTCCCTCTGTT--G
Hdiss5145       TCGGTACT---TGGAGGAATTC-GT-AACAAG-CTTCCCTCAGTT--G
Sbicolour9319  TCGTGACCC--TTAAACAAAACAGACCGTGAACATGTCTCTCATGT---
Hfilli5144     TCGTGATAG--CGGAGGAATTC-GT-AACAAG-CTTCCCTCTCAT--G
Hhirt6656      ?????????????????????????????????????????????????????????????
Hlongi6560     ?????????????????????????????????????????????????????????????
Hpilg4603      TCGTGAGCGG---AGGAATTC-GT-AACAAA-GTTTCCCTCCGTT--G
Hpilg4738      TCCTGACC--CGGAAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--C
Hquar5088      TCGTACCC---TGGAAGAAAACAGA-AACAAA-GCGTCCCTCTCAT--G
Hrud6576       TCGTGACCC---TTAAACAAAACAGACCGCGAACGCGTCTCTCATGT---
Htamb6585      TCGTGACC---CGGAGGAAATCCGT-AACAAG-CTTCCCTCTGAT--G
Icylin5114     TCGTGACCT---GAAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G
Icylin6632     TCGTGACC--G-GAAGA-ATTC-GT-AACAAA-GCTTCCCTCAGTT--C
Mcape6655      TCGTGAGG---CGAAGGATT-C-GT-AACAAG-GTTTCCCTCAGTT--G
Monocymbium    ?????????????????????????????????????????????????????????????
Pdilat6617     ?????????????????????????????????????????????????????????????
Pentastichtisar  
istifoliaSpies6 TCGTGACC---GAAACCAATGCAGACCGTGAACAAGTCACATTTGT---
Pfrank4705     TCGTGACC-----GAGGAATTC-GT-AACAAA-GTTTCCCTCAGTT--G
Pschinz6652    ?????????????????????????????????????????????????????????????
Sbic6588       TCGTGACC---TTAAACAAAACAGACCGTGAACATGTCTCTCATGT---
Sbic6589       TCGTGACC--CAGGAGGAATTC-GT-AACACA-GTTTCCCTCAGTT--C
Shalep9318     TCGTGACA---CGAAGGAAATC-GT-AACAAG-CTTCCCTACGTT--C
Snigros6567    ?????????????????????????????????????????????????????????????
Tleuc6627     ?????????????????????????????????????????????????????????????
Tspic4846     TCGTGAT-----GGAGGAATTC-GT-AACAAA-GCTTCCCTCAGTT--G
Ttria6555     TCGTGACC-GT-GAAGTAACG-CC-AACAAGAGTTTCCCTCAGTT--G
Zeadiplo       CTGTGACCC---TTAAACGAAACAGACCGCGAACGAGTACCCCGTG---C
ZmaysGfl       CCGTGACCC---TTAAACAAAACAGACCGCGAACGAGTACCCCGTG---C
Saccharumoffic  
inarum      TCGTGACC---TTAAACAAAACAGACCGCGAACGAGTCTCTCGTG--CC
Imperatacylind  
rica       TCGTGACCC---TTAAACAAAACAGACCGCGAACGAGTCTCTCGTG--CC
Dichanthiumann  
ulatum     TCGTGACCC---TTAAACAAAACAGACCGCGAACGAGTCTCTCGTG--CC

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Aadscen6713      ?????????????????????????????????????????????????????????????
Aameth6679      TCCC-GGGCTTC-----GGCTCGGC---CAAGGCACCCGAG
Aameth6685      CCCCCGGGCTTC-----GGCTCGGC---CAAGGCACCCAT
Aappen6543      CNCCNCGGTAC-----GGTTCGGC-CTCTGGCTCCCNAG
Aappen6946      ACCCGGGGCTAG-----GGCCCCGACTCCGGCCCCCTAG

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*Adist6684* CCCCCGGCCCTT-----GGTTCAC-CTTGGGCCCCCTT  
*Aeuco5869* CCACCGGGCTCC-----GGCCCGGC--AAAGGCTCCCGAG  
*Arundinellaneapalensis* -CGCCGGACTTC-----GGTCCGGT--TTAGGCCCCCGAG  
*Aschiren6581* ???  
*Bbladh5173* CGGCCGGGCTCC-----GGCCCGGC--CTAGGCCCCCGAG  
*Binsc5681* CCACCGGGCCTC-----GGCC-GGC--ACAGGCCCCCGAG  
*BBrachiaria* ???  
*Bradi4784* CGGCCGGGCTCC-----GGCCCGGC--AAAGGCTCCCGAG  
*Bserra6554* ???  
*Cexc6577* ???  
*Cdieter6642* CCCCCGGGCTTC-----GGTCCGGC--CAAGGCCCTCTAG  
*Cleistachnesorghoides* CGCCCGGGCCTC-----GGCNCGGC--ANNNGNCCCGAG  
*Cmarg4397* CGGCCGGGCTTC-----GGCCCGGC--CAAGGCCCCCGAG  
*Cplur6406* CTCCNCGGCTAG-----GG--CCGG--CTTGGCCCCCTAT  
*Cprolix6682* CGGCCGGGCTTC-----GGTCCGGC--CAAGGCCCCCGAG  
*Danthoniacalifornica* CGGCCCGGCTCC-----GGCCCGGC--AAAGGCTCCCGAG  
*Dfili5872* CCACCGGGCCTC-----GGCCCGGA--CAAGGCCCCCGAG  
*Emuti5074* CCACCGGGCCTC-----GGCCCGGA--CAAGGCCCCCGAG  
*Emuti6715* CCACCGGGCCTC-----GGCCCGGA--CAAGGCCCCCGAG  
*Eragrostisdielsii* CGTCCG--GTGATGGGGCTTG--CCTCGCTCCCGAC--TTAGGNCCCGAG  
*Eviol5085* CTGCCGGGCTCC-----GGTCCGGC--CTGAGGTCCCGAG  
*Hanam4807* TCGCCGGGCTTC-----GGCCCGGC--CAAGGTTCCCGAG  
*Hanam4841* TCGTGGGCTTC-----GGCCCGGC--CAAGGTTCCCGAG  
*Hcon4741* TCGTCCGAGCCTC-----GGTCCGGC--AAGGCCCCCGAG  
*Hdiss5145* TCGCCGGGCTTC-----GGCCCGGC--CAAGGTTCCCGAG  
*Sbicolour9319* -CGTCCGAGCTTT-----GGTCCGGC--CAAGGTTCCCGAG  
*Hfilli5144* TCGCCGGGCTTC-----GGCCCGGC--CAAGGTTCCCGAG  
*Hhirt6656* ACCCTGGGCTTC-----GGCCCGGC--CAAGGTTCCCGAG  
*Hlongi6560* ???  
*Hpilg4603* CGGCCGGGCTTC-----GGCCCGGC--CAAGGCCCCCGAG  
*Hpilg4738* CCCCCGGGCAAG-----GGCCCGGC--ACAAGGCCCCCGAG  
*Hquar5088* TCGCCGGGCTTC-----GGCCCGGC--CAAGGTTCCCGAG  
*Hrud6576* -CGCCGGGCTTC-----GGCCCGGC--CAAGGTTCCCGAG  
*Htamb6585* TCGCCGGGCTTC-----GGCCCGGC--CAAGGTTCCCGAG  
*Icylin5114* CCGCCGGGCTTC-----GGCCCGGC--CAAGGCCCCCGAG  
*Icylin6632* CCCCCGGGCTTC-----GGTCCGGC--CCCAGGCCCTAT  
*Mcape6655* CCCCCGGGCTAC-----GGTCCGGC--ACAGG--CCCCTAG  
*Monocymbium* ???  
*Pdilatum6617* ???  
*PentastichisaristifoliaSpies6* CGGCCACACGA-----GCTCACGCCGTGTGGC--CTAGGCCCCCGAG  
*Pfrank4705* CCGCCTGGCTTC-----GGCCTGGT--CAAGGCCCTCGAG  
*Pschinz6652* ???  
*Sbic6588* -CGTCCGAGCTTC-----GGTCCGGC--CAAGGTTCCCGAG  
*Sbic6589* ACCNCGGGCTGG-----GGTCCGGC--CTAGGCCCCCGAG  
*Shalep9318* CTCTGGGCTTC-----TGCCCGTG--CAAGGCCCTCGAG  
*Snigros6567* ???  
*Tleuc6627* ???  
*Tspic4846* CCACCGGGCTTC-----GGCCCGGC--AAAGGTTCCCGAG  
*Ttria6555* CCCCCGGGCTTC-----GGCCCGGC--CACGGCCCCCGAG  
*Zeadiplo* CGCCGG--GCTCC-----GGCCCGGC--CAAGGCCCCCGAG  
*ZmayGfl* CGCCGG--GCTCC-----GGCCCGGC--CAAGGCCCCCGAG  
*Saccharumofficinatum* -GCCGG--GCTTC-----GGCCCGGC--CAAGGCCCCCGAG  
*Imperatacylindrica* -GCCGG--GCTCC-----GGCCCGGC--CAAGGCCCCCGAG  
*Dichanthiumannulatum* -GCCGG--GCTTC-----GGTCCGGC--CAAGGCTCCCGAG

*Aadscen6713* ???  
*Aameth6679* -CCCCGTCCC-----GGGGCGGAGGTGCCGC--AAAAGAACCACGGC  
*Aameth6685* -CCTCGTCCC-----GGGGCGGAGGTGCCGC--AAAAGAACCACGGC  
*Aappen6543* -CCCCGTCCA-----GGGGCGGAAGGGCCGC--AAAACAACCACGGC  
*Aappen6946* GCCCCGTCCC-----GGGGCGGAGGGGCCGC--AAAACAACCACGGT  
*Adist6684* -TCCCCTCCC-----TGGGTAATCGCCCGAA--AA-ACAACCACGGC  
*Aeuco5869* -CCCCGTCCC-----GGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Arundinellaneapalensis* -CTCCGTCCC-----GGGGCAGAGGGCCCT--AACAGAACCACGGC  
*Aschiren6581* ???  
*Bbladh5173* -CCCCTTCCC-----GGGGCGGAGGAGCCGC--CAAAGAACCACGGC  
*Binsc5681* -CTCCGTCTC-----GGGGCGGAGGGGCCGC--AAAAGAACCACCGC  
*BBrachiaria* ???  
*Bradi4784* -CCCCGTCCA-----GGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Bserra6554* ???  
*Cexc6577* ???



*Cdieter6642* -CCCTGTCAC-----GGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Cleistachnesorghoides* -CCNGTCTNT-----GGGGCGGAGGGGCCAC--AAAAGAACCACGGC  
*Cmarg4397* -CCCTGTCAC-----GGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Cplur6406* -CCCCNTCTT-----GGGGTGGAGGGGCCGC--AAAATAACCACGGC  
*Cprolix6682* -CCCTGTCAC-----GGGGCGGAGGGGCCGC--AAAACAACCACGGC  
*Danthoniacalifornica* -CTCCCTCTCC-----GGGAGGGGAGCGGCCGT--AAAAGAACCACGGC  
*Dfili5872* -CCCCGTCCC-----GGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Emuti5074* -CTCCGCA-----GAGGGGCCGC--AACAGAACCACGGC  
*Emuti6715* -CTCCGCA-----GAGGGGCCGC--AACAGAACCACGGC  
*Eragrostisdielsii* -TTTCTTAC-----GGAGGGCTGGGGGCCGC--AACAGAATCCACGGC  
*Eviol5085* -CCCCGTCT-----GGGGCGGAGGGGCCGC--AAAAGATCCACGGC  
*Hanam4807* -CCCCGTCCC-----GGGGCGGAGGTGCCGC--AAAAGAACCACGGC  
*Hanam4841* -CCCCGTCCG-----GGGGCGGAGGTGCCGC--AAAAGAACCACGGC  
*Hcon4741* -CTTC-----GGCAGAGGGGCCGC--CACAGAACCACGGC  
*Hdiss5145* -CCCCGTCTCTCATGTTCGGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Sbicolour9319* -CTCCAAC-CT-----GGGGCAGAGGGGCCAC--AAAAGAACCACGGC  
*Hfilli5144* -CCCCGTCCC-----GGGGCGGAGGTGCCGC--AAAAGAACCACGGC  
*Hhirt6656* -CCCCGTCCG-----GGGGCGGAGGTGCCGC--AAAAGAACCACGGC  
*Hlongi6560* ???  
*Hpilg4603* -CCCTGTCAC-----GGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Hpilg4738* -CCCCGTAC-----GGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Hquar5088* CCCCCGTCCC-----GGGGCGGAGGTGCAGCC--AAAAGAACCACAGC  
*Hrud6576* -CCCCGTCTCTCATGTTCGGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Htamb6585* -CCCCGTCTCTCATGTTCGGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Icylin5114* -CTCCGTCCC-----GGGGCGGAGGGGCCGC--AACAGAACCACGGC  
*Icylin6632* -CTCCGTCCC-----GGGGCGGAGGGGCCGC--AACAGAACCACGGC  
*Mcape6655* -CCCCGTCT-----GGGGCGGAGGGGCCAC--AACAGAACCACGGC  
*Monocymbium* ???  
*Pdilatum6617* ???  
*PentaschistisaristifoliaSpies6* -CTCCGCAA-----GGACGGGGAGCGGCCGC--AAAAGAACCACGGC  
*Pfrank4705* -CACCGTCAA-----GGGGC-GAGGGGCCGC--AACAGAACCACGGC  
*Pschinz6652* ???  
*Sbic6588* -CTCCAAC-CT-----GGGGCAGAGGGGCCAC--AAAAGAACCACGGC  
*Sbic6589* -CTCCCTCT-----GGGGCAGAGGGGCCAC--AAAAGAACCACGGC  
*Shalep9318* -CTTC-TCAT-----GGAGGAGAGGGGCCAC--AAAAGAACCACAGGC  
*Snigros6567* ???  
*Tleuc6627* ???  
*Tspic4846* -CCCCGTCCC-----GGGGCGGAGGGGCCGC--AAAAGAACCACGGC  
*Ttria6555* -CTCCGTTC-----GGGGCGGAGGGGCCAC--AAAAGAACCACGGC  
*Zeadiplo* -GAACCTCCCG-----GGGGAAGGGGGG-CGCGAAAAGAACCACGGC  
*ZmaysGfl* -GAACCTCCCG-----GGGGAAGGGGGGCGCGAAAAGAACCACGGC  
*Saccharumofficinatum* CTCCGTCCC-----GGAGCGGAGGGGCCAC--AACAGAACCACGGC  
*Imperatocylindrica* CTCCGTCCC-----GGGGCGGAGGGGCCGC--AACAGAACCACGGC  
*Dichanthiumannulatum* CCTGTCTCC-----GGGGGGGGGGGCCCC--AAAAGAACCACGGC

*Aadscen6713* ???  
*Aameth6679* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*Aameth6685* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*Aappen6543* GCCTT---AAGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*Aappen6946* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*Adist6684* GCCTTCGTAAACATTTCAGGAACACTTA-TA-----TTG---CCTT---GC  
*Aeuco5869* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*Arundinellanepalensis* GCCAT---AGGCGTCAAGGAACACTCA-TA-----TTG---CCTT---GC  
*Aschiren6581* ???  
*Bbladh5173* GCCTT---AGGCGTCAAGGAACACTAC-TA-----TTG---CCTT---GC  
*Binsc5681* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*BBrachiaria* ???  
*Bradi4784* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*Bserra6554* ???  
*Cexc6577* ???  
*Cdieter6642* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*Cleistachnesorghoides* GCCTA---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTN---GC  
*Cmarg4397* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*Cplur6406* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*Cprolix6682* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT---GC  
*Danthoniacalifornica* GCCGAC---CGGCGCAAGGAACACTGTTA-----TTG---CCT---C  
*Dfili5872* GCCTT---AGGCGTCAAGGAACACTTG-TA-----TTG---CCTT---GC  
*Emuti5074* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---GCTC---GC  
*Emuti6715* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTC---GC  
*Eragrostisdielsii* GCCGA---CGGCGTCAAGGAACACTAA-TA-----TTG---CCTT---GC  
*Eviol5085* GCCTT---ACGAGTCAAGGAA-ACTTG-TA-----TTG---CCTT---GC

*Hanam4807* GCCTT---AGGCGTCAAGGAACACTTAATA-----TTG---CCTT--GC  
*Hanam4841* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Hcon4741* GCCTT---AGGCGTCAAGGAACACTTG-TA-----TTG---CCTT--GC  
*Hdiss5145* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Sbicolour9319* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Hfilli5144* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Hhirt6656* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Hlongi6560* ???  
*Hpilg4603* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Hpilg4738* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Hquar5088* GCCTT---AGGCGTCAAGGAACACTTAATA-----TTG---CCTT--GC  
*Hrud6576* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Htamb6585* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Icylin5114* GCCTA---AGGCGTCAAGGAACACTTC-TA-----TTG---CCTT--GC  
*Icylin6632* GCCTA---AGGCGTCAAGGAACACTTC-TA-----TTG---CCTT--GC  
*Mcape6655* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Monocymbium* ???  
*Pdilatum6617* ???  
*PentaschistisaristifoliaSpies6* GCCGTA---CGGCGTCAAGGAAACTGT-TA-----TTG---CCTT--GC  
*Pfrank4705* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Pschinz6652* ???  
*Sbic6588* GC---TT---AGGCGTCAAGGAACACTCA-TA-----TTG---CCTT--GC  
*Sbic6589* GCCTT---AGGCGTCAAGGAACACTCA-TG-----TTG---CCTT--GC  
*Shalep9318* GCCGA---AGGCGTCAAGGAACACTGA-TA-----TTG---CCTT--GC  
*Snigros6567* ???  
*Tleuc6627* ???  
*Tspic4846* GCCTT---AGGCGTCAAGGAACACTG-TA-----TTG---CCTT--GC  
*Ttria6555* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTT--GC  
*Zeadiplo* GCCTC---GGGCGCAAGGAACACCAG-TA-----CTA---CCTCCTGC  
*ZmaysGfl* GCCCC---GGGCGCAAGGAACACCAG-TA-----CTA---CCTCCTGC  
*Saccharumofficinarum* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTG---CCTTGC  
*Imperatacylindrica* GCCTT---AGGCGTCAAGGAACACTTC-TA-----TTG---CCTTGC  
*Dichanthiumannulatum* GCCTT---AGGCGTCAAGGAACACTTA-TA-----TTT---CCTTGCCC

*Aadscen6713* ???  
*Aameth6679* ACGGCGGGGCGGCTGGC-TTGCCCTCCGCTCCCGCGCAGC---GATCAT  
*Aameth6685* ACGGCGGGGCGGCTGGC-TTGCCCTCCGCTCCCGCGCAGC---GATCAT  
*Aappen6543* TCGGCGAGCCGCTCGGC-CTGCCTTCCGCTCCCGCGCAGC---GATGAT  
*Aappen6946* TCG-CGG-GCGGTCGGC-CTGCCTTCCGCTCCCGCGCAGC---GATGAT  
*Adist6684* ACTTCGGAGCGGCTGGC-TTGCCCTCCGCTCCCGCGCAAC---GATCAT  
*Aeuco5869* TCGGTGGAGCGGTCGGC-ATGCCTTCTGCTCCCGCGCAGC---GATGAT  
*Arundinellaneapalensis* ACGGTGGAGCGGTCGGC-CTGCCTTCTGCTCCCTGCGCAGC---GATGCT  
*Aschiren6581* ???  
*Bbladh5173* CCGGCGGAGCGGTCNNC-CTGCCTTCCGCTCCCGCGCAGC---GATGAT  
*Binsc5681* CCGGTGGAGCGGTCGGC-CTGCCTTCCGCTCCCGCGCA-C---GATGAT  
*BBrachiaria* ???  
*Bradi4784* TCGGCGGGGCGGTCGGC-CTGCCTTCCGCTCCCGCGCAGC---GATGAT  
*Bserra6554* ???  
*Cexc6577* ???  
*Cdieter6642* CCGGTGGAGCGGTCGGC-CTGCCTTCCGCTCCCGCGCAGC---GATGAT  
*Cleistachnesorghoides* NCGGCGGAGCGGTCGGC-NTGCCTTCCGCTCCCGCGCAGC---GATGAT  
*Cmarg4397* CCGGTGGAGCGGTCGGC-CTGCCTTCCGCTCCCGCGCAGC---GATGAT  
*Cplur6406* CCGGTGA-CGGTC--C-CTGCCTTCCACTCCCGCGCA-C---GATGAT  
*Cprolix6682* GCCGTGGAGCGTCAACTTCTCCCTCC-CTCCCGCGCAC---GATGAT  
*Danthoniacalifornica* GCGGCGGGCGGCCAGC-CTGCCGTCCTCACACGCGTA-----  
*Dfili5872* TCGGTGGAGCGGTCGGC-ATGCCTTCTGCTCCCGCGCAGC---GATGAT  
*Emuti5074* TCGGCGGAGCGGACGGC-CTGCCTTCCGCTCCCGCGCAGC---GATGTT  
*Emuti6715* TCGGCGGAGCGGACGGC-CTGCCTTCCGCTCCCGCGCAGC---GATGTT  
*Eragrostisdielsii* CCAATGACTCGACCGGC-CTGCCGGGCGCGTCTTGAGCAGC---GATACC  
*Eviol5085* TCGGTGGAGCGGTCGGC-GTGCCTTCCGCTCCCGGTGCAGC---GATAAT  
*Hanam4807* CCGGTGGAGCGGTCGGC-CTGCCTTCCGCTCCCGGTGCAGC---GATGAT  
*Hanam4841* CCGGTGGAGCGGTCAGC-TTGCCTTCCGCTCCCGGTGCAGC---GATGAT  
*Hcon4741* TCGGTGGAGCGGTCGGC-CTGCCTTCTGCTCCCGCGCGGC---GATGAT  
*Hdiss5145* AAGGTGGAGCGGTCGGC-CTGCCTTCCGCTCCCGGTGCAGC---GATGAT  
*Sbicolour9319* ACAGTGGAGCGGTCGGC-NTGCCTTGGGGTCCCGATGAAGAACAAGAT  
*Hfilli5144* CCGTGA--CGTCCGGC-CTGCCTTCCGCTCCCGGTGCAGC---GATGAT  
*Hhirt6656* CCGGTGGAGCGGTCGTC-TTGCCCTTCCGCTCCCGGTGCAGC---GATGAT  
*Hlongi6560* ???  
*Hpilg4603* CCGGTGGAGCGGTCGCC--TGCCCTTCCGCTCCCGCGCAGC---GATGAT  
*Hpilg4738* CCGGCGGAGCGGTCGGC-CTGCCTTCCGCTCCCGCGCAGC---GATGAT  
*Hquar5088* CCGGTGGAGCGGTCGGC-CTGCCTTCCGCTCCCGGTGCAGC---GATGAT

*Hrud6576* AAGGTTGGAACCAAGGC-TTCCGTGGGGGCACCGATGAAGA-CGGATGAT  
*Htamb6585* AAGGTGGAGCGGTCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Icylin5114* TCGGCGGAGCGGTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Icylin6632* TCGGCGGAGCGGCCG-C-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Mcape6655* TCGGCGG-ACGGTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Monocymbium* ???  
*Pdilatum6617* ???  
*PentaschistisaristifoliaSpies6* GCGTGGCGTGGCTGGC-CTGCCAGCCGCGCGCGCGCAGC---GATGAT  
*Pfrank4705* TCGACGGAGCGGTCGGC-CTGCCTTCCGCTCCCCGTGCAGC---GATGAT  
*Pschinz6652* ???  
*Sbic6588* GCAGTGGAG-GCTCGGC-CTCCGTGGTGCCGCGATGAAGAACGCAAGAT  
*Sbic6589* ACA-CGGAGTGGTCGGC-CTGCCTTCCGCTCCCTGAGCAGC---GATGAT  
*Shalep9318* TCGGGGCTTTGGTTGGC-TTGCCGATCATGCCTCGTGCAGC---GATGAT  
*Snigros6567* ???  
*Tleuc6627* ???  
*Tspic4846* CCGGCGGAGCGGTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Ttria6555* TCGGCGGAGCGGTCGGC-TTGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Zeadiplo* CCGGCGGAGCGGTCGGC-CCGCCTTCCGCTCCCCGGGCAGC---GGTTA-  
*ZmaysGfl* CCGGCGGAGCGGTCGGC-CCGCCTTCCGTTCCAGGGCAGC---GGTTA-  
*Saccharumofficinatum* ---GGTGGAGCGGTCGGC-CCGCCTTCCGCTCCCCGCGCAGC---GATGAT  
*Imperatacylindrica* ---GGCGGA-CGGTCGGC-CTGCCTTCCGCTCCCCGCGCAGA---GTGCT  
*Dichanthiumannulatum* ---GGTGGAGCGGTCGGC-CTGCCTTCCGCTCCCCGCGCAGC---GATGAT

*Aadscen6713* ???????????  
*Aameth6679* ATAC-TAATC  
*Aameth6685* ATAC-TAATC  
*Aappen6543* ATCT-TAATC  
*Aappen6946* ATCT-TAATC  
*Adist6684* ATCTCTAATC  
*Aeuco5869* ATCT-TAATC  
*Arundinellanepalensis* ATGT--AAC  
*Aschiren6581* ???????????  
*Bbladh5173* ATCTTTAATC  
*Binsc5681* ATCT-TAATC  
*BBrachiararia* ???????????  
*Bradi4784* ATCT-TAATC  
*Bserra6554* ???????????  
*Cdieter6642* ATCT-TAATC  
*Cexc6577* ???????????  
*Cleistachnesorghoides* ATCT-TAATC  
*Cmarg4397* ATCT-TAATC  
*Cplur6406* ATCT-TAATC  
*Cprolix6682* ATCT-TAATC  
*Danthoniacalifornica* ???????????  
*Dfili5872* ATCT-TAATC  
*Emuti5074* ATCT-TAATC  
*Emuti6715* ATCT-TAATC  
*Eragrostisdielsii* TTTTGTAATC  
*Eviol5085* ATCT-TAATC  
*Hanam4807* ATCTT-AATC  
*Hanam4841* ATCTT-AATC  
*Hcon4741* ATCT-TAATC  
*Hdiss5145* ATCTT-AATC  
*Sbicolour9319* ATCA-TCC--  
*Hfilli5144* ATCTT-AATC  
*Hhirt6656* ATCTT-AATC  
*Hlongi6560* ???????????  
*Hpilg4603* ATCT-TAATC  
*Hpilg4738* ATCT-TAATC  
*Hquar5088* ATCT-TAATC  
*Hrud6576* ATCT-TAATC  
*Htamb6585* ATCTTTAATC  
*Icylin5114* ATCT-TAATC  
*Icylin6632* ATCT-TAATC  
*Mcape6655* ACCT-TAAAC  
*Monocymbium* ???????????  
*Pdilatum6617* ???????????  
*PentaschistisaristifoliaSpies6* ATAC-TAATC  
*Pfrank4705* ATCT-TAATC  
*Pschinz6652* ???????????

<i>Sbic6588</i>	ATCT-TAATC
<i>Sbic6589</i>	ATCT-TAATC
<i>Shalep9318</i>	ATCTT-AATC
<i>Snigros6567</i>	??????????
<i>Tleuc6627</i>	??????????
<i>Tspic4846</i>	ATCT-TAATC
<i>Ttria6555</i>	TTAT-TAATC
<i>Zeadiplo</i>	CACCTTAGTC
<i>ZmaysGfl</i>	CACCTTAATC
<i>Saccharumofficinatum</i>	ATCT-TAATC
<i>Imperatacylindrica</i>	ATC-TTAATC
<i>Dichanthiumannulatum</i>	ATAC-TAATC
<i>Aadscen6713</i>	??
<i>Ameth6679</i>	--AA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Ameth6685</i>	--AA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Aappen6543</i>	--CAACACG-ACTCT--CGG-CAACGG-ATATCT-
<i>Aappen6946</i>	--CA-CACG-ACTCT--CGG-CAACGG-ATATCT-
<i>Adist6684</i>	--CA-CACG-ACTCT--CGGTCAACGG-ATATCT-
<i>Aeuco5869</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Arundinellaneapalensis</i>	--AA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Aschiren6581</i>	??
<i>Bbladh5173</i>	--CA-CACGAACCTCT---GG-GAACGGTATATCT-
<i>Binsc6681</i>	--CA-CA-C-ACTCT--CG--CAA-CGGATATCT-
<i>BBrachiaria</i>	??
<i>Bradi4784</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Bserra6554</i>	??
<i>Cdieter6642</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Cexc6577</i>	??
<i>Cleistachnesorghoides</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Cmarg4397</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Cplur6406</i>	--CA-CACG-ACTCT--CGG-CAACGG-ATATCT-
<i>Cprolix6682</i>	CACA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Danthoniacalifornica</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Dfili5872</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Emuti5074</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Emuti6715</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Eragrostisdielsii</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Evill5085</i>	--TA-CATG-ACTAT--CCG-CAA-CGGATATCT-
<i>Hanam4807</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Hanam4841</i>	--CA-CGCG-ACTCT--CGG-CAA-CGGATATCT-
<i>Hcon4741</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Hdiss5145</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Sbicolour9319</i>	C-----
<i>Hfilli5144</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Hhirt6656</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Hlongi6560</i>	??
<i>Hpilg4603</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Hpilg4738</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Hquar5088</i>	--CA-CACC-ACTCT--CAG-TAA-CGGATATCT-
<i>Hrud6576</i>	CACACGACTCTCGGCAACGGATATCTCGGC-----
<i>Htamb6585</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Icylin5114</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Icylin6632</i>	--CA-CACG-ACTCT--CGG-CAACGG-ATATCT-
<i>Mcape6655</i>	--CA-CACG-ACTCT--CGG-CAACGG-ATATCT-
<i>Monocymbium</i>	??
<i>Pdilatum6617</i>	??
<i>PentaschistisaristifoliaSpies6</i>	??
<i>Pfrank4705</i>	--GA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Pschinz6652</i>	??
<i>Sbic6588</i>	AA-----
<i>Sbic6589</i>	--CA-CACG-ACTCT--CGG-CAACGG-ATATCT-
<i>Shalep9318</i>	--CA-CACG-ACTCT--CGG-CAACGG-ATATCT-
<i>Snigros6567</i>	??
<i>Tleuc6627</i>	??
<i>Tspic4846</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Ttria6555</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Zeadiplo</i>	--GA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>ZmaysGfl</i>	--GA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Saccharumofficinatum</i>	--GA-CACG-ACTCT--CGG-CAA-CGGATATCT-
<i>Imperatacylindrica</i>	--CA-CACG-ACTCT--CGG-CAA-CGGATATCT-

*Dichanthiumannulatum*

--CA-CACG-ACTCT--CGG-CAA-CGGATATCN-

*Aadscen6713* ???  
*Ameth6679* -CGGCTC-TCGCATCGATGAAGAA-----  
*Ameth6685* -CGGCTC-TCGCATCGATGAAGAA-----  
*Aappen6543* -CGGCTC-TCGCATCGATGAAGAA-----  
*Aappen6946* -CGGCTC-TCGCATCGATGAAGAA-----  
*Adist6684* -CGGCTC-TCGCATCGATGAAGAA-----  
*Aeuco5869* -CGGCTC-TC-----  
*Arundinellanepalensis* -CGGCTC-TCGCATCGATGAAGAACGTAGCAAAA  
*Aschiren6581* ???  
*Bblad5173* -GGGCTC-TCGCATCGAAGAA-----  
*Binsc5681* -CGGCTC-TCGCATCGATGAAGAA-----  
*BBrachiaria* ???  
*Bradi4784* -CGGCTC-TCGCATCGATGAAGAA-----  
*Bserra6554* ???  
*Cdieter6642* -CGGCTC-TCGCATCGATGAAGAA-----  
*Cexc6577* ???  
*Cleistachnesorghoides* -CGGCTC-TCGCATCGATGAAGAACGTAGCAAAA  
*Cmarg4397* -CGGCTC-TCGCATCGATGAAGAA-----  
*Cplur6406* -CGGCTC-TCGCATCGATGAAGAA-----  
*Cprolix6682* -CGGATC-TCGCATCGATGAAGAA-----  
*Danthoniacalifornica* -CGGCTC-TCGCATCGATGAAGAACGTAGCAAAA  
*Dfili5872* -CGGCTC-TCGCATCGATGAAGAA-----  
*Emuti5074* -CGGCTC-TCGCATCGATGAAGAA-----  
*Emuti6715* -CGGCTC-TCGCATCGATGAAGAA-----  
*Eragrostisdielsii* -CGGCTC-TCGCATCGATGAAGAACGTAGCAAAA  
*Evill5085* -CGGCTC-TTGTATCGATGAAGAA-----  
*Hanam4807* -CGGCTC-TCGCATCGATGAAGAA-----  
*Hanam4841* -CGGCTC-TCGCATCGATGAAGAA-----  
*Hcon4741* -CGGCTC-TCGCATCGATGAAGAA-----  
*Hdiss5145* -CGGCTC-TCGCATCGATGAAGAA-----  
*Sbicolour9319* ???  
*Hfilli5144* -CGGCTC-TCGCATCGATGAAGAA-----  
*Hhirt6656* -CGGCTC-TCGCATCGATGAAGAA-----  
*Hlongi6560* ???  
*Hpilg4603* -CGGCTC-TCGCATCGATGAAGAA-----  
*Hpilg4738* -CGGCTC-TCGCATCGATGAAGAA-----  
*Hquar5088* -CGGTTT-TCGCATCGATGAAGAA-----  
*Hrud6576* ???  
*Htamb6585* -CGGCTC-TCCATTCGATGAAGAA-----  
*Icylin5114* -CGGCTC-TCGCATCGATGAAGAA-----  
*Icylin6632* -CGGCTC-TCGCATCGATGAAGAA-----  
*Mcape6655* -CGGCTC-TCGCATCGATGAAGAA-----  
*Monocymbium* -----  
*Pdilatum6617* ???  
*PentastichistisaristifoliaSpies6* ???  
*Pfrank4705* -CGGCTC-TCGCATCGATGAAGAA-----  
*Pschinz6652* ???  
*Sbic6588* ???  
*Sbic6589* -CGGCTC-TCGCATCGATGAAGAA-----  
*Shalep9318* -CGGCTC-TCGCATCGATGAAGAA-----  
*Snigros6567* ???  
*Tleuc6627* ???  
*Tspic4846* -CGGCTC-TCGCATCGATGAAGAA-----  
*Ttria6555* -CGGCTC-TCGCATCGATGAAGAA-----  
*Zeadiplo* -CGGCTC-TCGCATCGATGAAGAACGTAGCAAAA  
*ZmaysGf1* -CGGCTC-TCGCATCGATGAAGAACGTAGCAAAA  
*Saccharumofficinatum* -CGGCTC-TCGCATCGATGAAGAACGTAGCAAAA  
*Imperatacylindrica* -CGGCTC-TCGCATCGATGAAGAACGTAGCAAAA  
*Dichanthiumannulatum* -NNNNNN-NNNNNNNNNNNNNNNNNNNNNNNNNNNAAAA

*Aadscen6713* TGCG-ATACCTGGTGT-GAA-TTGC  
*Aameth6679* ?????????????????????????????????????  
*Aameth6685* TGCG-ATACCTAGTGT-GAA-TTGC  
*Aappen6543* TGCGGATACGTGGCGTGCAA-TTGC  
*Aappen6946* ?????????????????????????????????????  
*Adist6684* TNCCGATACCTAGCGCGAATTCGC  
*Aeuco5869* TGCG-ATACCTGGTGT-GAA-TTGC

*Arundinellanepalensis*  
*Aschiren6581*  
*Bbladh5173*  
*Binsc5681*  
*BBrachiaria*  
*Bradi4784*  
*Bserra6554*  
*Cdieter6642*  
*Cexc6577*  
*Cleistachnesorghoides*  
*Cmarg4397*  
*Cplur6406*  
*Cprolix6682*  
*Danthoniacalifornica*  
*Dfili5872*  
*Emuti5074*  
*Emuti6715*  
*Eragrostisdielsii*  
*Eviol5085*  
*Hanam4807*  
*Hanam4841*  
*Hcon4741*  
*Hdiss5145*  
*Sbicolour9319*  
*Hfilli5144*  
*Hhirt6656*  
*Hlongi6560*  
*Hpilg4603*  
*Hplig4738*  
*Hquar5088*  
*Hrud6576*  
*Htamb6585*  
*Icylin5114*  
*Icylin6632*  
*Mcape6655*  
*Monocymbium*  
*Pdilatum6617*  
*PentaschistisaristifoliaSpies6*  
*Pfrank4705*  
*Pschinz6652*  
*Sbic6588*  
*Sbic6589*  
*Shalep9318*  
*Snigros6567*  
*Tleuc6627*  
*Tspic4846*  
*Ttria6555*  
*Zeadiplo*  
*ZmaysGfl*  
*Saccharumofficinarum*  
*Imperatacylindrica*  
*Dichanthiumannulatum*

TGCG-ATACCTGGTGT-GAA-TTGC  
ATGCGATACCTAGCGT-CAA-TTCC  
TGCCGT--CCTAGCGTTNA-TTCCC  
TGCGGATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAAATTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGCGGATACCTGGTGT-GTAATTGC  
????????????????????  
GTGCCATACCTAGTGT-CAA--TTC  
TGCG-ATACCTGGTGT-GAA-TTGC  
-ATCGATACCTAGTGT-GAA--TTC  
TGCCGATACCTAGCGGTGAA-TTCC  
TGCGGATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TACGGATACCTGGTGT-GAA-TGCC  
TGCCGATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
ATNCGATACCTAGTNT-GAA-TTGC  
TGCGGATACCTGGTGT-GAA-TTGC  
TGCT-ATACCTCGAGT-GAA-CTGC  
TGCGGATACCTGGTGT-GAA-TTGC  
TACGGATACCTAGCGCTCAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGGCCCCACATGCCTATAC----GC  
TGCGGATACCTGGTGT-GAA-TTGC  
TGCGGATACCTGGTGT-GAA-TTGC  
TGCGNATACCTGGTGT-GAA-TTGC  
????????????????????  
TGCGGATACCTGGTGT-GAA-TTGC  
TACG-ATACCTGGTGTGTACTTGC  
TCCCGATACCTAGCGGCTA--TTGC  
TGCGGATACCTAGTGT-GAA-TTGC  
TGCGGATACCTGGTGTGCTGAA-TTGC  
TGCGGATACCTGGTGT-GAA-TTGC  
T-CG-ATACCTAGTGT-CAA-TTGC  
TCG--ATACCTGGTGT-GAA-TTGC  
????????????????????  
????????????????????  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGCGGATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAC-TTGC  
TTCCACGTCGACCTATGACACTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
---GGATACTGGTGTGCTGAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC  
TGCG-ATACCTGGTGT-GAA-TTGC

*Aadscen6713*  
*Aameth6679*  
*Aameth6685*  
*Aappen6543*  
*Aappen6946*  
*Adist6684*  
*Aeuco5869*  
*Arundinellanepalensis*  
*Aschiren6581*  
*Bbladh5173*  
*Binsc5681*  
*BBrachiaria*  
*Bradi4784*  
*Bserra6554*  
*Cdieter6642*  
*Cexc6577*  
*Cleistachnesorghoides*

ATAAATCCCG--TGTAACCATCG--AGTTTTTGAA-CGCAAGTTGCGCCC  
????????????????????  
ATACATCCCG--CTGAACCATCT--AGTTTTTAAACGCAAGTTGCGCCC  
ATAAATCCCG--CGAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCGCCC  
????????????????????  
ATTCATGTGCCCTCGAATGGATCCATTATTTCCGACTGCAACTTGCCGCC  
A--AATCCCG--CGAA-CCATCG--AGTTTTTGAA-CGCAAGTTGCGCCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCGCCC  
ATAAATCCCG--TCGAATGCATCCC--ATTTTTTCAACTCAACTTGCCCC  
--TAATGTGC-TC-AATGGGTCC--TTATGGCCGAATACACCG-GCCGCG  
ATAAATCCCG--CGAAACCATCG--AGTTTTTGAA-CGCAAGTTGCGCCC  
ATAAATCCCG--CGAAACCATCG-AGTGTGTGAA-CGCAAGTTGCGCCC  
ATTAATCCCG--CGAAACCATCG--AGTTTTTTAA-CGCAACTTGCCCC  
ATAA-TCCCG--CGTAACCATCG--AGTTTTTGAA-CGCAAGTTGCGCCC  
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A-TAATCCCG-TC--AACGATCCC--ATTTGTTCAAACCAACTTGCCCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAA-CGCAAGTTGCGCCC

Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Emuti5074  
Emuti6715  
Eragrostisdielsii  
Evill5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdiss5145  
Sbicolour9319  
Hfilli5144  
Hhirt6656  
Hlongi6560  
Hplig4603  
Hplig4738  
Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Monocymbium  
Pdilatum6617  
PentastichistisaristifoliaSpies6  
Pfrank4705  
Pschinz6652  
Sbic6588  
Sbic6589  
Shalep9318  
Snigros6567  
Tleuc6627  
Tspic4846  
Ttria6555  
Zeadiplo  
ZmaysGfl  
Saccharumofficinatum  
Imperatocylindrica  
Dichanthiumannulatum

CATAATCCCG--CGAACC-ATCGA---GTTTTTGAACGCAAGTTGCCCC  
AT-AATCCCG--C-NAATCATCN--ATTTTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--C--AACCATCGG-AGTTTTTGAACGCAAGTTGCCCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
ATAAATCCCN--CCAA-CCATCG--AGTTTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--C-GAACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--C-AACCATCGC-AGTTTTTGAACGCAAGTTGCCCC  
AG-AATCCCG--T-GAACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
ATGAATCCCG--TCCAATC-ATCGCAGTTTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--C--AACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
AGAAATCCCG--ACGAACCATCGA-GATTCTTC-AACGCAAGTTGCCCC  
ATAAATCCCG--CGAA-CCATCGC-AGTTTTTGAACGCAAGTTGCCCC  
ATTAATCCCG--GTCAACCATCGC-ATTTTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--CA-AACCATCGG-AGTTTTTGAACGCAAGTTGCCCC  
--TACTGTGC-TC-CATAGTTC---TTATGTCCACGAATCTAGCCCC  
ATAA-TCCCG--CGAA-CCATCG--AGTTTTTGAACGCAAGTTGCCCC  
AGCAATCCCG--CGGAACCATCG--AGTCTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--CG-AACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
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ATAAATCCCG--CGAA-CCATCG--AGTTTTTGAACGCAAGTTGCCCC  
ATAAATCCCT--C-CAACCATCGA-AGTTTTTGAACGCAAGTTGCCCC  
--TAATCCCG-TC-AATCGATTCC---TTTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--C-CAATCATCC--ATTTTTTGAACGCAAGTTGCCCC  
AT-AATCCCG--C-GAACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--CGCAACCATCN--AGTTTTTGAACGCAAGTTGCCCC  
ATAA--TCTG--GTCAATGGTCC--ATTTTTTCAA-CTCAAGTTGCCCC  
ATAA-TCCCG--CGAA-CCATCC--AATTTTTGAACGCAAGTTGCCCC  
??  
??  
ATAA-TCCCG--CG-AACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--CG-AACCATCGG-AGTTTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--CG-AACCATCGG-AGTTTTTGAACGCAAGTTGCCCC  
ATAA-TCCCG--CGAA-CCATCG--AGTTTGTGAACGCAAGTTGCCCT  
--TAATTCG--TC-AATGGATTGC---ATTTGTCAATAAACCA-GCTACA  
AGAAATCCCG--CG-AACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
AT-AATCCCG--CGCAACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
AGAA-TCCCG--CGCAACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
AG-AATCCCG--C-GAACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
AG-AATCCCG--T-GAACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
AG-AATCCCG--CG-AACCATCG--AGTTTTTGAACGCAAGTTGCCCC  
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AG-AATCCCG--CG-AATCATCG--AGTTTTTGAACGCAAGTTGCCCC

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Aameth6685  
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Aeuco5869  
Arundinellaneapalensis  
Aschiren6581  
Bbladh5173  
Binsc5681  
BBrachiaria  
Bradi4784  
Bserra6554  
Cdieter6642  
Cexc6577  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Emuti5074  
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Eragrostisdielsii  
Evill5085  
Hanam4807

G-AAGCC-TTCTGGTCA-GGGCAGCCTGCCTGGGCG--TCACGCC---  
??  
C-AGGCC-TTCTGGTCA-GGACACCTCTTCTGGGCCG-TCACGCC---  
G-AGGCC-TTCTGGTCA-GGGCACCCTGCCTGGGCG--TCACGCC--  
??  
TGACCCCGTTCCTTTCACACGACACCAATTGCATGTGCGGTGAT  
G-AAGCC-TTTTGGTCA-GGGCAGCCTGCCTGGGCG--TCACGCC---  
G-AAGCC-TCCAGGCCG-GGGCAGCCTGCCTGGGCG--TCACGCC---  
C-ACGCC-TTCTTTGAC--GACACAAGTGTGCGCGTCAACAAA--  
T-ACCGGGTCTTTCAC-CGACAACAATGGGA-GTCGCC-CACCAG-T  
G-AGGCC-TTCTGGTCA-GGGCAGCCTGCCTGGGCG--TCACGCC---  
C-ACGCC-TTCTGGTCA-GGGCAGCCTGCCTGGGCG--TCACGCC---  
T-ATGCC-TTCTGNTCAA-GGGCACCCTGCCTGGGCG--TCACGCC--  
G-AGGCC-TTCTGGTCA-GGGCAGCCTGCCTGGGCG--TCACGCC---  
??  
T-ACGCC-TTCTTGCAA-CGACACCTATTGCA-TGGCCGTACCCCA--  
G-AGGCC-TTCTGGCCG-GGGCAGCCTGCCTGGGCG--TCACGCC---  
GGACGCC-TTCTGTTCAA-CGACACCTTGC--TGGCGTCAACCCA--  
G-ACGCC-TTCTTGTCAA-GGGCACATCTGCCTGGGCG--TAACAC---  
G--AGCC-TTCTGGTC-A-GGGCACCTGCCTGGGCG--TCACGCC--  
G-AGGCC-TTCTGGCCG-GGGCAGCCTGCCTGGGCG--TCACGCC---  
G-AAGCC-TTTTGGTCA-GGGCAC-TCTGCCTGGGCG--TCACGCC---  
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G-AGGCC-TTCTGGTCA-GGGCAGCCTGCCTGGGCG--TCACGCC---  
G-ACGCC-TTCTGGTCA-GGGCAGCCTGCCTGGGCG--TCACGCC---  
A--CGCC-TTCTGGTCAA-GGGCACCTGCCTGGGCG--TCACGCC---

Hanam4841  
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Hdiss5145  
Sbicolour9319  
Hfilli5144  
Hhirt6656  
Hlongi6560  
Hpilg4603  
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Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Monocymbium  
Pdilatatum6617  
PentaschistisaristifoliaSpies6  
Pfrank4705  
Pschinz6652  
Sbic6588  
Sbic6589  
Shalep9318  
Snigros6567  
Tleuc6627  
Tspic4846  
Ttria6555  
Zeadiplo  
ZmaysGfl  
Saccharumofficinarium  
Imperatacylindrica  
Dichanthiumannulatum

C-ACGTC-TTCTGGTCTGA-CGGCACCTCTGCCTGAGCG--TCAA-TCT--  
G-AGGCC-TTCTGGTCTGA-GGGCACGCTGCCTGGGCG--TCACGCC---  
TGACGCC-TTCTGGTCTGA-CGACACATCTGATGGGCA--TCACAGCC--  
G-AGGCC-TTCCGGCCGA-GGGCACGCTGCCTGGGCG--TCACGCCA--  
C-ACGAC-TTCTGGTCTGC-ACACACCTTTC---TTCTCGCCCCAAGTT  
G-A-GCC-TTCTGGTCTGA-GGGCACGCTGCCTGGGCG--TCACGCC---  
G-AAGCC-TTCTGGTCTGA-GGGCACGCTGCCTGGGCG--TCACGCC---  
G-AGGCC-TTCTGGTCTGA-GGGCACGCTGCCTGGGCG--TCACGCC---  
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G-ATGCC-TTCTGGTCTGA-GGGCACGCTGCCTGGGCG--TCACGCC---  
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C-ACGCC-TTCTGGTCTGA-GGGCACCTCTGCCTGGGCG--TCACGCCA--  
GTAGGCC-TTCTGGTCTGA-GGGCACGCTGCCTGGGCG--TCACGCC---  
G-AAGCC-TTCTGGCCGA-AGGCACNTCTGCCTGGGCG--TCACGCC---  
T-ACGCC-TTCTGGTCTGA-CGACACCTATGCCTGGGCG--TCACGCCA--  
G-AGGCC-TTCTGGTCTGA-GGGCACGCTGCCTGGGCG--TCACGCC---  
??  
??  
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G-AG-CC-TTCCGGCCGA-GGGCACGCTGCCTGGGCG--TCACGCC---  
G-AGGCC-TTCCGGCCGA-GGGCACGCTGCCTGGGCG--TCACGCCA--  
G-ACGCC-TTCTGGTCTGA-GGGCACGCTGCCTGGGCG--TCACGCC---  
TGACGCC-TTCTGGTCTGA-GGCAACACCTACCATGTGCCACCACCG-T  
G-AGGCC-TTCTGGCCTA-GGGCACGCTGCCTGGGCG--TCACGCC---  
GAAGGCC-TTCTGGTCTGA-GGGCACGCTGCCTGGGCG--TCACGCC---  
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G-AGGCC-TTCTGGTCTGA-GGGCACGCTGCCTGGGCG--TCACGCC---

Aadscen6713  
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Aeuco5869  
Arundinellaneapalensis  
Aschiren6581  
Bblad5173  
Binsc5681  
BBrachiaria  
Bradi4784  
Bserra6554  
Cdieter6642  
Cexc6577  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Emuti5074  
Emuti6715  
Eragrostisdielsii  
Evill5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdiss5145  
Sbicolour9319  
Hfilli5144  
Hhirt6656  
Hlongi6560  
Hpilg4603  
Hplig4738  
Hquar5088

AAAAGAC-ACTCCC-AACCCA---TCCATGGGG---ACGG-ACGTGGT-  
??  
ATAGCAC-ACTCACCAGATGAC---CCTCGGCC---ACAGATTTGGT-  
AAAACAC-ACTCCC-AACCCA---AATCGGGG---AGGGATGTG-T-  
??  
GAATGAC-ACGTCGACACGATCTTTC-GGCAGACGG---CACACTTTGG-  
AAAAGAC-ACGTCCCAAGCCAC---AATCAAG---GAGGTA-CTGGC-  
AAAAGAC-ACTCCC-AACCCAACC---CAGGGG---AGGGATGTGGC-  
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GC-TGTG-CC---CGGCACGGCTATCC-GGCAACC---CAAATCTGG-  
AAAAGAC-ACTCCCAACCAC---CCTCGGGG---AGGGCACGTGGT-  
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AAATATG-ACTCGGAGACCATTCCT-CGCAGCA---CGACCTTCG-  
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AAAAGAC-GCTCCC---ACCCAACC---CCGGT---AGGACCGGGC-  
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AAAANAC-ACTCCC-AAACCAC---AATCGTGG---AGGGAC-TGGT-  
AAAACACTACTCCC-AACCCAC---NCTCGTGG---AGGGACATGTT-  
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-AAAGAC-ACTCCAGTCCCTACCCTTGACAGGGCA---TCTACTTCTT-  
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TATCTAC-AAGTCG-ACGTAACCGTC---GGGA---CGGACCTGGT-  
AAAACAC-A-CTCCCAACCCAA---AATCGGG---CGGGATGTGGT-  
TAATAAC-ACTCAC-AACATAACAATC---CGCG---AGCACCTTCT-  
ACATTAAC-ACTCCG-AACCCA---CCCTCGGG---AGGGATGTGGT-  
TC-TGTG-CC---CAATCCACTTGTG-GCCAGG---CCCACTTTTG-  
AAAA-AC-ACTCCC-AACCCAC---CCTCGGG---ACGGACATGGT-  
-AAACAC-GCTCCC-ACCCACTAACCTGTGGT---GGGATCGGGC-  
AAAAGAC-ACTCCCAACCCGT---ACTCGGG---AGGGACGTGGT-  
??  
AAAAGAC-AGTCCC-AACCCAC---CATCGGG---ACGGACGTGGT-



Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Monocymbium  
Pdilatam6617  
PentaschistisaristifoliaSpies6  
Pfrank4705  
Pschinz6652  
Sbic6588  
Sbic6589  
Shalep9318  
Snigros6567  
Tleuc6627  
Tspic4846  
Ttria6555  
Zeadiplo  
ZmaysGfl  
Saccharumofficinatum  
Imperatacylindrica  
Dichanthiumannulatum

AAAATAC-ACTCCC-AACACACCATC----GGGG-----AGGACCTGGT-  
GCTGTAC-ACTGCCACAAC-CACTAT-GGCAGTAG----CTACCTTCTG-  
AAATGAT-ACTCGCCAACGCGCAT--CCTCGGGG----AGGACATTT-TC  
AAAAGAC-ACTCCC-AACCCACCC----TCGGGG----AGGGACATGGT-  
AAAACAC-ACTCCCAAACCCAC----ACAAGG-----AAAGGACATTTT-  
TAATGAT-ACTCCG-AACCCCTCG--CAAGGCGG----ACGGACATTTT-  
AAAAGAC-ACTCCC-AACCCA---TCATAGGGT-----GGGACGATTTA  
??  
--GTACC-CGCTCAGATACTAG----CAGCAA---ATAGGACTTCT-  
AAAAGAC-ACTCCT-ACCCCAT--ACTATGGGGT---ACGG-ATGTGGT-  
ACAGTAC-ACTCCC-AACCCA----GCCTCGGGG---AGGG-ATGTGGT-  
ACAG-AC-ACTCCC-AACCCA----GGCTCGGGG---AGGG-ATGTGGT-  
AAAAGAC-ACTCCC-AACCCA---TCCCTGGGT----AGGG-ACCTGGT-  
GC--GGC-GCTCGAACAAGTTTGTCT--GGCAGCC-----CAAACCTCG--  
AAAAGAC-GCTCCC-AACCCA---ACCAAGGG---AGGG-ACGCTGT-  
AAAAGAC-ACTCCC-AACCCACCC----TCGGGGG---AGGGACGTGGT-  
AAAAGAC-ACTCCCAAACCCAC-----CCTAGGGG---AGGG-ACGTGGT-  
AAAAGAC-ACTCCC-AACACCCCC--CCTGCGGGG---GAGGGACGTGGC-  
AAAAGAC-ACTCCCAAACCCAC----CCGAGGGG---AGGG-ACGTGGT-  
AAAAGAC-ACTCCCAAACCCAC----CCTCGGGG---AGGG-ACGTGGT-  
AAAAGAC-ACTCCCAAACCCAA----CCTCGGGGGG-----ACGTTGT-

Aadscen6713  
Aameth6679  
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Aappen6543  
Aappen6946  
Adist6684  
Aeuco5869  
Arundinellanepalensis  
Aschiren6581  
Bbladh5173  
Binsc5681  
BBrachiaria  
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Cdieter6642  
Cexc6577  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Emuti5074  
Emuti6715  
Eragrostisdielsii  
Eviol5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdiss5145  
Sbicolour9319  
Hfilli5144  
Hhirt6656  
Hlongi6560  
Hpilg4603  
Hplig4738  
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Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Monocymbium  
Pdilatam6617  
PentaschistisaristifoliaSpies6  
Pfrank4705  
Pschinz6652

GTTTGGCT--CCTCGTGC-CGTAAGGT---GCG-GTGGGCCG-AAGTT--  
??  
GTTTCGCC--CCTCGTGG-CACGAGGGGCGAGC---ACCACAAGTT--  
GTTTGGCC--CCTCGTGC-CGCAAGG---CGCG--TGGGCG-AAGTT--  
??  
-TTTACC-CCTCCTTGC-GGCAAGGA---GGGGCCGAACACAACGTG--  
GTATGGCC--CCCCGTGC-CGCACGG---CGCGG-TGGGCC-AAATT--  
GTATGGCC--CCCCGTGC-CGCATGG---CGCGG-TGGGCC-AAATT--  
-TT-TACC-GCCCCCTGC-CGCACGG---GG-GCCGAACACCAAGTC--  
-CT-CACC-TCGCCCTGC-GGCA-GGG---GGAGCCAACACCCAC-TG--  
GTTTGGCT--CCCCGTGC-CGCAGGG---CGCAG-TGGGCC-AAATT--  
GTTTGGCC--CCCCGTGC-CGCAAGGA---CGCGT--GGGCC-AAATT--  
-TTTTCGCTCCCTTCC-GGCAAGGG---CGAGGCTGAAACACAACCTC-  
GTTTGGCT--CCCCCTAC-CGCAAGG---GCGGT--GGGCC-AAATT--  
??  
-GTCTAGC-GCCCCCTGC-CGCACGG---GG-GCCAGACCCCAACTT--  
GTTTGGCC--TCCCGTGC-CTCACGG---CGCGG-TGGGCC-AAATT--  
-TT-GACC-GCGCCCTGC-GGCAAGGG---GGGGCCAGACACACTG--  
GTCTGGCC--CCCCGCCCCGGCAGGG---CGAAG-TGGGCCCAATAT--  
GTCTGGCC--CCCCGTGC-CGCAAGG---CGGGG-TGGGCCAAAGTT--  
GTTTGGCT--CCCCGTGC-CGCAGGG---CGCGG-TGGGCC-AAATT--  
GTATGGCC--CCCCGTGC-CGCACGG---GGCGG-TGGGCC-AAATT--  
GTTTGGCC--CCTCGCGC-CTCAGG---CGCGG-TGGGCC-AAATTT-  
GTTTGGCC--CCTC-CGC-CTCANGG---CACGG-GGGCC-AAATTT-  
ATTTGGCT--CCCCGTGC-AACACGA---CACGG-TGGGCC-AAATT--  
-TCCTTCC-TCCCTTGC-AGGA-CGA---GGAACCAACACCATGAT--  
GTTTGGCC--CCCCGTGC-CGCAAGG---CGGGG-TGACCACAACCT--  
NTT-----  
GTTTGGCC--TCCCGTGC-CGCAAGG---CGCGG-TAGGCC-AAATT--  
CTTTGCC--CCCCGTAC-CATACGA---CGGGGTGAACACAACCTT--  
TTTTGGCT--CCCCGATGCCGATGGC---GTGGTGGGCC-AAATT--  
-GCCAAC-CCACCTTAT-TGAACGTT-----GGGCAACACCCCTT--  
GTTTGGCC--CCCCATGC-CGTATGG---CGCGG-TGGGCC-AAATT--  
ATGTGGCT--CCCCGT-CTCGCAAGGG---GCGGTG-GGCCTC-AAGAT--  
GTTTGGCC--CCCCGTGC-CGCAGGG---CGCGG-TGGGCC-AAATT--  
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GTTTGGCC--CCCCGTGC-CGTACGG---CGCGG-TGGGCC-AAATT--  
GTTTGGCC--CCCCCTGC-CATACGG---CGGGC-GGACACGAAATT--  
-CTCTACC-ACCCCTGC-CGCACGA---CGG-GGCATACCAACAT--  
GTTTGGAC-CCCCCTGCCGGCAGGG---GGCGGTGGGACACAAGTTG-  
GTTTGGAC--CCCCGCTC-CGCAGGG---CGCGG-TGGGCC-AAATT--  
GTTTCGTC--CCCCCTC-CGCAAGG---CGCGA-TGGGCC-AAATT--  
GTTTGGAC--CCCCGTG-CGCAAGG---GGGGTTCAGACACAACCTG--  
GTTTGGCC--TCTCGTGC-CGTGTGC---ACAGG-TGGGCC-AAATC--  
??  
GTTTGGAT--CCCCGTGC-CGCATGGGCACAGGT-CGGGCC-AAAGTT-  
GTTTGGCT--CCCCCTGC-CTTGGC---CATG-GTGGGCTA-AAGTT--

*Sbic6588*  
*Sbic6589*  
*Shalep9318*  
*Snigros6567*  
*Tleuc6627*  
*Tspic4846*  
*Ttria6555*  
*Zeadiplo*  
*ZmaysGfl*  
*Saccharumofficinarum*  
*Imperatacylindrica*  
*Dichanthiumannulatum*

TTTTGGCT--CCCCGATGCCGATGG---CGCG-GTGGGCC--AAGTT--  
TTTTGGCT--CCCCGA-GCCGATGG---CGCG-GTGGGCC--AAGTT--  
GTTTGGCT--CCCTGTGC-TGCAAAG---CGCG-GGGGGCTG-AAATT--  
-TTCCACC-CCACCATCT-AGCT-CG---AGCAGCCAACACCACGTG--  
GTTTGGCT--CCCCGTGC-CGCAAGG---CGCG-GTGGGCC--AAGAT--  
GTTTGGCC--CCCCGTGC-CGCAGGG---CGCG-GTGGGCC--AATT--  
GTTTGGCT--CCCCGTGC-CGCAGGG---CGCG-GGGGCCA-AAATT--  
GTCTGGCC--CCCCACGC-CGCAGGG---CGAGG-CGGGCCG-AAGCA--  
GTCTGGCC--CCACGCG-CGCAGGG---CGAGG-TGGGCCG-AAGCA--  
GTTTGGTC--CCCCGTGC-CGCAGGG---CGCG-GTGGGCC--AAGTT--  
GTTTGGAC--CCCCCGC-CGCAGGG---CGCG-GTGGGCC--AAGTT--  
GTTTGGCC--CCCCNNTC-CCNAAGG---GGNG-GGGGCC--AAGTT--

*Aadscen6713*  
*Aameth6679*  
*Aameth6685*  
*Aappen6543*  
*Aappen6946*  
*Adist6684*  
*Aeuco5869*  
*Arundinellaneapalensis*  
*Aschiren6581*  
*Bbladh5173*  
*Binsc5681*  
*BBrachiaria*  
*Bradi4784*  
*Bserra6554*  
*Cdieter6642*  
*Cexc6577*  
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*Cplur6406*  
*Cprolix6682*  
*Danthoniacalifornica*  
*Dfili5872*  
*Emuti5074*  
*Emuti6715*  
*Eragrostisdielsii*  
*Evill5085*  
*Hanam4807*  
*Hanam4841*  
*Hcon4741*  
*Hdiss5145*  
*Sbicolour9319*  
*Hfilli5144*  
*Hhirt6656*  
*Hlongi6560*  
*Hpilg4603*  
*Hplig4738*  
*Hquar5088*  
*Hrud6576*  
*Htamb6585*  
*Icylin5114*  
*Icylin6632*  
*Mcape6655*  
*Monocymbium*  
*Pdilatum6617*  
*PentaschistisaristifoliaSpies6*  
*Pfrank4705*  
*Pschinz6652*  
*Sbic6588*  
*Sbic6589*  
*Shalep9318*  
*Snigros6567*  
*Tleuc6627*  
*Tspic4846*  
*Ttria6555*  
*Zeadiplo*  
*ZmaysGfl*  
*Saccharumofficinarum*

GGGGCTGCCGGC--ATATCTT-GCCGGG-CACC-GCAGC-TGGTGGGGC  
??  
CTAGCTGACGGGTGATACTCGT-GACTGT-CTTTGGAGC-TGACTGGCC  
GGGGCTGCCG-G-CTAATCG-TGCCGGG-CACA-TCACG-TGGTGGGGC  
??  
CCCTCTCCAGG--GTAG-TCGTGTGCGAGTG-ACTTTGGCTG-TGAGC  
GGGGCTGCCGGC--T-AATCC-TGCCAGG-CACA-GCACC-TGGTGGGC-  
GGGGCTGCCGG--CAAACCGT-TCCGGG-CACA-GCATG-TGGTGGGGC  
CC-TCCTGCAAG--GCTAATCTGGG--AGTG-TATTTAGCG-T-GACG  
CCCTC-CCCTTG--GGAG-GCTTGG--GAGTG-ACTTTGGCTG-GACG  
GGGGCTGCCGGC--G-AATCG-TGTCGAG-CACA-GCAGC-TGGTGGGGC  
GGGGCTGCCGGC--ATAACTT-GACTGG-CACC-TCACG-TGGTGGGCA  
CGGTCTGCCGAGCGTAATCTTGGCAGGTGACA-TTACG-TGGTTGACG  
GGAGCTGCCGGC--GTACCTT-GCCGGG-CACC-GCAGC-TGGTGGGGC  
??  
CCCTCTCCAGG--GTAG-TGTGCG--GAGTG-TCTTTGGCGTT-GAGC  
GGGGCTGCCGG--CGAATCGT-TCCGGG-CACA-GCAGC-TGGTGGGGC  
CCCTT-GCCGAG--GGAG-TGTTGG--GAGTG-ACTTTCCGGTG-GACG  
GGGGCTGCCAGC--TAATCT-CGCCAGG-CACA-TTACC-TGGTCCGGC  
GGGGCTGCCGGC--GGAATCGTCCGGGGGACA-TCACG-TGGTGGGGC  
GGGGCTGCCGG--CGTACCGT-CGGGG-CGCA-GCACA-AGGTGGGGC  
GGGGCTGCCAGC--A-AATCT-TGCCAGG-CACA-TCACG-TGGTGGGGC  
GGGGCTGCCGG--CATATCG-CGTCGGG-CACA-TCACG-TGGTGGGGC  
GGGGCTGCCGGA--CAGATCG-CGTCGGA-CACA-TCTT--TGGTGGGGC  
GGGGCTGCCGG--CTTACGGTA-CCGGT-CACA-GCACA-AGGTGGGTG  
CCCTC--CCGG--GGAGTGCTGGC--AGTG-TCTTTCCGGT-T-GACG  
GCGGCTCCGAC--GGAATCGTCCGAGTGACA-TTACG-TGGTGAGCC  
??  
GGGGCTGCCGGC--G-AATCG-TGTCGGG-CACA-GCAGC-TGGTGGGC-  
CGGCTTCCAGA-TTATCTATCTTA-GTGAATTTT--TCGGTATGAC  
GGGGCTGCCGGC--GATTCTG-TCCGGG-CACA-TCACG-TGGTGGGGC  
CCTTC-CCCAAG--GTAC-CCTTGG--AATTT-ACTTTGTGATT-AACC  
GGGGCTGTGC-A--CGAATCG-TGTCGGG-CACA-TCACG-TGGTGGGGC  
CGGCTGCCGGT--GTATCGTGACCGGA-CACA-GCGCG-TGGTAGCGC  
GGGGCTGCCGGC--GGAATCG-CGCCGGG-CACA-GCAGC-TGGTGGGGC  
??  
GGGGCTGCCA-A--CGAATCG-TGTCAGG-CACA-GCAGC-TGGTGGGGC  
GGCTCTCCATGC-CTATCTGTGGCAGTGCATTTAGCTAGTACGCN  
CTCTCTGCCATG--CATGGCCGTGCCAGGTGCATTTAGCT-T-AACG  
GGTCTGCCGAG--GGAGCCTTGGCGAGTGA--CTTTTGGTGGTTGACG  
GGGGCTGCCGG--CGAA-CGTG-TCCGGG-CACA-TGACG-TGGTGGGGC  
GGGGCTGCCAGC--GAATCGT-GTCAGA-CACA-TTATG-TGGTGGACC  
-GGGTTGCCCTT--GGAGTCTTGGCGAGTGACTTTT--GGTGGT-GACG  
TTGGCTGCCGG--TGATCTT-GCTGGG-CACC-TCATA-TGGTGGGGC  
??  
GGGGCTGCCGGC--ATAACCG-TGTCGGG-CACA-GCATT-CTTGGGGC  
GTGGCTGCCGGC--ATACCGN-GTCGGG-CACCTGCATG-TGGNGGGCC  
GGGGCTGCCGGC--GATTCTG-TCCGGG-CACC-TCACG-TGGTGGGGC  
GGGGCTGCCGGC--GATTCTG-TCCGGG-CACA-TCACG-TGGTGGGGC  
GGGGCTGCCGGA-TATACATT-GCCGGG-CACC-GCATG-TGGTGGGGC  
CT-TC-GCCATG--GATG-CGTGG--GAGTG-TCTTTAGTGC-T-GACG  
GGGGCTGCCGGC--GGGACCGT-GCCGGG-CACC-GCAGC-TGGTGGGGC  
GGGGCTGCCAG--TGAATCAGCCGGTACACA-TCACG-TGGTGGGGC  
GGGGCTGCCCGC--GAATCGT-GTCG-G-CACA--CAC--TGGTGGGGC  
GGTCTGCCGGC--CGAACCGCG-CGGGG-CGCA-GCAGC-TGGTGGGGC  
GGGGCTGCCGGC--CGAACCGCG-CGGGG-CGCA-GCAGC-TGGTGGGGC  
GGGGCTGCCGGC--GAATCG-TGTCGGG-CACA-GCAGC-TGGTGGGGC

*Imperatacylindrica*  
*Dichanthiumannulatum*

GGGGCTGCCGGC---GAACCGC-GCCGGG-CACA-GCACG-TGGTGGGCG  
GGGGCTGCTTCCGGCGAATCG-TGTCGGG-CGCA-GCACA-TGGTGGGCG

*Aadscen6713*  
*Aameth6679*  
*Aameth6685*  
*Aappen6543*  
*Aappen6946*  
*Adist6684*  
*Aeuco5869*  
*Arundinellanepalensis*  
*Aschiren6581*  
*Bbladh5173*  
*Binsc5681*  
*BBrachiaria*  
*Bradi4784*  
*Bserra6554*  
*Cdieter6642*  
*Cexc6577*  
*Cleistachnesorghoides*  
*Cmarg4397*  
*Cplur6406*  
*Cprolix6682*  
*Danthoniacalifornica*  
*Dfili5872*  
*Emuti5074*  
*Emuti6715*  
*Eragrostisdielsii*  
*Evill5085*  
*Hanam4807*  
*Hanam4841*  
*Hcon4741*  
*Hdiss5145*  
*Sbicolour9319*  
*Hfilli5144*  
*Hhirt6656*  
*Hlongi6560*  
*Hpilg4603*  
*Hplig4738*  
*Hquar5088*  
*Hrud6576*  
*Htamb6585*  
*Icylin5114*  
*Icylin6632*  
*Mcape6655*  
*Monocymbium*  
*Pdilatum6617*  
*PentaschistisaristifoliaSpies6*  
*Pfrank4705*  
*Pschinz6652*  
*Sbic6588*  
*Sbic6589*  
*Shalep9318*  
*Snigros6567*  
*Tleuc6627*  
*Tspic4846*  
*Ttria6555*  
*Zeadiplo*  
*ZmaysGfl*  
*Saccharumofficinarum*  
*Imperatacylindrica*  
*Dichanthiumannulatum*

--ACAT---TAGTTGTT---CTC---GGTG-CAA-CATCCCGGTAT---A  
??  
AGGCATCA--CGTTGTT---CTCG--ACTGACAAAGCCCCGGGAC---G  
--ACATC---AGTTGTT---CTC---GGTG-CAT-GACCTCGGCAC---G  
??  
CCA-AGG--CAG-ACGTTCTCTC---TGACACACGAACGCC--TCA---GG  
G-ACATC---AGTTGT---CTC---CGTG-CAG-TGACCCGTCAC---A  
--ACAAC---AGTTGTT---CTC---GATG-CAG-TGGCTCGACAC---G  
CCC-ACG---TAACTTCTCTC---AA---CAAAAACCTCC--TCG---TG  
CCA-ATG---AAGATGTTCTCTC---TG-CACAAAAGGGCGG--TCAT---GG  
C-ACATC---GTTGTT---CTC---GGTG-CAG-CGCCCCGGCAT---G  
--ACATC---AGTTGTT---CTC---AGTG-TAA-TGTCTCGCAC---G  
CCACATC---AGATGTT---CTC---TCTGACATACA-CTCCTCAC---G  
--ACACT--CAGGTGTT---CTC---TGTG-TAA-TGCCCTGGCAA---G  
??  
CCA-AGT--CAG-ATGTTCTCTC---TG---CACAGAAGCC--TCAC---GG  
--ACACCT--AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCAC---G  
CCA-AGG--CAGTACGTTCTCTT---TGACACATGAAGGCGTAC---GG  
C-ACATC---AATTGTT---CTCT--CGTACCA--TACCCNTNCGC-G-  
--ACATC---AGATGTT---CTC---TGTG-CAATGCCCCGGCAC---G  
--ACCA---ACGTTGT---CAC---GGTG-CTG-CGCCCCGGCCCC---G  
--ACATC---AATTGT---CTC---GGTG-CAA-TACCCTCAAAA---G  
--ACATC---AAGTTGTT---CTC---GGTG-CAG-CGCTCGGCAC---G  
C-ACATC---AAGTTGTT---CTC---TGTG-CAG-CGACTCGGCAC---G  
--ACAC---ACGGTGT---CTC---GCTG-CTG-CGAACTGGTCCA-AG  
CCACAG---CAG-ACTTTT-CTT---AAT--CCAAAACCC--TCG---GG  
C-ACATC---AGATGTT---CTC---TGTGACAC-TGACCCCTCAC---GG  
??  
G-ACATT--AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCAC---G  
C-ACAACA--TTTGTTCCTCGSCGATAAAA-ATCCCTCCCGCCGCT---  
--ACAC-T-TAATTGTT---CTC---TGTG-CAA-CACCCC-ACAC---G  
CCA-AGA---AAA-ACTTCCCTC---AA---CAAAAAGCCT--TCG---GG  
--ACATT--AGTTGTT---CTC---GGTG-CAT-TGCCTCGGCGC---G  
--ATCT---CGCTATA---CTA---AACG-CAA-TGCATCGGGG---C  
--ACATCA--G-TTGT---CTC---GGTG-CAG-TGCCCGGGCAC---G  
C-ACATT--AGTTGTT---CTC---GGTG-CAT-TGCTTCGGAGC---C  
C-ACGTC--AACTTCTCNTCGCAGAGACTCCATCATGGCTGCCAGT-TA  
CCCACGT--CAA-AC-TTCTCTC---GG---CGACATTGCC--TCT---GC  
CCACATCA--AGATGTT---CTC---TGTGACAGACGACGGCTCGCGG  
--ACATC---AGTTTTT---CTC---GGTG-CA---CACCCGGCGC---G  
--ACATC---ATTGTT---CTC---GTG-CAG-CGCCCCGGCAC---G  
CCA-ATGC--AGATGTT---CTC---GGTG-CACACGAACGCGGTCA-GG  
--ACAT-G-AAGTTGTT---CTC---ACTG-CAA-CATCGCATCAT---G  
??  
G-AGATCT--ACTTGT---GCC---GGTG-CAG-CGCCCCGGCAC---G  
--ACAT-T--AGTTGTT---CTC---GGTG-CAA-TGTCCTGACAC---G  
--ACAC-T-TAATGTT---CTC---TGTG-CAG-CGCCCC-ACAC---G  
--ACAC-T-TAATGTT---CTC---TGTG-CAG-CGCCCCGACAC---G  
--ACATG-TAGGTGTT---CTC---TGTG-CAA-TATCGGCACAT---G  
CCACAGG--CAG-GCGTTCCCTC---AG---CCAGAAGCC--TCGT---G  
--ACTC--AAAGTTGTT---CTCGTTAGTG-CAG--GCCTCGGCAC---G  
--ACATC---AGTTGT---CTC---GGTGACAC-TGCCCGGGCAC---G  
--ACATCT--AATTGTT---CTC---GGTG-CA---CGCCCCGGCAC---G  
--ACATC---AAGTTGTT---CTC---GGTG-CAG-CGTCCCGGGC---G  
--ACATC---AAGTTGTTGTTCTC---GGTG-CAG-CGTCCCGGGC---G  
--ACACTC--AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCAC---G  
--ACATC---AGTTGTT---CTC---GGTG-CAG-CGCCCCGGCGC---G  
--ACATC---AGTTGTT---CTC---GGTG-CAG-CACCCCCGGCACAC-G

*Aadscen6713*  
*Aameth6679*  
*Aameth6685*  
*Aappen6543*  
*Aappen6946*  
*Adist6684*

-AGCTA--TCTTGTGGCCCTAAGG-ACCCATACT-TGACC--GCAGCGC  
??  
CAACTGGCGTTATCGAAACCTAATGACTCATCGAATTA---T-GCAC  
CG-CTAT-CGT-ATTGGCCCTTACG-ACCCATTGA-GCACAGTAAGC-C  
??  
CG-CAACTGTCAT-TCAAA-ATCTCGATCCATCGAGCG---ATTCTGCGC

Aeuco5869  
Arundinellaneapalensis  
Aschiren6581  
Bblad5173  
Binsc5681  
BBrachiaria  
Bradi4784  
Bserra6554  
Cdieter6642  
Cexc6577  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfili5872  
Emuti5074  
Emuti6715  
Eragrostisdielsii  
Evill5085  
Hanam4807  
Hanam4841  
Hcon4741  
Hdiss5145  
Sbicolour9319  
Hfilli5144  
Hhirt6656  
Hlongi6560  
Hpilg4603  
Hplig4738  
Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114  
Icylin6632  
Mcape6655  
Monocymbium  
Pdilatatum6617  
PentaschistisaristifoliaSpies6  
Pfrank4705  
Pschinz6652  
Sbic6588  
Sbic6589  
Shalep9318  
Snigros6567  
Tleuc6627  
Tspic4846  
Ttria6555  
Zeadiplo  
ZmaysGfl  
Saccharumofficinorum  
Imperatacylindrica  
Dichanthiumannulatum

AGGCCG---GCGCATCGCCCTAAGG-ACCCATCGA-GCACC--GGAGCGC  
CGGCCAG--CGTCTTGGCCC-TAAGGACCCATGGA-GCACC--GCAGCGC  
CG-CAACTTGCAT-TCAANN-----  
CG-CAACTTGTATCTGAAA-CTATGAATCCATCGAGCA---ATTATGCAC  
CGGCCG--CGT-ATCGGCCCTAAGG-ACCCATCAA-GCACC--TAAGCGC  
GAACTG--GCTTGTGGCCCTAAA--ACCCAT-GA-CGACC--TTAGCAC  
CGCTAT--TGATTAACCTTACG-ATCCATTACGAACT---CAGCAC  
TACCTGG-CTTG-TCCGCTCTAACG-ACCCATAAA-GCACC--GTAGCCA  
??  
CG-CAACTTGCAT-TCAAC-CTCTCGATCGATCGAGCAC--ATTAGCAC  
CGGCCG--CGCATCGGCC-TAAGGACCCATGGA-GCACC--GCAGCGC  
CGGCAACTTGGCANTGAAA-----  
CGGCCTT---CGCTTGACCTAAAG-ATCCATCGA-GCACC--TAAGCCC  
C-GCCG--TCCGACTGCCCTAACG-ACCCATCGA-GCAAT--TAAGCAC  
CAGCCG--CGAGACGGCCCTCA-GGACCCCTCGC-ATACC--GCAGCGC  
CGCGC-----ATCTGCCCTAAGG-ACCCATCGA-GCACC--TGAGCGC  
CAACCG--CTTATCGGCCCTAAGG-ACCCATTNT-GCACA--CCANCNC  
CAACNG--CTTATCTGCCCTAACG-ACCCATTGT-GCACA--TCAGCGC  
TATCAGGTACGGAATGGCCCTCA-TGACCCATGA---ACC--GAAGCGC  
AC-CTACTTACGT-TCNNA-AACTCGATGGTTCGCGGN---ATTAGCAA  
CAGCCG--TTGCGACTGCCNTAACG-ATCCATCA--GCAAT--TAAGCAA  
??  
CGGACG--CAA-ATCGGCCCTTAGG-ACCCATCGA-GCACC--GCAGCGC  
??  
TGGCCG--GTGCATCTGCCCTAAAG-ACCCTTCNG-GCACT--GCAGCCC  
CC-CAGTT-----  
TGGTCGG-TGT-GTCTGCCCTAAGG-ACCCATCGA-GCACC--TAAGCGC  
GTATCCG-ACGCAATGCCTTCAATG-GACCC--A-TTACC--G-AGCGC  
CGGCCG--CGC-GACGGCCCTAAGG-ACCCATCGA-GCACC--GAAGCGC  
??  
CGCTCG--GT-ATTNGCCCTAAG--ACCCATCGA-GCACC--TAAGCGC  
ANTTCNAANNATAATGGACCCATCNAGCACATTAAGCGCATCACC-----  
TG-CAGCTTGTCTTTCTGCGCAATCGATGGATCTAGAGC--ATTATGCAC  
CGGCCGTTGCGTTCGCCCTATCGGATCCATCGA-GCAAT--TCAGCGC  
C-GCCAG--TGCGTTGGCC--TAA-----  
CGGNCGT--TGATCTGCCCTAAGG-ATCCATGGAAGCACN--TCTTCAC  
CGCAAGTGTGATTTGAAAATAATGGATGGATCGA-GCACAT-TAAGCGC  
TACCTGG-TACTATGGCCTTTACG--ACCGATCAT-CGACC--TCAGCGC  
??  
CTGCCGA-AGCGATTGACCCCTAAGGACCCATCA-CCACC--GAAGCGC  
CAGCTAT-CTCCATAACCTCAAGG-ACCCATTTA-CGACC--GTAGCGC  
TGGCCG--GTGCATCTGCCCTAAG-ACCCTTCGG-GCA-T--GC-GCCC  
TGGCCG--GTGCATCTGCCCTAAG-ACCCTTCGG-GCACT--GC--GCA  
TACCTA--TCTATTGCGCTTAA--ACCCATATG-TGAAC--TAAGCGT  
CG-CAACTTTCGT-TCAAA-AACTCGATGGTTGCGGG---ATTCTGCAA  
CAGCCG--GCGCAGCGCCCTAAGG-ACCCACACG-CGACC--GCAGCGC  
CGGCCG--GCGTAGGCC--TAAGGACCCATCGA-GCACC--GAAGCGC  
CGGCCG-CGT-ATNTGCCCTTATG-ANCCATCTA-GCACC--TAAGCGC  
TGGCCG--CCATTGCGCC--TACGGACCCATCG----AGC--G-ACCGA  
CGGCCG--CCATTGCGCC--TAAGGACCCATCG----AGC--G-ACCGA  
CGGCCGCGC-C-ATCG-----  
CGGCCGCG-TG-TCGGCCCTAAGG-ACCCATCGA-GCACC--GCAGCGC  
--ACCGCG--CATCGGCCCTAAGG-ACCCATCGA-GCACC--GAAGC--

Aadscen6713  
Aameth6679  
Aameth6685  
Aappen6543  
Aappen6946  
Adist6684  
Aeuco5869  
Arundinellaneapalensis  
Aschiren6581  
Bblad5173  
Binsc5681  
BBrachiaria  
Bradi4784  
Bserra6554  
Cdieter6642  
Cexc6577

TTTGC-CGCTC-GGA-CCGCGACCCCAT--GTCAG-CTGG-ACTACCCGC  
??  
TTCATC-CCTC-GGA-TCGCAAT-----  
ATCGCC-GCTC-GGA-TCGCGACCCCAT--GTCATTCTCC-ACTACCCGC  
-----  
TTCACACC-TC-GTA-TCGCAATTGCT--GCCATCTTCAT-CTATCCGA  
ATCCC-GCTC-GGA-CCGCGACCCAG--GTCAGTGGG-ACTACCCGC  
ATCGCC-GCTC-GGA-CC-----  
??  
TTCNCCCTC-GGA-TCGCAATTTCAT--GACATTTGGA-CTATCCGA  
CTCGCC-GCTC-GGA-CCGCGACCCAG--GTCAGTCTGG-ACTACCCGC  
TTTGT-TGCTC-GGA-CCGCGACCCCAT--GTCATACTGG-ACTACCCGC  
ATCAC-CCTCGGAT--CGCAACCCCT--NCCNTNTTCG-ACTACCCGA  
CTTAAGTCTC-GGA-CCGCGACCCCAT--GTCATGCTGG-ACTACCCGC  
??  
TTCACCC-TC-GTA-TCGCAATTGCT--GCCATCTCAA-CTATCCGA

*Cleistachnesorghoides* ACCGCC-GCTC-GGA-CCGCGACCCAG--GTCAGTCGGG-ACT-----  
*Cmarg4397* ???  
*Cplur6406* ATCNCC-NCTC-GNA-TCNCCACCNCAT--GTCAATCTGG-ACTACC--GN  
*Cprolix6682* ATCGCC-CCTC-GGA-CCGCAACCCCAT--GCCATTCTGGTACTACCCGC  
*Danthoniacalifornica* T-CGTC-GCTC-GG-----  
*Dfili5872* ATCACC-CCTC-GGA-CCGC-ACCCCA---GTCATTCTGG-ACTACCCGC  
*Emuti5074* ATTGCC-NCTC-GGA-CCNCAACCCAG--GNNAGCCC-----  
*Emuti6715* ATTTC-CCTC-GGA-TCGCGATCCCT--GTGATTTTGG-ACTATACGC  
*Eragrostisdielsii* A-TCTC-GCTC-GGA-CC-----  
*Evill5085* TTCCTGCGAGG-GGA-TCGCATTTTCCATANCCTTCTTACATACTATNCCA  
*Hanam4807* NATCNCCCTC-GGA--CGCAANCCCT--GNCATNTTGN-ACTANNCGN  
*Hanam4841* ???  
*Hcon4741* ATCGCC-GCTC-GGA-CCGCGACCCAG--GTCAGTCGGG-ACTACCCGC  
*Hdiss5145* ???  
*Sbicolour9319* CATCCATGCTC-GGA-CCGCGACCCAN--GTCAGTCNGG-ACCACCCGC  
*Hfili5144* ???  
*Hhirt6656* ATCC--GCTC-GGA-CCACGACCCAG--GTCAGTCTGG-ACTACCCGC  
*Hlongi6560* ACA---CCTT-CTA-CCGC-ACCCCAT--GTCA-GCTGG-ACTACCCGC  
*Hpilg4603* ATCGCC-GCTC-GGA-CCGCGACCCAG--GTCAGTCGGG-ACTACCCGC  
*Hplig4738* ???  
*Hquar5088* ATCC--GCTC-GGA-CCGCGACCCAG--GTCATTCTGG-ACTACCCGC  
*Hrud6576* ???  
*Htamb6585* TTCACCGCTCG-GTA-TCGCATCTTCT--GCTATCTTCTG-CGACCCGA  
*Icylin5114* ATCGCC-GCTC-GGA-TCACGATCCCAT--GTCATCCTGG-ACTATCCGC  
*Icylin6632* ???  
*Mcape6655* ATCACC-GCTC-GGATCCGA--CCCAT--GTCATTCTGG-ACTACCCGC  
*Monocymbium* TTCGCC-GCTC-GGA-TCGCAATTTTCAT--GTCATTTTGGTACTATCCGC  
*Pdilatum6617* TTGTA-CACTC-GGA-CCGC-ACCCCAT--GTGATGCTGG-ATTACCCGC  
*PentastichisaristifoliaSpies6* ???  
*Pfrank4705* ATCGCC-GCTC-GGA-CCGCGACCCAG--GTCAGTCGGG-ACTACCCGC  
*Pschinz6652* CTTGG-TGCTC-GGA-CCGCGACCCCAA--GTCATGCGGG-ACTACCCGC  
*Sbic6588* ANTCCATGCTC-GGA-CCGCGACCCAN--GTCANNNGG-ACCACCCGC  
*Sbic6589* CATCGATGCTC-GGA-CCGCGACCCAG--GTCAGTCTGG-ACCACCCGC  
*Shalep9318* TTCCT-CGCTC-GGA-CCGC-ACCCCAT--GTCATACCTGG-ACTACCCGC  
*Snigros6567* TTCACACCAT--GTA-TCGCATTTTGCT--ACGTTCTTTCAT-ACATGCCA  
*Tleuc6627* CAGCC--GCTC-GGA-CCGCGACCCAG--GTCAGACGGG-ACTACCCGC  
*Tspic4846* ATCGCC-GCTC-GGA-CCGCGACCCAG--GTCAGTCGGG-ACTACCCGC  
*Ttria6555* TTTTCC-GCTTTGGA-ACGCGACCC-AN--GTCA-TCTGG-ACTACCCGC  
*Zeadiplo* GCTTGT-CCTC-GGA-CCGCGA-----  
*ZmaysGfl* GCTTGC-CCTC-GGA-CCGCGA-----  
*Saccharumofficinatum* -----  
*Imperatacylindrica* ATCGCC-GCTC-GGA-CC-----  
*Dichanthiumannulatum* ATCACC-GCTC-GGA-CCGCGA-----

*Aadscen6713* GAA-TCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Aameth6679* ???  
*Aameth6685* ???  
*Aappen6543* ----TCGG-TATA-GGTTTTGTGG-----TTTGGGGG-----  
*Aappen6946* GAA-TGGA-AACCTTGCTAGGTTCTAATCCCN--AAATCCAGAGCCCC  
*Adist6684* ???  
*Aeuco5869* ???  
*Arundinellanepalensis* ???  
*Aschiren6581* ???  
*Bbladh* GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Binsc5681* ???  
*BBrachiararia* GAA-TCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Bradi4784* ???  
*Bserra5074* GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Bserra6654* GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Bserra6954* GAA-TCGG-TAGA-CGATACGTT-AATG--GGT--ATATCCCAGCCTGAA  
*Cdieter6642* ???  
*Cexc6577* -----  
*Cleistachnesorghoides* ???  
*Cmarg4397* GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Cplur6406* ???  
*Cprolix6682* GAA-TCGG-ACC--GCTACGTTGGAAGTCCCA--ATTTCAGAGAAACCC  
*Danthoniacalifornica* ???  
*Dfili5872* GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Eragrostisdielsii* ???  
*Evill5085* ???

*Hanam4841* GAA-TCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Hcon6942* ???  
*Hdiss5145* GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Hemarthria9319* ???  
*Hfilli5144* ???  
*Hhirt6550* GAA-TCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCACCAAACC  
*Hhirt6656* ???  
*Hlongi6560* GAA-TCGG-TAGA-CGCTACGTT-AAGT--CCC--TCCTCCCAGCCGAAA  
*Hpilg4603* GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Hpilg4738* GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Hquar5088* ???  
*Hrud6576* ???  
*Htamb6585* ???  
*Icylin5114* GAAATCGG-TAGA-CGCTACGTT-CAATTCCT--CCTTCCCAGCCGAAA  
*Icylin6632* GAA-TCGG-TAGA-CGCTACGTT--TATTCCT--CCTTCCCAGAAGAAC  
*Mcape6655* GTA-TGGG-AAC--TGCTAAGTG-TAACTCCA--ATTCCCAGAAC--  
*Monocymbium* ???  
*Pdilatum6617* GAA-TCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Pentaschistis aristifol* ???  
*Pfrank4705* GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Pschinz6652* GAAATCGG-TAGA-CGCTACGTT-CAAGTCCCT--CCTTCCCAGCCGAAA  
*Sbic6588* ???  
*Sbic6589* GAA-T-GG-T-GA-CGCTACGTT--ACTCCCT--CCTTCCCAGCCGAAA  
*Shalep9318* GAA-TCGG-AACC-CGCTACGTT-AAGT--CCC--ATTC--AGC-GAAA  
*Snigros6567* GAAATCGG-TAGA-CGCTACGTT-TAATTCCT--CATTCAGCCGAAA  
*Tleuco6627* ???  
*Tspic4846* GAAATCGG-TAGA-CGCTACGTT-CAATTCCT--CCTTCCCAGCCGAAA  
*Ttria6555* GAAAT-GG-TAG--CGGTAGCGTTCAAGTCCCGACATAAGCTGCCACT  
*Zeadiplo* ???  
*Zea* TTCT-AT-CTA--AGTGAACCTCAATTT-AGAAGTAGT-TAATAACT  
*Sacharum officinarum* -----TTCCA--AATTCAGAGAA-ACC  
*Icylindrical* -----TTCCA--AATTCAGAGAA-ACC  
*Dichanthium annulatum* ???

*Aadscen6713* AAACAA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Aameth6679* ???  
*Aameth6685* ???  
*Aappen6543* AAAGAAAGCGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Aappen6946* AAACAA-GCGGTTCT--CAAAC TAGAA-CCCAAAGGAAA-GGATAGGTGC  
*Adist6684* ???  
*Aeuco5869* ???  
*Arundinellane palensis* ???  
*Aschiner6581* ???  
*Bblad* AAACAA-GCGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Binsc5681* ???  
*BBrachiaria* AAACAA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Bradi4784* ???  
*Bserra5074* AAAC TT-GCGG-TCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Bserra6654* AAACAA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Bserra6954* AAACAA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Cdieter6642* ???  
*Cexc6577* -----  
*Cleistachnesorghoides* ???  
*Cmarg4397* AAAC TT-GCGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Cplur6406* ???  
*Cprolix6682* AAACAA-GC-GGTT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Danthoniacalifornica* ???  
*Dfilli5872* AAAC TA-GCGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Eragrostisdielsii* ???  
*Evill5085* ???  
*Hanam4841* --ACCA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Hcon6942* ???  
*Hdiss5145* AAACAA-GCGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Hemarthria9319* ???  
*Hfilli5144* ???  
*Hhirt6550* AAACAA-GCG-----CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Hhirt6656* ???  
*Hlongi6560* AAACAACGAGGTTCT--CGAAC TAGAA-TACAAAGGAAAAGGATAGGTGC  
*Hpilg4603* AAACCA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAAATGC  
*Hpilg4738* AAAC TA-GCGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC

*Hquar5088* ???  
*Hrud6576* ???  
*Htamb6585* ???  
*Icylin5114* AAACAA-GCGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Icylin6632* AAACAA-GCGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Mcape6655* TGACTA-CATGGTCTACCTAATGATGA-TCCAAAGGTAATGGATAGGTGC  
*Monocymbium* ???  
*Pdilatum6617* AA-CAA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Pentaschistisaristifol* ???  
*Pfrank4705* AAACAA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Pschinz6652* AAACAA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Sbic6588* ???  
*Sbic6589* AAA---GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Shalep9318* AA-CAA-GTGGTT-T--CCAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Snigros6657* AAACAA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Tleuco6627* ???  
*Tspic4846* AAACAA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Ttria6555* GGACAAAGA---TC---CATATTGGTC-----GGTCCCGTAGCG-TCT  
*Zeadiplo* ???  
*Zea* TGTACATT---TC---TGTTATGTAA-----GCCCACTTAG---CTC  
*Sacharumofficinatum* AAACAA-GTGGTTCT--CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Icylindrical* AAACAA-GTG-----CAAAC TAGAA-CCCAAAGGAAAAGGATAGGTGC  
*Dichanthiumannulatum* ???

*Aadscen6713* AGAGACTCAATGGAAGCTATTCTA--ACGAATCGAAGTAATTACGTTGTG  
*Aameth6679* ???  
*Aameth6685* ???  
*Aappen6543* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Aappen6946* AGAGACTCTATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Adist6684* ???  
*Aeuco5869* ???  
*Arundinellanepalensis* ???  
*Aschiren6581* ???  
*Bbladh* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Binsc5681* ???  
*BBrachiaria* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAATTACGTTGTG  
*Bradi4784* ???  
*Bserra5074* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Bserra6654* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAATTACGTTGTG  
*Bserra6954* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Cdieter6642* ???  
*Cexc6577* -----  
*Cleistachnesorghoides* ???  
*Cmarg4397* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Cplur6406* ???  
*Cprolix6682* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Danthoniacalifornica* ???  
*Dfilli5872* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Eragrostisdielsii* ???  
*Evill5085* ???  
*Hanam4841* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Hcon6942* ???  
*Hdiss5145* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Hemarthria9319* ???  
*Hfilli5144* ???  
*Hhirt6550* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Hhirt6656* ???  
*Hlongi6560* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAGTTAATTACGTTGTG  
*Hpilg4603* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Hpilg4738* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Hquar5088* ???  
*Hrud6576* ???  
*Htamb6585* ???  
*Icylin5114* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Icylin6632* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----  
*Mcape6655* AGAGACTCCCATG--GCTGTTCTAGAATGGATCTGAGCGAT-----  
*Monocymbium* ???  
*Pdilatum6617* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAATTGCGTTGTG  
*Pentaschistisaristifol* ???  
*Pfrank4705* AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----

<i>Pschinz6652</i>	AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAATTACGTTGTG
<i>Sbic6588</i>	??
<i>Sbic6589</i>	AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----
<i>Shalep9318</i>	AGAGACTCAATGGA-GCTGTTCTA--ACGAATCGAAGTAAT-ACGTTGTG
<i>Snigros6567</i>	AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAATTACGTTGTG
<i>Tleuco6627</i>	??
<i>Tspic4846</i>	AGAGACTCAATGGAAGCTTCTA--ACGAATCGAAGTAAT-----
<i>Ttria6555</i>	ACCGATTTTCGCGAAGCTGTCCTA--CCGGTTCGAAGTATT-----
<i>Zeadiplo</i>	??
<i>Zea</i>	AGAGGTTAGAGCATCGCATTTGTA-ACGAATCGGTTCAAAT-----
<i>Sacharumofficinatum</i>	AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----
<i>Icylindrical</i>	AGAGACTCAATGGAAGCTGTTCTA--ACGAATCGAAGTAAT-----
<i>Dichanthiumannulatum</i>	??
<i>Aadscen6713</i>	TTGGTAGTGGAACTCCCTCGAAATATAGAAAGAAGGGCTTTATA-----
<i>Aameth6679</i>	??
<i>Aameth6685</i>	??
<i>Aappen6543</i>	-----
<i>Aappen6946</i>	-----
<i>Adist6684</i>	??
<i>Aeuco5869</i>	??
<i>Arundinellaneapalensis</i>	??
<i>Aschieren6581</i>	??
<i>Bbladh</i>	-----
<i>Binsc5681</i>	??
<i>BBrachiaria</i>	TTGGTAGGGGAACCTCGAAATCTAGAAAGAAGGGCTTTATATATA
<i>Bradi4784</i>	??
<i>Bserra5074</i>	-----
<i>Bserra6654</i>	TTGGTAGTGGAACTCCCTCGAAATATAGAAAGAAGGGCTTTATTT-ATA
<i>Bserra6954</i>	-----
<i>Cdieter6642</i>	??
<i>Cexc6577</i>	-----
<i>Cleistachnesorghoides</i>	??
<i>Cmarg4397</i>	-----
<i>Cplur6406</i>	??
<i>Cprolix6682</i>	-----
<i>Danthoniacalifornica</i>	??
<i>Dfilli5872</i>	-----
<i>Eragrostisdielsii</i>	??
<i>Evill5085</i>	??
<i>Hanam4841</i>	-----
<i>Hcon6942</i>	??
<i>Hdiss5145</i>	-----
<i>Hemarthria9319</i>	??
<i>Hfilli5144</i>	??
<i>Hhirt6550</i>	-----
<i>Hhirt6656</i>	??
<i>Hlongi6560</i>	TTGGTAGTGGAAATTCCTCTAAATTTGAGAAAGAAGGGCTTTATA-----
<i>Hpilg4603</i>	-----
<i>Hpilg4738</i>	-----
<i>Hquar5088</i>	??
<i>Hrud6576</i>	??
<i>Htamb6585</i>	??
<i>Icylin5114</i>	-----
<i>Icylin6632</i>	-----
<i>Mcape6655</i>	-----
<i>Monocymbium</i>	??
<i>Pdilatum6617</i>	TTGGTAGTGGAACTCTTCGAAATTAGAGAAAGAAGGGCTTTATA-----
<i>Pentaschistisarisistifol</i>	??
<i>Pfrank4705</i>	-----
<i>Pschinz6652</i>	TTGGTAGTGGAACTCCCTCGAAATATAGAAAGAAGGGCTTTATA-----
<i>Sbic6588</i>	??
<i>Sbic6589</i>	-----
<i>Shalep9318</i>	TTGGTAGTGGAACTCCCTCGAAATACTAGAAAGAAGGGCTTTATA-----
<i>Snigros6567</i>	TTGGTAGTGGAACTCCCTCGAAATACTAGAAAGAAGGGCTTTATA-----
<i>Tleuco6627</i>	??
<i>Tspic4846</i>	-----
<i>Ttria6555</i>	-----
<i>Zeadiplo</i>	??
<i>Zea</i>	-----



Sacharumofficinarum -----  
Icylindrical -----  
Dichanthiumannulatum ???

Aadscen6713 CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--  
Aameth6679 ???  
Aameth6685 ???  
Aappen6543 -----AACGATTAATCACA--  
Aappen6946 -----AACGATCAATCACA--  
Adist6684 ???  
Aeuco5869 ???  
Arundinellaneapalensis ???  
Aschiren6581 ???  
Bbladh -----AACGATTAATCACA--  
Binsc5681 ???  
BBrachiaria CATTTAATACACACGTATAGATACTGACATAGCAAACGATTAATCAGC--  
Bradi4784 ???  
Bserra5074 -----AACGATTAATCACA--  
Bserra6654 CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--  
Bserra6954 -----AACGATTAATCACA--  
Cdieter6642 ???  
Cexc6577 -----  
Cleistachnesorghoides ???  
Cmarg4397 -----AACGATTAATCACAAA  
Cplur6406 ???  
Cprolix6682 -----AACGATTAATCACA--  
Danthoniacalifornica ???  
Dfilli5872 -----AACGATTAATCACA--  
Eragrostisdielsii ???  
Evill5085 ???  
Hanam4841 -----AACGATTAATCACA--  
Hcon6942 ???  
Hdiss5145 -----AACGATTAATCACA--  
Hemarthria9319 ???  
Hfilli5144 ???  
Hhirt6550 -----AACGATTAATCACA--  
Hhirt6656 ???  
Hlongi6560 CACCTAATAAACACGTATAGATACTGGCATAGCAAACGATTAATCAGC--  
Hpilg4603 -----AACGATTAATCACA--  
Hpilg4738 -----AACGATTAATCACA--  
Hquar5088 ???  
Hrud6576 ???  
Htamb6585 ???  
Icylin5114 -----AACGATTAATCACA--  
Icylin6632 -----AACGATTATTCACA--  
Mcape6655 -----GCAGCTTAATCACA--  
Monocymbium ???  
Pdilatatum6617 CATCTAATATACACGCATATATACTGACATAGCAAACGATTAATCAGC--  
Pentaschistisaristifol ???  
Pfrank4705 -----AACGATTAATCACA--  
Pschinz6652 CATCTAATACACACGTATAGATACTGACATAGCAAACGATTAATCATA--  
Sbic6588 ???  
Sbic6589 -----AACGATTAATCACA--  
Shalep9318 CATTTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--  
Snigros6567 CATTTAATACACACGTATAGATACTGACATAGCAAACGATTAATCACA--  
Tleuco6627 ???  
Tspic4846 -----TACGATTAATCACA--  
Ttria6555 -----ACGAGT-TTCAGG--  
Zeadiplo ???  
Zea -----CCGATT-TTTTTT--  
Sacharumofficinarum -----AACGATTAATCACA--  
Icylindrical -----AACGATTAATCACA--  
Dichanthiumannulatum ???

Aadscen6713 GAACCCATATCATAATATAGGTTCTTTATTTTATAT-----TTTTTTT  
Aameth6679 ???  
Aameth6685 ???  
Aappen6543 GAACCCATATTATAATATAGGTTCTTTATTTGATTT-----TT  
Aappen6946 GAACCCATATTATAATATAGGTTCTTTATTTGATTT-----TT



Bserra6654 AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTATG---A  
 Bserra6954 AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCAT---A  
 Cdieter6642 ???  
 Cexc6577 ---CTGAAACGCTCCCAAGAACTCTCGCTATCGCCAGATAAATTCGGTGGGA  
 Cleistachnesorghoides ???  
 Cmarg4397 AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCAT---A  
 Cplur6406 ???  
 Cprolix6682 AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCAT---A  
 Danthoniacalifornica ???  
 Dfilli5872 AGAATGAAATT-TGG-AAGGATTATGAA-----ATAGAAAATTCAT---A  
 Eragrostisdielsii ???  
 Evill5085 ???  
 Hanam4841 AGAATGAAATT-TGG-AAGGATTATGAA-----ATAGAAAATTCAT---A  
 Hcon6942 ???  
 Hdiss5145 AGAATGAAATT-TGG-AAGGATTATGAAGAGGTATAGAAAATTCAT---A  
 Hemarthria9319 ???  
 Hfilli5144 ???  
 Hhirt6550 AGAATGAAATT-TGG-AAGGATTATGAAGAGGTATAGAAAATTCAT---A  
 Hhirt6656 ???  
 Hlongi6560 AAATTTGAAAT-AGA-AATGATTATGAA-----ATAATAAATTCCTG---A  
 Hpilg4603 AGAATGAAATT-TGG-AATGATTATGAA-----ATAGAAAATTCAT---A  
 Hpilg4738 AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCAT---A  
 Hquar5088 ???  
 Hrud6576 ???  
 Htamb6585 ???  
 Icylin5114 AGAATGAAATT-AG--AATGATTATGAA-----ATAGAAAATTCAT---A  
 Icylin6632 AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCAT---A  
 Mcape6655 TGAATGAAATT-CGG-AATGATTATGAA-----ATACAACAACAGC---A  
 Monocymbium ???  
 Pdilatum6617 -GAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCCTG---A  
 Pentaschistisaristifol ???  
 Pfrank4705 AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCAT---A  
 Pschinz6652 AGAATGAAATT-AGG-AATGATTATGAA-----ATATAAATTCCTG---A  
 Sbic6588 ???  
 Sbic6589 AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCAT---A  
 Shalep9318 AGAATGAAATT-AGG-AATGATTAGGAA-----ATAGAAAATTCCTG---A  
 Snigros6567 AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCCTG---A  
 Tleuco6627 ???  
 Tspic4846 AGAATGAAATT-TGG-AAGGATTATGAA-----ATACAAAATTCAT---A  
 Ttria6555 AAATTTGAAATT-AGG-AATGATTATGAA-----ATAAAA-ATTCAT---A  
 Zeadiplo ???  
 Zea CGAATAAAATG-GGG-AATCAGGGTCTA-----TTAGTC-GTTCAA---G  
 Sacharumofficinatum AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCAT---A  
 Icyllindrical AGAATGAAATT-AGG-AATGATTATGAA-----ATAGAAAATTCCT---A  
 Dichanthiumannulatum ???

Aadscen6713 ATTTTTT-GAGAATTA----TTGT-GAATCC-ATTCCAA-TCGA-ATATT  
 Aameth6679 ???  
 Aameth6685 ???  
 Aappen6543 ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
 Aapen6946 ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
 Adist6684 ???  
 Aeuco5869 ???  
 Arundinellaneapalensis ???  
 Aschiren6581 ???  
 Bbladh ATTTTTT-GAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
 Binsc5681 ???  
 BBrachiararia ATTTTTT-TAGAATTA----TTGT-GAATCC-ATTCCAA-TCGA-ATATT  
 Bradi4784 ???  
 Bserra5074 ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
 Bserra6654 ATTTTTT-CATAATTA----TTGT-GAATC--ATTCCAA-TCGA-ATATT  
 Bserra6954 ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
 Cdieter6642 ???  
 Cexc6577 CTATGATCGTCTGACTA-TTTCGCGCTGTCT-ACGCCAA-CAGAGTCATT  
 Cleistachnesorghoides ???  
 Cmarg4397 ATTTTTT-GAGAATTAATTTATGTT-GAATCT-ATTCCAA-TCGA-ATATT  
 Cplur6406 ???  
 Cprolix6682 ATTTTTT-GAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
 Danthoniacalifornica ???  
 Dfilli5872 ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT

*Eragrostisdielsii* ???  
*Evill5085* ???  
*Hanam4841* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Hcon6942* ???  
*Hdiss5145* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Hemarthria9319* ???  
*Hfilli5144* ???  
*Hhirt6550* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Hhirt6656* ???  
*Hlongi6560* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Hpilg4603* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Hpilg4738* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Hquar5088* ???  
*Hrud6576* ???  
*Htamb6585* ???  
*Icylin5114* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Icylin6632* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Mccape6655* TTAAATG-AAGAAATC----ATGT-GAAGCT-ATTGCTT-GCAATATATT  
*Monocymbium* ???  
*Pdilatum6617* ATTTTTT-TAGAATTA----TTGT-GAATCC-ATTCCAA-TCGA-ATATT  
*Pentaschistisaristifol* ???  
*Pfrank4705* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Pschinz6652* ATTTTTT-TAGAATTA----TTGT-GAATCC-ATTCCAA-TCGA-ATATT  
*Sbic6588* ???  
*Sbic6589* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Shalep9318* ATTTTTT-TAGAATTA----TTGT-GAATCC-ATTCCAA-TCGA-ATATT  
*Snigros6567* ATTTTTT-TAGAATTA----TTGT-GAATCC-ATTCCAA-TCGA-ATATT  
*Tleuco6627* ???  
*Tspic4846* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Ttria6555* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Zeadiplo* ???  
*Zea* ACTCCAT-TTGTCATA----TATT-CCATAT-ATCACAT-TCGA-AGATA  
*Sacharumofficinatum* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Icylindrical* ATTTTTT-TAGAATTA----TTGT-GAATCT-ATTCCAA-TCGA-ATATT  
*Dichanthiumannulatum* ???

*Aadscen6713* -TCATAATCA-AATCCTT--CAATTCATTGTTTTG-----AGATCTTTA  
*Aameth6679* ???  
*Aameth6685* ???  
*Aappen6543* -GAGTAATCA-AATCCTT--CAA-TCATTGTTTTCG-----AGATCTTTT  
*Aappen6946* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTTC-----GAGATCTTTT  
*Adist6684* ???  
*Aeuco5869* ???  
*Arundinellanepalensis* ???  
*Aschiren6581* ???  
*Bbladh* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
*Binsc5681* ???  
*BBrachiararia* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTG-----AGATC-----  
*Bradi4784* ???  
*Bserra5074* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
*Bserra6654* -GAGTAATCA-AATCCTT--CAACTCATT---TTG-----AGATCTTTA  
*Bserra6954* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
*Cdieter6642* ???  
*Cexc6577* AGGGGAGACGGATCACGCACATATATATATTTCATTTCA----ATATC-TAT  
*Cleistachnesorghoides* ???  
*Cmarg4397* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATC-TTT  
*Cplur6406* ???  
*Cprolix6682* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTTC-----GAGATCTTTT  
*Danthoniacalifornica* ???  
*Dfilli5872* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
*Eragrostisdielsii* ???  
*Evill5085* ???  
*Hanam4841* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
*Hcon6942* ???  
*Hdiss5145* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
*Hemarthria9319* ???  
*Hfilli5144* ???  
*Hhirt6550* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTCG-----AGATCTTTT  
*Hhirt6656* ???  
*Hlongi6560* -TAGTAATCA-AATCCTT--CAATTCAGT---TTG-----GAGTCTTTA

*Hpilg4603* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Hpilg4738* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Hquar5088* ???  
*Hrud6576* ???  
*Htamb6585* ???  
*Icylin5114* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Icylin6632* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Mcape6655* -GACTAATCACAATGGAC--TAATACATTGTGTT-----GAGATCATT  
*Monocymbium* ???  
*Pdilatum6617* -GAGTAATCA-AATCCTT--CAATTCATTATTTTGTTTTTCGAGATCTTAA  
*Pentaschistis aristifol* ???  
*Pfrank4705* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Pschinz6652* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTCA  
*Sbic6588* ???  
*Sbic6589* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Shalep9318* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Snigros6567* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Tleuco6627* ???  
*Tspic4846* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Ttria6555* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Zeadiplo* ???  
*Zea* -TCATATTCA-TGGAATA--CGATTCACCT--TTCA-----AGATGCCCT  
*Sacharum officinarum* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Icylindrical* -GAGTAATCA-AATCCTT--CAATTCATTGTTTTTCG-----AGATCTTTT  
*Dichanthium annulatum* ???

*Aadscen6713* AAAAGTGG-ATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Aameth6679* ???  
*Aameth6685* ???  
*Aappen6543* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Aappen6946* CACTGAGTGGATTAATC--GGACGAGGATA--AAGAGA--GACTAACA  
*Adist6684* ???  
*Aeuco5869* ???  
*Arundinellane palensis* ???  
*Aschirene6581* ???  
*Bbladh* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Binsc5681* ???  
*BBrachiaria* AAAAAGTGG-ATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Bradi4784* ???  
*Bserra5074* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Bserra6654* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Bserra6954* TAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Cdieter6642* ???  
*Cexc6577* ATAAGATGTGATAGTC--GAAAGTATACTGG-AAAAGAATGACGTACC  
*Cleistachnesorghoides* ???  
*Cmarg4397* TAAAAGCGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Cplur6406* ???  
*Cprolix6682* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Danthoniacalifornica* ???  
*Dfilli5872* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Eragrostisdielsii* ???  
*Evill5085* ???  
*Hanam4841* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hcon6942* ???  
*Hdiss5145* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hemarthria9319* ???  
*Hfilli5144* ???  
*Hhirt6550* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hhirt6656* ???  
*Hlongi6560* AAACAGGTGATTAATCCGGGACGAGGACA--AAGAGA--GAGTCCCA  
*Hpilg4603* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hpilg4738* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Hquar5088* ???  
*Hrud6576* ???  
*Htamb6585* ???  
*Icylin5114* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Icylin6632* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA  
*Mcape6655* AAAAAGTGGATAAATC--GGCGGAGGATT--AAGCGT--GGTTCATA  
*Monocymbium* ???  
*Pdilatum6617* AAAAAGTGGATTAATC--GGACGAGGATA--AAGAGA--GAGTCCCA

Pentaschistis aristifol ???? ???? ???? ???? ???? ???? ???? ???? ???? ???? ???? ???? ???? ???? ???? ????  
Pfrank4705 AAAAAGTGGATTAATC--GGACGAGGATA---AAGAGA--GAGTCCCA  
Pschinz6652 AAAAAGTGGATTAATC--GGACGAGGATA---AAGAGA--GAGTCCCA  
Sbic6588 ???  
Sbic6589 AAAAAGTGGATTAATC--GGACGAGGATA---AAGAGA--GAGTCCCA  
Shalep9318 GAAAGTGG--TATACTC---GAATCGGACG---AGGATAA--GAGTCCCA  
Snigros6567 AAAAAGTGG--ATTAATC---GACGAGGATA---AAGAGA--GAGTCCCA  
Tleuco6627 ???  
Tspic4846 AAAAAGTGGATTAATC--GGACGAGGATA---AAGAGA--GAGTCCCA  
Ttria6555 AAAAAGTGGATTAATC--GGACGAGGATA---AAGAGA--GAGTCCCA  
Zeadiplo ???  
Zea GA---TGGTGAAT---GGTAGAC-----ACGCGA--GACTCAAA  
Sacharum officinarum AAAAAGTGGATTAATC--GGACGAGGATA---AAGAGA--GAGTCCCA  
Icylindrical AAAAAGTGGATTAATC--GGACGAGGATA---AAGAGA--GAGTCCCA  
Dichanthium annulatum ???

Aadscen6713 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Aameth6679 ???  
Aameth6685 ???  
Aappen6543 TTCTA--CATGTCAAT--ACTGA--CAACATT--AATTTCTAGTAAAA  
Aappen6946 TTTTG--AATCCCAAC--GGAAT--CGGG--ACACATAACTTTCAAAAT  
Adist6684 ???  
Aeuco5869 ???  
Arundinellane palensis ???  
Aschiren6581 ???  
Bbladh TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Bnsc5681 ???  
BBrachiaria TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Bradi4784 ???  
Bserra5074 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Bserra6654 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Bserra6954 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Cdieter6642 ???  
Cexc6577 GTCAACGANTTTCNGG---NNN--TATAGTCTTTATTATTTTANANA  
Cleistachnesorghoides ???  
Cmarg4397 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Cplur6406 ???  
Cprolix6682 TTCTA--CATGTCAAT--ACTGA--CGGC--AGATGGATTGAATCCC  
Danthoniacalifornica ???  
Dfilli5872 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Eragrostisdielsii ???  
Evill5085 ???  
Hanam4841 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Hcon6942 ???  
Hdiss5145 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Hemarthria9319 ???  
Hfilli5144 ???  
Hhirt6550 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Hhirt6656 ???  
Hlongi6560 TTCTA--CATGTCAAT--ACTG---AACAATGAAATTTCTAGTAAAA  
Hpilg4603 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Hpilg4738 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Hquar5088 ???  
Hrud6576 ???  
Htamb6585 ???  
Icylin5114 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Icylin6632 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Mcape6655 TTCGG--TATGTCAAT--GCGAT--CGTC--TGACTTTACTTTCTAAC  
Monocymbium ???  
Pdilatatum6617 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Pentaschistis aristifol ???  
Pfrank4705 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Pschinz6652 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Sbic6588 ???  
Sbic6589 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Shalep9318 TTCTA--CATGTCAAT--ACTGA--CAACAATGAA--TTTCTAGTAAAA  
Snigros6567 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Tleuco6627 ???  
Tspic4846 TTCTA--CATGTCAAT--ACTGA--CAACAATGAAATTTCTAGTAAAA  
Ttria6555 TCCTA--CATGTCAAT--ACTGGACAACAATGAAATTTCTAGTAAAA

<i>Zeadiplo</i>	??
<i>Zea</i>	ATCT---CGTG-CTAA--AGAGCGTGGAGGTTCGAGTCCT-CTTCAA
<i>Sacharumofficinarum</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Icylindrical</i>	TTCTA--CATGTCAAT--ACTGA-CAACAATGAAATTTCTAGTAAAA
<i>Dichanthiumannulatum</i>	??

<i>Aadscen6713</i>	GGA
<i>Aameth6679</i>	???
<i>Aameth6685</i>	???
<i>Aappen6543</i>	GGA
<i>Aappen6946</i>	TCC
<i>Adist6684</i>	???
<i>Aeuco5869</i>	???
<i>Arundinellanepalensis</i>	???
<i>Aschiren6581</i>	???
<i>Bbladh</i>	GGA
<i>Binsc5681</i>	???
<i>BBrachiaria</i>	GGA
<i>Bradi4784</i>	???
<i>Bserra5074</i>	GGA
<i>Bserra6654</i>	GGA
<i>Bserra6954</i>	GGA
<i>Cdieter6642</i>	???
<i>Cexc6577</i>	ATG
<i>Cleistachnesorghoides</i>	???
<i>Cmarg4397</i>	GGA
<i>Cplur6406</i>	???
<i>Cprolix6682</i>	TCC
<i>Danthoniacalifornica</i>	???
<i>Dfilli5872</i>	GGA
<i>Eragrostisdielsii</i>	???
<i>Evill5085</i>	???
<i>Hanam4841</i>	GGA
<i>Hcon6942</i>	???
<i>Hdiss5145</i>	GGA
<i>Hemarthria9319</i>	???
<i>Hfilli5144</i>	???
<i>Hhirt6550</i>	GGA
<i>Hhirt6656</i>	???
<i>Hlongi6560</i>	GGA
<i>Hpilg4603</i>	GGA
<i>Hpilg4738</i>	GGA
<i>Hquar5088</i>	???
<i>Hrud6576</i>	???
<i>Htamb6585</i>	???
<i>Icylin5114</i>	GGA
<i>Icylin6632</i>	GGG
<i>Mcape6655</i>	ACC
<i>Monocymbium</i>	???
<i>Pdilatum6617</i>	GGA
<i>Pentaschistisaristifol</i>	???
<i>Pfrank4705</i>	GGA
<i>Pschinz6652</i>	GGA
<i>Sbic6588</i>	???
<i>Sbic6589</i>	GG-
<i>Shalep9318</i>	GGA
<i>Snigros6567</i>	GGA
<i>Tleuco6627</i>	???
<i>Tspic4846</i>	GGA
<i>Ttria6555</i>	GGG
<i>Zeadiplo</i>	???
<i>Zea</i>	GGC
<i>Sacharumofficinarum</i>	GGA
<i>Icylindrical</i>	GGA
<i>Dichanthiumannulatum</i>	???

<i>Aadscen6713</i>	?????
<i>Aameth6679</i>	GACCC
<i>Aameth6685</i>	GACCC

Aappen6543 ?????  
 Aappen6946 GATCC  
 Adist6684 GACCC  
 Aeuco5869 GACCC  
 Arundinellaneapalensis ?????  
 Aschiren6581 GACCC  
 Bbladh GACCC  
 Binsc5681 GACCC  
 BBrachiaria ?????  
 Bradi4784 ?????  
 Bserra5074 GACTC  
 Bserra6654 ?????  
 Bserra6954 GACTC  
 Cdieter6642 GACCC  
 Cexc6577 GACCC  
 Cleistachnesorghoides ?????  
 Cmarg4397 ?????  
 Cplur6406 GACCC  
 Cprolix6682 GACCC  
 Danthoniacalifornica ?????  
 Dfilli5872 CCTCC  
 Eragrostisdielsii ?????  
 Evill5085 ?????  
 Hanam4807 GACCC  
 Hcon6942 GACCA  
 Hdiss5145a ACCCC  
 Hemarthria9319 GACCC  
 Hfilli5144 GCCCC  
 Hhirt6650 ?????  
 Hhirt6656f GAACC  
 Hlongi6560 ?????  
 Hpilg4603 GACCC  
 Hpilg4738 GACCC  
 Hquar5088 GACCA  
 Hrud6576 GACCA  
 Htamb6585 GACCC  
 Icylin5114a GACCC  
 Icylin6632a GACCA  
 Mcape6655 GACCC  
 Monocymbium ?????  
 Pdilatatum6617 GACCC  
 Pentaschistisaristifol ?????  
 Pfrank4705 GACCA  
 Pschinz6652 ?????  
 Sbic6588 ?????  
 Sbic6588 GACCA  
 Shalep9318 GACCC  
 Snigros6567 ?????  
 Tleuco6627 ?????  
 Tspic4846 GACCA  
 Ttria6555 GACCC  
 Zeadiplo ?????  
 Zea ATAAT  
 Sacharumofficinarum AAATC  
 Icyindrical AAATC  
 Dichanthiumannulatum ?????

Aadscen6713 ???  
 Aameth6679 AGTT-CAGGGAGATCGGGGGT-TGGGAGAAACTAAGGG-TTGGTA  
 Aameth6685 AGTT-C-GTGAGATAGGGGGT-TGGGAGAAAATAAGGGATTGGTA  
 Aappen6543 ???  
 Aappen6946 ATATTTTGGGACCAAAATTCG---GGAGA---TAGGGG---ATAT  
 Adist6684 CGTTACAGGGAAGACGGGGT-TGGGAGAAAATAAGGGCTTGGTA  
 Aeuco5869 CGGT--CGGGAGATCGGGGGT-TGGGAGAAAANAATGGGGTTGGTA  
 Arundinellaneapalensis ???  
 Aschiren6581 A-GTTCAGGGAGATAGGGGGTGTGGGAGAAAATAAGGGATTGGTN  
 Bbladh AAATNGNCGAGATACCTTTGTTGGGAGAAAACATCGGGTAGAAT  
 Binsc5681 AAAATNG-GGAGAGACGGG--TTGGGAGAAAATAAGGCATTGGTA  
 BBrachiaria ???  
 Bradi4784 ???



Bserra5074 CGATTGTCGGACCAAGTTCAG---GGAGA---TAGGGG---AAAA  
 Bserra6654 ???  
 Bserra6954 CGTTTGTTCGGACCAAGTTCAG---GGAGA---TAGGGGN-TATT  
 Cdieter6642 AAGTTCAGGGAGATAGGGGTT-TGGGAGAAAATAAGGGATTGGTA  
 Cexc6577 AAGT-CAGGGGGATACGGGTT-TGGGAGCAAATAACGGGTCGGTA  
 Cleistachnesorghoides ???  
 Cmarg4397 ???  
 Cplur6406 C---TCGGGAAAATAGGGGTGTGCGGAGGAAAATACGGGTTTGGGA  
 Cprolix6682 CAGT--CGGGAGATAGGGGTT-TGGGAGAAAATAAGGGATTGGTA  
 Danthoniacalifornica ???  
 Dfilli5872 CNTTTTTAGGACCAAGTTCAG---GGAGA---TAGGGGTTGGAA  
 Eragrostisdielsii ???  
 Evil5085 ???  
 Hanam4807 ACCTTCCGCAA-TAGGCCTCCCCCGCCCTTCGGCACACCCA  
 Hcon6942 AGGTTACAGGAGATAGGGGTT-TGGGGCAAATAACGGATAGGTA  
 Hdiss5145a TTATAGCTTACACACGTCCC---TGCACA---AAAGGCACTCCCT  
 Hemarthria9319 G---TCCGGGAATTAGGGGT-CGGAGAACATCAAGGGTTGGGA  
 Hfilli5144 TCTTAGCCCGACCAAGTTTC--GGCACA---TAGGGCCCTATT  
 Hhirt6650 ???  
 Hhirt6656f A-GTTTCGGGAATAGGGGTG--GTGGGAGAACATAAGGGATTGGTA  
 Hlongi6560 ???  
 Hpilg4603 AANTTTAGGGAGATAGGGGTT-TGGGAGGAAAATAAGGGATTGAAA  
 Hpilg4738 TCTTACCGGACCC-AAATTTC--GGGAGA---TAGGGTTATTT  
 Hquar5088 AGTTCAGGAGATAGGGGGTG--TGTTT---TAAGGGTCAGGTA  
 Hrud6576 AAGTTCAGGG-GATAGCGGGTTCGGGAGAAAAGAACCGGATTGAT  
 Htamb6585 AAGTTCAGGG-GATAGCGGGTTTGGGAGAAAATACCGAATTTCGTA  
 Icylin5114a AANTTTCAGGAGATAGGGGGTTTGGGAGAAAATAAGGGTTCCGTA  
 Icylin6632a --GTTACGGGAAAATACGGGGGT--GGGGAAAATAAGGGATTGGTA  
 Mcape6655 ACGTTCAGGAAAATAAGGGGGTTGGGAGAAAATAAGGGATTGGTA  
 Monocymbium ???  
 Pdilatun6617 AGCTTCCGGGAATTAGGGGCCCTGGGGCCAACATTTAGCCCAA  
 Pentaschistisaristifol ???  
 Pfrank4705 AAGTTTTGAGCACCAGTTCG--GGGAGA---TAGGGGTCATTGA  
 Pschinz6652 ???  
 Sbic6588 ???  
 Sbic6588 G---TTCGGGAATAGGGGT-TGGGGAAAATAAGGGATTGGGA  
 Shalep9318 TCTTACCGGACCCCATATA---CCGATT---TCCCGTTATTTT  
 Snigros6567 ???  
 Tleuco6627 ???  
 Tspic4846 TCGTTGTCCANAAAAGTTTC--GCANA---TAGGGGTTATTT  
 Ttria6555 AAGTTCAGCAGATAGGGGT--GTGGGAGAAAATAAGGGATTGGTA  
 Zeadiplo ???  
 Zea C-GGAGAATGCTCATTTGAT--GAGCATTCCTCGTAGAAGTATTC--  
 Sacharumofficinatum CGTCGACTTATA-AGTCGT-GAGGGTTCAAGTCCCTCTATCCC  
 Icyindrical CGTCGACTTATA-AGTCGT-GAGGGTTCAAGTCCCTCTATCCC  
 Dichanthiumannulatum ???

Aadscen6713 ???  
 Aameth6679 TCC--CTTTAGGAAAACAAGAATA---TAGTTACCCA-CACTC  
 Aameth6685 TCC--CTTTAGGAAAACAAGAAA---TAGTTACCCA-CACTC  
 Aappen6543 ???  
 Aappen6946 TCTTCGC---ACACA-AGG-----CA-ATAAAC---CTTG  
 Adist6684 TCC--CATTAGGAAAACAAGAATA---TAGTTACCCA-CATTC  
 Aeuco5869 TCN-GTTATTGTTCAAGTAGAAAAC--TAGTTACCCATCATCC  
 Arundinellaneapalensis ???  
 Aschirene6581 TCA-ACNTTG-GAGATAGGGGA---TAGTTACCC-GGATTC  
 Bbladh CA---CATCA-TCTTACCGAAAGACATAGTTACCCA-TACTC  
 Binsc5681 TCAAATA-TAGGAAA---CAAAGA--TAATAGTTACCAATTC  
 Bbrachiaria ???  
 Bradi4784 ???  
 Bserra5074 ACTGGCG---ACACG-GGC-----CA-ATAAAC---CTTG  
 Bserra6654 ???  
 Bserra6954 TTTCGGG---TAAAG-AGA-----AA-ATAAAC---CTTC  
 Cdieter6642 TCA-GCTATCGGAAACGTAGAAA--TAGTTACCCA-AATTC  
 Cexc6577 CCA-TCATTCTGAAACATAGAAA--TAGTAACCCA-ACTTC  
 Cleistachnesorghoides ???  
 Cmarg4397 ???  
 Cplur6406 TCAA-CTTTAGGAAA--AAGGAA--ATAGGTTTCCC-GATTC  
 Cprolix6682 TCA-GCTATNGGAAAATAAGAAA--TAGTTACCCC-AATTC  
 Danthoniacalifornica ???

Dfilli5872  
*Eragrostisdielsii*  
Evill5085  
Hanam4807  
Hcon6942  
Hdiss5145a  
Hemarthria9319  
Hfilli5144  
Hhirt6650  
Hhirt6656f  
Hlongi6560  
Hpilg4603  
Hpilg4738  
Hquar5088  
Hrud6576  
Htamb6585  
Icylin5114a  
Icylin6632a  
Mcape6655  
Monocymbium  
Pdilatatum6617  
Pentaschistisaristifol  
Pfrank4705  
Pschinz6652  
Sbic6588  
Sbic6588  
Shalep9318  
Snigros6567  
Tleuco6627  
Tspic4846  
Ttria6555  
Zeapdipo  
Zea  
Sacharumofficinatum  
Icylindrical  
Dichanthiumannulatum

G-----AAAATAAGG-----GA-----TTGG  
????????????????????????????????????  
????????????????????????????????????  
GAT-AAAAATCTAGACTCGGA-AAGGCTAGNCCA-CAGTT  
TCA-NCTATAGGAAAAATAGAAAA--TAGTTACCCA-CATTC  
TTCTCCC----ATCTTTTGG-----CACC-AAA----ATTC  
GTCA-ACTTAGGAACA-AAGG----ATAGGTTCCCC-A-TTC  
TTTCGGA----CACCATTG-----CAGATAA-----ATAA  
????????????????????????????????????  
CCA-ACNTTAGGAGAAAAGAGTA--TAGTTACCCAGTATTC  
????????????????????????????????????  
TA--TCATAAATAGGAGAAAAGAAAATAGTTACCCA-AATTC  
TCGGGTC----AATGCGAG-----GATAA-----ATAG  
CATGGGT----TTCGGGACC-----AAGGTTTCAG---GGAG  
GCA-ACATT-GTAAATCTAG-AAA-GTAGTTAGCCG--ATTC  
GCA-CCATTCGTAACATAG-AAA-CTAGTTACCCC-CATTC  
TCATAGAGTATGCAAAAAAGAGTAATAGTTACCCA-AATTC  
TCTG-CTATGGGAAAAAGAAA--ATAGTTACCCA-AATTC  
TCAA-CTTTAAGAAAAGAGAC----AAGTTTCGGG-ACTTC  
????????????????????????????????????  
GCCTTGCTCAGTTATTCTGGTAACAGTAGATATCCCAATTA  
????????????????????????????????????  
TCGGGTC----AATAGGACC-----ANGTTTCAG---GGAG  
P????????????????????????????????????  
????????????????????????????????????  
TCAA-CTTTAGGAAAA-AAGG----NTTGTACCCC-AATTC  
TTTGTA---GACCAAAGT-----CAGATAA-----CATG  
Sn????????????????????????????????????  
????????????????????????????????????  
TCGGACC----AAGTGCAGG-----GAG-----ATAG  
TCC--CATTCGCAAAACAAGAAA--TAGTTACCA--AATTC  
Zeap????????????????????????????????????  
AAT--CTGGCGCTCTCCTCTATC---TAATGAATA--CTTTA  
AAA-CCCTCTTTTATCCCTAACC-ATAGTTGTAT-CCTTT  
AAA-CCCTCTTTTATCCCTAACC-ATAGTTGTAT-CCTTT  
????????????????????????????????????

Aadscen6713  
Aameth6679  
Aameth6685  
Aappen6543  
Aappen6946  
Adist6684  
Aeuco5869  
Arundinellaneapalensis  
Aschiren6581  
Bbladh  
Binsc5681  
BBrachiararia  
Bradi4784  
Bserra5074  
Bserra6654  
Bserra6954  
Cdieter6642  
Cexc6577  
Cleistachnesorghoides  
Cmarg4397  
Cplur6406  
Cprolix6682  
Danthoniacalifornica  
Dfilli5872  
*Eragrostisdielsii*  
Evill5085  
Hanam4807  
Hcon6942  
Hdiss5145a  
Hemarthria9319  
Hfilli5144  
Hhirt6650  
Hhirt6656f

????????????????????????????????????  
TGA-GTG-ATC-GAAAGAG-TAAG-AGGAGAAAG-TGTTTCCTCACGC-  
TAA-GTG-ATC-GAAAGAC-TAAG-AGGAGAAAG-TGTTTCCTCACGC-  
????????????????????????????????????  
ACCCCGGT--G-CTCCTAAATATC----GGACC--ACGTTTCAGGGAGAT  
TAAATGTG-ATC-GAAAGACGTAAG-ATGAGAAAG-TGTTTCCTCACGC-  
TGGTGTGGAGC-GAAAGAA-TAAG-ATGAGAAAGTGTGTTCTTCGCGC-  
????????????????????????????????????  
TAG-GTGA-TC-GAAATGAGTAAGATGAGAAATG-AGTTTCCTCACGC-  
TAAATGTA-ATCTNAAANAC-TAATCATGCCATCGGTCTTTCCTCCCC  
TAA-GTG-ATC-GAATCAG-TAAG-ATGAGATAG-TGTCTCCTCACGC-  
????????????????????????????????????  
--CACTGT--G-CTCA-AAATATC----GGACC--ACGTTTCAGGGAGAT  
????????????????????????????????????  
A-CNCNGT--G-CTCATAAAAATC----AGACC--ACGTTTCAGGGAGAT  
TGA-GTG-ATC-GAAAGAG-TAAG-ATGAGAA-AGTGTTCCTCACGC-  
TTT-GTG-ATC-TAAAGAG-TAAG-ATGAGTT-AGCGTTTCTCACGC-  
????????????????????????????????????  
????????????????????????????????????  
TGA-GTG-ATC-GAAA-ACGTAAG-ATGAGAAA-GAGTTTCCTCACCT-  
TCA-GTG-ATC-GAAAGAG-TAAG-ATGAGAG-AGTGTTCCTCACGC-  
????????????????????????????????????  
TATAAAAATAAGGGATTGGTATGGA----TTG----GTATT-----AT  
????????????????????????????????????  
????????????????????????????????????  
TAAGGACAATAGGAGTTATTTTC--GGGAAATATATATATCCTAGGTCT  
TAAATGTG-ATC-GAAAGAA-TAAG-ATGAGAA-TGTGTTTCCTCGCGC-  
AAGGAAATAGGGGTTTGGGACAAA----ATAAC--GGATTGGT----AT  
TAA-GTA-ATC-GATAAAGTACTA-A-GAGAAA-GTGTACCTCAGC-  
CAGTGTGT--GAGATAGAAATAAT----TACCA--AGTTTCAT---CTA  
????????????????????????????????????  
TGA-GTGGATC-GAAAGACGTAAGATGGAGAAAG-TGTTTCCTCAC-C-

*Hlongi6560* ???  
*Hpilg4603* TA-AGTA-ATC-GAAAGAG-TAAG-ATGAGAAAG-TGTTTCCTCACGC-  
*Hpilg4738* CATGGGTTGGAGGATAAAAAATTAA----GTACC--AAGTTTGT---CCT  
*Hquar5088* CGTAGGTGTTGGGAGAGAATAAG----GGATA--GGTATCAT---CAT  
*Hrud6576* TTA-GCG-ATCGAGGAGAG-TAAG-ATGAGAAAG-CGTTTCCTCACGC-  
*Htamb6585* TAA-GTT-ATCGAA-AGCG-TAAG-ATGAGATTG-CGTTCCCTCACGC-  
*Icylin5114a* TAA-GTG-ATC-GAATCAG-TAAG-ATGAGAAAG-TGTCTCCTCACGC-  
*Icylin6632a* NAA-GGNA--T-CAAANA-GNAAG-GG-AGNAAANAGTTTCCTCCC-C-  
*Mcape6655* TCA-TTG-ATC-GAAATAG-TAAG-AGGAGACCG-AGTTTCCTCACGC-  
*Monocymbium* ???  
*Pdilatum6617* AAAGCCCATATAAATGGGCTTAA---GAGTCACTAGCTTTATCATTCT  
*Pentaschistis aristifol* ???  
*Pfrank4705* ATAGGGGTTGGGAGAAAA-TAAG----GGATT--GGTACCAT---CAA  
*Pschinz6652* ???  
*Sbic6588* ???  
*Sbic6588* TAA-GGG-ATC-GAAAGATGTAAG-G-GAGAAA-GTGTTCCTCACGC-  
*Shalep9318* GTGGCGTT-GGAAAAGAAATAAC----AGGTT--TGTCTCAC---CT-  
*Snigros6567* ???  
*Tleuco6627* ???  
*Tspic4846* CGGGGGGGGAGGGAGAAAA--TAG---GTACC--GGTATCGT---CAT  
*Ttria6555* TTG-GTGG-TC-GAAAGA-GTAAGT--GAGGATG-AGCAGTCTCACGC-  
*Zeadiplo* ???  
*Zea* TTG-ATTT-TT-TAGTCC-CTTTAA--TTGAAT--AG-----ATGC  
*Sacharum officinarum* TTT-TCT-TTT-ATCAAT-GGG--TTT----AAGA--TTC-A-CTAGC-  
*Icylindrical* TTT-TCT-TTT-ATCAAT-GGG--TTT----AAGA--TTC-A-CTAGC-  
*Dichanthium annulatum* ???

*Aadscen6713* ???  
*Aameth6679* -TGTTTCG-TGAGTTACTCA-GAATAC-GATCA-GTAATTTAT-CTACTAA  
*Aameth6685* -TGTTCT-TGAGTTACTCA-GAATAC-GATAA-GTAGTTAT-CTACTAA  
*Aappen6543* ???  
*Aappen6946* GGGGTGGGGAGAAAAATAAGGGATTGGTATCAACTTTCGGAA---AC-AA  
*Adist6684* -TGTTCT-TGAGTTACTTA-GAATAC-GATAA-GTAAGTTAT-CTTCTAA  
*Aeucu5869* -TGTTCAGAATGACTGA-GAATAC-GATCG-CCAAGGTAT-GTACGAA  
*Arundinellane palensis* ???  
*Aschiren6581* -TGTTT-TGGAGTTACTTA-GAGTAC-GATAA-GTAATTTAT-CTGCTAA  
*Bbladh* -TTCTCACGGCGTTACT-A-GAATCCCGATAACGTAATTTCTTCTTCTAA  
*Binsc5681* -TGTTCT-TGAGTTACCGA-GAATAC-GATGA-GTGAGTTAT-CTACGAA  
*BBrachiaria* ???  
*Bradi4784* ???  
*Bserra5074* GGGGTTGGGGAGAAAAATAAGGGATTGGTATCACCAATAGGAA---ACGAA  
*Bserra6654* ???  
*Bserra6954* GGGGTTGGGGAGAAAAATAAGGGATTGGTATCACCTATCGGAA---ACTAA  
*Cdieter6642* -TGTTCTTGA-GTTACTTA-GAATAC-GATCA-GTAATTTAT-CTACTAA  
*Cexc6577* -TGTTCTTGA-GTTACTTA-GA-TAC-TATC--CTAAGTTT-CTTCTAG  
*Cleistachnesorghoides* ???  
*Cmarg4397* ???  
*Cplur6406* -GTT-CT-TGAGTTCTGATGGATAC-GATAA-GTTGTTTAT-CTACTAA  
*Cprolix6682* -TGTTCTTGA-GTTACTTA-GAATAC-GATAA-GTAAGTTAT-CTACTAA  
*Danthoniacalifornica* ???  
*Dfilli5872* ---CATCAATA-GGAA-AATAGTTACCCAAATTCTAAG-TAA---TCGAA  
*Eragrostis dielsii* ???  
*Evill5085* ???  
*Hanam4807* AGGGTTGGAAATAAGAAGA-TAATACCGAGTTTGTCTCTCGGTGCT-CGAG  
*Hcon6942* -TGTTCTATA-AACTTTAC-CACTGG-CTC--CTGAGTTAT-CG-CTGA  
*Hdiss5145a* ---CATCAATA-GGAA-AATAGTTACCCAAATTCTAAG-TAA---TCGAA  
*Hemarthria9319* -CCT-CT-GCGGTTTTTTA-GAATAC-GACAA-CTAATTTAT-CTACTAG  
*Hfilli5144* TAGCTACAAAA-AAAATAATACTAACCCAAAGTTCC-CT-AAG---CCTAA  
*Hhirt6650* ???  
*Hhirt6656f* -TGTTCTGACGTTGCTGA-AGATAC-GATAAAGTAAATTTAT-CT-AGTA  
*Hlongi6560* ???  
*Hpilg4603* -TTCTCT-TGAGTTACCTA-GAAT-----A-GGATCTTAT----ATAA  
*Hpilg4738* AAGGTGATAGA-AAGAATA-AGATGCCAAAGTTAGCCTTG---GCTAG  
*Hquar5088* TAGGAAAAAGAGAAGAACATAGTTTCCAGATTCTACG-TAG---TCGAA  
*Hrud6576* -TGTTCT-TGAGTTACTTA-GAATAC-GATAG-GTAATTTT--CTGCTAG  
*Htamb6585* -TGTTCT-TGAGTTACTTA-GAATAC-GATAA-GTAATTTT-CTACTAA  
*Icylin5114a* -TGTTCT-TGAGTTACTTA-GAATAC-GATAA-GTAAGTTAT-CTACTAC  
*Icylin6632a* -TGNCT-TCAGTTTCTTA-GAATCC-GATCA-GTTATTTAT-CTACTAA  
*Mcape6655* -TGCTCT-TGCGTTACTTA-GAATACGA-TAG---TATTTTACCTAGTCA  
*Monocymbium* ???

*Pdilatum6617* CTAT----GGAGCAAAGGA-GAGTGACGAGAAGCTCAATGTATCTTATGCT  
*Pentasthisisaristifol* ???  
*Pfrank4705* TAGG--AAAAA-AAGAAAATAGTTACCCAAAGTCTAAG-TAA---TCGAA  
*Pschinz6652* ???  
*Sbic6588* ???  
*Sbic6588* -CGTTCT-TGGGTTTCTTA-GAATAC-GATAA-GTAATTTAT-CTACTAG  
*Shalep9318* TCCAAAAATA-AAAAAATACCTACCCAAATTCTACT-TCT---TCGAA  
*Snigros6567* ???  
*Tleuco6627* ???  
*Tspic4846* AGGA--AAAAA-AAGAATATAGTTACCCAAATTCTAAG-TAG---TCGAG  
*Tria6555* -AGTTC-TGA-GTTACTTA-GAGTAC-GATAA-GTAATTTAT-TTGAGCT  
*Zeadipl* ???  
*Zea* -AAATA---CATTTACTAA-GAGATG-CACAA-GAAAGGG---TCAGGAT  
*Sacharumofficinarum* --TTTC-TCA--TT-CT-AC-TCTTT-CACAAAGGAGTGCACAAAGAACT  
*Icylindrical* --TTTC-TCA--TT-CT-AC-TCTTT-CACAAAGGAGTGCACAAAGAACT  
*Dichanthiumannulatum* ???

*Aadscen6713* ???  
*Aameth6679* AGTTAAACT-----AAAGAAAAAATCAA-TAATCTCATCTCAT--  
*Aameth6685* AGATATTCT-----ACAGAAAAAATCTA-TAATCTCATCTCAT--  
*Aappen6543* ???  
*Aappen6946* GA-----AACCAATAT-----AATCCCATCTCTTAG  
*Adist6684* AGCTTATCT-----ACAGAAAAAATCTA-TAATCTCATCTCAT--  
*Aeuco5869* GG-----AATCTAGCATA--AGGCTCATA-----  
*Arundinellaneapalensis* ???  
*Aschiren6581* AGCCAAGGTC-----TGGATGAAGAAATCTA-TAATCTCGTCTCAA--  
*Bbladh* AA-----AACCAAT-----AATCTCAT  
*Binsc5681* TGT-----AATGATAAAGATTA-TAATCTCATCTCAA--  
*BBrachiaria* ???  
*Bradi4784* ???  
*Bserra5074* GA-----AAAAATAT-----AATCTCATCACTTAG  
*Bserra6654* ???  
*Bserra6954* GA-----AACCAATAT-----AATCCCATCACTTAG  
*Cdieter6642* AG-----ACAT-AATATA-TAGTCTCATA-----  
*Cexc6577* AT-----ACCTTAATCTA-TAGTCTCAA-----  
*Cleistachnesorghoides* ???  
*Cmarg4397* ???  
*Cplur6406* ACTAAGGCT-----CGAGAGAAGAACTCA-CAATCTCGTCTCTG--  
*Cprolix6682* AG-----ACTT-AATCTA-TAGTCTCATA-----  
*Danthoniacalifornica* ???  
*Dfilli5872* AG-----AGTAAGATG-----AGAAAGTGTTTCC  
*Eragrostisdielsii* ???  
*Evill5085* ???  
*Hanam4807* GGTTACAGACGATACGATGTGTGCCTTAGCTTGCATAAAAAAACCTAGCA  
*Hcon6942* AC-----AGCGAGCACA--AGCTTCGTA-----  
*Hdiss5145a* AG-----AGTAAAATG-----AGAAAGTGTTTCC  
*Hemarthria9319* AGAGCTTATA----G-ACGAAAGAAAAACAAA-TAATCTCGTGTCAA--  
*Hfilli5144* AT-----AAAAAACCTA--GAATAAAGTAATTTCC  
*Hhirt6650* ???  
*Hhirt6656f* AGCCTAGGCT-----CGAGAGAAGAAATCTA-GAATCTCGTCTCTA--  
*Hlongi6560* ???  
*Hpilg4603* AG-----AAAAAAT-----AATCTCATAGCC--  
*Hpilg4738* AA-----TGAAGAAACCT--AGAATAGGGTAACTA  
*Hquar5088* AG-----AGTATGTTG-----AGACAGGGTTTCC  
*Hrud6576* AGT-----GAAGAAATAAA-TAATCGCTCTCA--  
*Htamb6585* AGT-----AAAAAATAAA-TAATCTCGTCTCAT--  
*Icylin5114a* AGG-----AATGATGAAGCCTA-TAATCTCATCTCAT--  
*Icylin6632a* ATAAA-----GCATGAAGGAAAAACCTA-CTATATAGTCTCAT--  
*Mcape6655* ACCTAAGATC-----TAGTGAAGAAATCCA-GAATCTCATCTCCT--  
*Monocymbium* ???  
*Pdilatum6617* AC-----ACACGGAAT-----AGATGGTATTTT--  
*Pentasthisisaristifol* ???  
*Pfrank4705* AG-----AGTAAGA-----AGAGAAAGGGTATCC  
*Pschinz6652* ???  
*Sbic6588* ???  
*Sbic6588* AGAGCCTATA---G-AGGAAAGAAAAATCAA-TAATCTCGTGTCTT--  
*Shalep9318* AG-----AGCCAGATGAG--CAATAAATCGTTTCC  
*Snigros6567* ???  
*Tleuco6627* ???  
*Tspic4846* AG-----AGTAAAATCCG--AGAGTAAGGGAAGCC

*Ttria6555* ACTCAAGGCC-----CGATTGAGGAAACCTA-GAATCTCGTCTCGA--  
*Zeadiplo* ???  
*Zea* ACTCATGGTA-----GAGCAGAGGAC-TGAA-AATCCCTCGTGCAC--  
*Sacharumofficinatum* A-TGAATCTTATGCTATTCATTAATA-----GATGATTTCTTTTTTA--  
*Icylindrical* A-TGAATCTTATGCTATTCATTAATA-----GATGATTTCTTTTTTA--  
*Dichanthiumannulatum* ???

*Aadscen6713* ???  
*Aameth6679* AGCCTTTCCTTAGAG-CTACTAATTC-ATCCATTAATAATTCATAA  
*Aameth6685* AGCCTTTCCTTAGAG-CTACTAATTC-ATCTATTAATAATTCATAA  
*Aappen6543* ???  
*Aappen6946* CCGTTCCTTAGAGATAATAATTAAGC-TAATAAAATTCATAATAC  
*Adist6684* AGCCTTTCCTTAGAG-CTAATAATTC-ATCCAAAAAATTCATAA  
*Aeuco5869* --TCTTTCCTTAGAG-CTAACATTC-CGCTCATAGGGTCATAA  
*Arundinellaneapalensis* ???  
*Aschiren6581* AGCCTTTCCTTAGAG-CTACTAATCC-GGCTATTAACATTCATAA  
*Bblad* ---CTTTATGTAGAC-CTGTAATCA-ATTTATTANTCTTCATAA  
*Binsc5681* AGCCTTTCCTTAGAG-CTAATTATCA-GACTATTAACATTCATAA  
*BBrachiaria* ???  
*Bradi4784* ???  
*Bserra5074* CCGTTCCTTAGAGATAACTAATTAAGC-TACTAAAATTCATAATAA  
*Bserra6654* ???  
*Bserra6954* CCGTTCCTTAGAAATAATAATTAAGC-TACTAAAATTCATAATAA  
*Cdieter6642* -GCCTTTCCTTAGAG-CTAATAATTA-AGCTATTAACATTCATAA  
*Cexc6577* -GCCTTACCTTAGACTCTACTAATTT-ATCTATTACATTCATAA  
*Cleistachnesorghoides* ???  
*Cmarg4397* ???  
*Cplur6406* AGTCTTCCCTTAGAG-CTACTTATCC-GTCTATTAGCATTTCATAA  
*Cprolix6682* -GCCTTTCCTTAGAG-CTACTAATCC-AGCTATTAACATTCATAA  
*Danthoniacalifornica* ???  
*Dfilli5872* TTACGCTGCTTTTGGAGTTACTTAGAA-TACGATAAGTAAT-TTAT  
*Eragrostisdielsii* ???  
*Evill5085* ???  
*Hanam4807* CGGCGCAAGTACGTTCTAGTACAACAGTCTTTCCTGTTCTCGGAT  
*Hcon6942* -GTCTTCCCTGAAAC-CTCACAAATC-TGCTCTAATCCGTGATAA  
*Hdiss5145a* TCACCCTTCTTGGAGTTTCTTAAAA-TACGATAAGTAAT-TTAT  
*Hemarthria9319* AGTCGTTCCTTAGAG-CTACTAATTA-AGCCAATAAAATTCATAA  
*Hfilli5144* TCTAACTTCTCTAATATCACTTAC-A-TTCATTAGGGATAATAT  
*Hhirt6650* ???  
*Hhirt6656f* AGCCTTCCCGTATAG-CTACTTATCC-GTCCATTAAGATTCATAA  
*Hlongi6560* ???  
*Hpilg4603* ---CTTCTTAGGGT-CAATAAGTGAGATAAAATAGTCATAATAA  
*Hpilg4738* TGTTCCTCGTATAAATATCATATCCG-TTCATTAGGCATA-ATCT  
*Hquar5088* TCACACTGCTCTCGTGTGCTTAGAA-ATCGATACGTTAT-TTGC  
*Hrud6576* AGCCTTTTCCTTAGAG-CAACTTATTT-GTTTATTAGCATTTCATAA  
*Htamb6585* AGCCTTTTCCTTAGAG-CTACT-ATTA-ATCTATTAACCTTCATAA  
*Icylin5114a* AGCCTTCCCTTAGAG-CTACTTATTC-ATCCATTAAGATTCATAA  
*Icylin6632a* AGCCGTCCTTAGAG-CTACTTATCA-ATCTATAAACATTCATAA  
*Mcape6655* AGCCGTTTCTTAGAG-CTACTCATTA-GTCTATAAGGTTTCATAA  
*Monocymbium* ???  
*Pdilatum6617* ----TTTGGATAGAGACGCGT--TTA-GGC---AAGGGATCATGA  
*Pentaschistisaristifol* ???  
*Pfrank4705* TCACGCCGCGCATGAGATACTTAGCA-TACGATAAGTAAT-ATAT  
*Pschinz6652* ???  
*Sbic6588* ???  
*Sbic6588* AGCCGTTTCCTTAGAG-CTACTAATTA-AGCCAATAAGATTCATAA  
*Shalep9318* TCACGCTGCTCTTTTTTTCCTTAGTA-TACGATAAGCATT-TTAT  
*Snigros6567* ???  
*Tleuco6627* ???  
*Tspic4846* TCTCGCCGCGCATGAGTTCCTAACCG-TTCAATAGGTATT-TTAT  
*Ttria6555* AGCCTTCCCGTAGAG-CTACTTATCC-GTTCATTAGGGATTCATAA  
*Zeadiplo* ???  
*Zea* CAGTTCAA-TCTGG-TTCCTCAGA----AAAAAAGGATCATAA  
*Sacharumofficinatum* TTTCTTTTTTTTATTTATTTAGAGTAGAGT--ATCGGCAA--GGAA  
*Icylindrical* TTTCTTTTTTTTTTTTATTT-ATTAGAGT--ATCGGCAA--GGAA  
*Dichanthiumannulatum* ???

*Aadscen6713* ???  
*Aameth6679* TAAATCAT-GCATGAGGT---GTCCGTATCCTGAAGGGGAGGC

*Aameth6685* TAATTCATTGCAAACGGT---GTTCCGTATCCTGAAGGGGAGGC  
*Aappen6543* ???  
*Adist6946* TTCATTC-GGTAGGTGTT-----ACGTATCCTGATGC-GAGGG  
*Adist6684* TACATCAT-GCGTACGGT---GTTACGTATCCTGAGGGGGAGGC  
*Aeuco5869* TTACTTTACCCTATCGGT---GTTTCATACACTGAGGACGAGCC  
*Arundinellaneapalensis* ???  
*Aschiren6581* TTATTCAT-GCAATAGGT---GTTTCGTATCCTGAGGGGGAGGC  
*Bblad* CACTTTACTTCCATGGGT---TTTCCGTCTGGGCTAATACCTA  
*Binsc5681* TTAATCAT-TCAATAGGT---GTTTCGTATACTGAGGGGGAGGG  
*BBrachiaria* ???  
*Bradi4784* ???  
*Bserra5074* TTCATGC-GGTAGGTGTT-----ACCTATCCTGATGC-GAGGG  
*Bserra6654* ???  
*Bserra6954* CTCATTC-GGTAGGTGTT-----ACCTATCCTGATGC-GAGGA  
*Cdieter6642* TAATTC-AGGCGGTAGGT---GTTTCGTATCCTGAGGGGGATGG  
*Cexc6577* TTAATGCAACCTATC-GT---GTTTCATTTACTCAAGCCATCC  
*Cleistachnesorghoides* ???  
*Cmarg4397* ???  
*Cplur6406* TTAAGTAACCAGTCGTT---GTTTCGTATACTCAAGCCGAGGG  
*Cprolix6882* TTATTC-ATGCAGTAGGT---GTTTCGTATACTGAGGCGGACGG  
*Danthoniacalifornica* ???  
*Dfilli5872* CTAATA-AAGAAAAA-----TAACTATCCTGATGGGGACGG  
*Eragrostisdielsii* ???  
*Evoll5085* ???  
*Hanam4807* AATATTCACCTCAGTAAGT---GTCATGTCCAGTAAGGG-GAAGG  
*Hcon6942* TAACAA-ACGGGGTGCGT---GTTGCGTACTCTTAGGCTGATCC  
*Hdiss5145a* CTACTA-AAAAA-----GAAACTATCCTTATGGGGGGAG  
*Hemarthria9319* T-AATTCACCTCAATGTT---GTTTCCTATCCTGAGGCGGACGA  
*Hfilli5144* TCATTC-AGTAACGTGTT-----ACGTATACCAATAG-GACTG  
*Hhirt6650* ???  
*Hhirt6656f* TTCTTTCT-GCGGGAGTT---GGTTCGTGTCCTCAGGCGGAGGG  
*Hlongi6560* ???  
*Hpilg4603* TTCTTCGGTACATGTGTAGGTATCTTGATGGGGGGGTTAAAG  
*Hpilg4738* TTCATTC-AGTAAGTGT-----AGGTATAGTAATAG-GACGG  
*Hquar5088* CTACTA-TAGAAA-----TAAGTACTCTCATCG---CCG  
*Hrud6576* TTAATTAG-GCCATCGTT---GGTTCGTACTCAAGCGGAGGG  
*Htamb6585* TTCTTTAA-TCAATCGTT---GTTTCGTATCCTCAAGCGGAGCG  
*Icylin5114a* TTCATCAA-TCAATAGGT---GTTTCGTATCCTAAGGCGGAGGG  
*Icylin6632a* T-TATTCATCTATCGTT---GTTTCGTATACTCAAGCCGAGCG  
*Mcape6655* T-AATTTAAGCAGTAGTT---GTTTGGTATCCTCAAGCGGAG-C  
*Monocymbium* ???  
*Pdilatum6617* TTAAGATAGGAAA-----GTAAATGTATAATAAAGA-GAGAG  
*Pentasthisaristifol* ???  
*Pfrank4705* CTAATA-AGAAAATGTT-----AACTATCCTAATGA-GGCGG  
*Pschinz6652* ???  
*Sbic6588* ???  
*Sbic6588* T-ACTTCATTCGGTAGGT---GTTTCGTATCCTGATGGGGAGGG  
*Shalep9318* CTGCTA-AAGGAAGCT-----AATCCATCCGAGTGA-AACCG  
*Snigros6567* ???  
*Tleuco6627* ???  
*Tspic4846* CTCCTAC-AGTAAGAGTT-----AGCTATCCTGATAG-GACGG  
*Ttria6555* TTCATTCA-GCAATGTT---GTTTCGTGTCCTCAGGCGGAGGG  
*Zeadiplo* ???  
*Zea* TGAATAGA-TCAATGAT---TACAAATATTTGAGATGGAGGG  
*Sacharumofficinarum* T--CTCGATTATTAATTCGATTTTAAAGTATTATTA-GTAAGCC  
*Icylindrical* T--CTCGATTATTAATTCGATTTTAAAGTATTATTA-GTAAGCC  
*Dichanthiumannulatum* ???

*Aadscen6713* ???  
*Aameth6679* -GTAAGG-A-TTAAA--ACTTACCTTATGA--AT-AACTAAACAA--TC  
*Aameth6685* -GTAAGG-A-TTACA--TCTTACCTTATCA-CAT-TACTAAACAA--TC  
*Aappen6543* ???  
*Aappen6946* -GTAAGG-ATTAAAA--CTTTTCTTACCGA-AAT-AACTAAAA--AATC  
*Adist6684* -GTAAGG-ATTAAAA--ACTTACCTTATGA-AAT-AAATAAACAA--TC  
*Aeuco5869* -GTAAGG-CATCTTAA---CTTATCTTGTGA-AAT-TCCTAATCA--GGC  
*Arundinellaneapalensis* ???  
*Aschiren6581* -GTAACA-ATTAAAA---CTTATCTTACGA-ACT-AACTAAACAA--TC  
*Bblad* -CTCAATT-ATTCATC---TT---CTTTCCCTTTT-ACCTAAACAA--T-  
*Binsc5681* -GGAAGG-ATTAAAA---CTCACCTTATGA-AAT-AACTAAAAAA--GC  
*BBrachiaria* ???

<i>Bradi4784</i>	??
<i>Bserra5074</i>	-GTATAGG-AGTAAAA--CTTATCTTAC-GA-AAT-AACTAAAA--AATC
<i>Bserra6654</i>	???
<i>Bserra6954</i>	-GTATAGG-ATTAAAA--CTTTTCTTAC-GA-AAT-AACTAAAA--AATC
<i>Cdieter6642</i>	-GTAAGG-GATTAAAA---CTTACCTTACGA-AAT-AACTAAACA--ATC
<i>Cexc6577</i>	-TTAAACCAATTCAAAA---CTCATCTTACGC-AATTAACCTAAACAT-AGC
<i>Cleistachnesorghoides</i>	???
<i>Cmarg4397</i>	???
<i>Cplur6406</i>	-GTAACA-ATTAAAA--CTTATCTTACGC-AAT-AACTCAAAATAGCA
<i>Cprolix6682</i>	-GGAAAC-AATTAAAA---CTCACCTTACGA-AAT-AACTAAACA--ATC
<i>Danthoniacalifornica</i>	???
<i>Dfilli5872</i>	-GTAATG-GTTTTAAA--CCTTA-----TGA-AAT-AACTAAAA--AATT
<i>Eragrostisdielsii</i>	???
<i>Evill5085</i>	???
<i>Hanam4807</i>	--TAAAGG-ATTCAGG---CTTAACTCTTATGTAT-AAATAAAAA--ATC
<i>Hcon6942</i>	-TTACT-TATCTCGC---CTTATCTTCCGA-ATT-GTCTAAACA--AGG
<i>Hdiss5145a</i>	GGTAAAGG-GTTTTAAA--CCTTACCTTATGA-AAT-AACTAAAAT-AATC
<i>Hemarthria9319</i>	-GTATACG-ATTAAAA---CTTATCTTATGA-AAT-AACTAAACA--ATC
<i>Hfilli5144</i>	--TATAAG-ATTCAGA--CCTTACCCTTATA-AAT-AAATAAAA--AATC
<i>Hhirt6650</i>	???
<i>Hhirt6656f</i>	-GTAACA-ATTAAAA--CTTATCTTATGA-AAT-AACTAATCAG--TC
<i>Hlongi6560</i>	???
<i>Hpilg4603</i>	-GTTAAAT-CTTAAAC-TTTATGAAATAAAT---T-AACTAAAAAA--ATC
<i>Hpilg4738</i>	--TGATG-CTTCAGG--CCATACCTTTAGG-AAT-AACTAAAA--AATC
<i>Hquar5088</i>	----TTC-CTTAGAG--CCTTT----TATT-AAT-G-CTAAAA--CTTC
<i>Hrud6576</i>	-GGAAACA-ATTAAAA---CTTATCTTACGA-AGT-AACTAAACAT-AGC
<i>Htamb6585</i>	-GTACAGA-ATTCAAA---CTTATCTTATGA-AAT-AACTAACAAAT--GC
<i>Icylin5114a</i>	-GTAACA-ATTAAAA---CTTACCTTATGA-AAT-AACTAATCAA--GC
<i>Icylin6632a</i>	-GTAACA-ATTCAAC---CTCATCTTACGA-AAT-AACTAAACA--ATC
<i>Mccape6655</i>	-TTATACA-ATTCAAA---CTCATCTTACGCTAT-CACTACACAA--GC
<i>Monocymbium</i>	???
<i>Pdilatum6617</i>	-TTATTCAAAGTATAG-TCATTCCTTAC---ATTTCCCTAAGTTCGAAT
<i>Pentaschistisaristifol</i>	???
<i>Pfrank4705</i>	--GTAAAG-GTTCAAA--TCTTACCTT-AGA-AAT-AACTAAAA--AATC
<i>Pschinz6652</i>	???
<i>Sbic6588</i>	???
<i>Sbic6588</i>	-GTAAGG-ATTAAAA--CTTACCTTATGA-AAT-AACTAAACA--ATC
<i>Shalep9318</i>	---ATACA-GTTCAAA--CCCTACCTTTAGA-AAT-AACTAAAA--AATC
<i>Snigros6567</i>	???
<i>Tleuco6627</i>	???
<i>Tspic4846</i>	--TGTAAG-GTTCAGG--TCTTACCTTTTGG-AAT-AATAAAA--AATC
<i>Ttria6555</i>	-GTAACA-AGTAAAC---CTCATCTTATGA-AAT-AACTAATCAG--TC
<i>Zeadiplo</i>	???
<i>Zea</i>	-GTAATA-TTTAAAA--ATCTACTTAGTC-TTT-AGTCTAGAGT--AA
<i>Sacharumofficinatum</i>	-ATCCACA-ATGCATAGGACT-ACCCC-TCCCATTTCCCTAATTTGGAAT
<i>Icylindrical</i>	-ATCCACA-ATGCATAGGACT-ACCCC-TCCCATTTCCCTAATTTGGAAT
<i>Dichanthiumannulatum</i>	???

<i>Aadscen6713</i>	???
<i>Aameth6679</i>	AGGGA-AATTAACCTG-TAT--CTTCGTTACCG-AGATGATT-CTACTA-C
<i>Aameth6685</i>	AGGGA-AATTAGCTC-TAT--CTTCGTTACCC-AATGGACA-CTATTA-C
<i>Aappen6543</i>	???
<i>Aappen6946</i>	AGGGA-AATT-GACTGTAT--GTTGTTTT-ACGAAAGGATACCTACTG-C
<i>Adist6684</i>	AGGGA-AATTAACCTG-TAT--CTTCGTTACCGAGATGATA-CTACTA-C
<i>Aeuco5869</i>	AGGGA-AACTGTCT-GTAT--CTTCGTTT-ATGAGATG-ATTCTACTA-C
<i>Arundinellanepalensis</i>	???
<i>Aschiren6581</i>	AGGGA-AACTAACT-GTAT--CTTCGTTTAC-CAGATGATA-CTACTA-C
<i>Bbladh</i>	AGAGG-AGGTAATTA-TAT--CTTTATACTTGT--ATGATT-CTATTA-T
<i>Binsc5681</i>	AGGGA-GATTGACTG-TAT--CTTCGTTCAAG-AGATGATA-CTACTA-C
<i>BBrachiararia</i>	???
<i>Bradi4784</i>	???
<i>Bserra5074</i>	AGGGA-AATT-TACTGTAT--GTTGTTTT-CCGAGATGACAACCTACTA-C
<i>Bserra6654</i>	???
<i>Bserra6954</i>	AGGGA-AACT-GACTGTAT--GTACGTTT-ATGAGATGATC-CTACTA-C
<i>Cdieter6642</i>	AGGGAG-ATTAACCT-GTAT--CTACGTTTACCGAGATGGATACTACTA-C
<i>Cexc6577</i>	ACTTGAAATTTATCT-CTATT-CTTCGTTT-ACCAATG-ATCCTACTA-C
<i>Cleistachnesorghoides</i>	???
<i>Cmarg4397</i>	???
<i>Cplur6406</i>	CGGGG-CCTTGG-GCGCATATCTTCTTAG-ACCAATGGACA-CTTTTA-C
<i>Cprolix6682</i>	AGGGAGGATTGACT-GTAT--CTACGTTT-CCGAGATG-ATACTACTA-C

<i>Danthoniacalifornica</i>	??
<i>Dfilli5872</i>	AGGGA-TGTTTAAACGGTCT--CTTCGTCT-ACGATTGGCAC-CTTTTGGG
<i>Eragrostisdielsii</i>	??
<i>Evill5085</i>	??
<i>Hanam4807</i>	AGGGA-AATTAACGT-GTAT--CTACGTTC-ATGAGATGATA-CTACTA-C
<i>Hcon6942</i>	AGCGA-CTGTGCGT-ATA---CCTTGTCT-TCGCGCGG-ATTONTCTN-N
<i>Hdiss5145a</i>	ACCGA-AATTTACGGTCT--TTATTCCC-ACCGTTTGCCA-CTTTTGTGTA
<i>Hemarthria9319</i>	AGGGA-AATTGGACTGTAT--CTTCGTT--ACCAAAGGATA-CTACTA-C
<i>Hfilli5144</i>	AGCGA-AATTGACCTAT----CTTTTCCC-ACCATGTGCCA-CTTTTGTGTC
<i>Hhirt6650</i>	??
<i>Hhirt6656f</i>	AGGGA-ACTGTCT-CTAT--CTTCGTTCCCGAGATGATC-CTACTA-C
<i>Hlongi6560</i>	??
<i>Hpilg4603</i>	AGGGA-AATTAACGGGTAT--CTAGGTTTATGA-GAGGACA-CTTTTA-C
<i>Hpilg4738</i>	CGGGA-AATT-AACTGTAT--CTACGTTC-ATGAGATGATC-CTACTG-C
<i>Hquar5088</i>	AGGGA-AATT-AACTGTAT--CTACGTTC-ATGAGATGATC-CTACTA-C
<i>Hrud6576</i>	AGGGA-AACTGTCTG-TAT--CTTCGTTCA-CCAAATGATA-CTACTA-C
<i>Htamb6585</i>	ATGGA-AACTAATG-TAT--CTTCGTTAC-CGATATGATA-CTCTTA-C
<i>Icylin5114a</i>	AGGGA-AATTAACGC-TAT--CTTCGTTCCACG-ATAGGACA-CTACTG-C
<i>Icylin6632a</i>	AGGGA-AATTGA-CTGTAT--CTTCGTTCC-CCAAATGATAACTACTA-C
<i>Mcape6655</i>	AGTTG-GACCTGCC-CTAT--CTTCGTTTCATC-TATTGTCC-CTACTA-C
<i>Monocymbium</i>	??
<i>Pdilatum6617</i>	AGAGA--ATGCATATTTATTTTTTAGTCCCTTTAATTGACA--TAGAAGC
<i>Pentaschistisaristifol</i>	??
<i>Pfrank4705</i>	AGGGA-AATT-AGCGGTAT--CTACGTTC-ATGAGAGGACC-CTACTG-C
<i>Pschinz6652</i>	??
<i>Sbic6588</i>	??
<i>Sbic6588</i>	AGGGA-AATTAG-CTGTAT--CTACGTTC-ATGAGAGGATT-CTACTA-C
<i>Shalep9318</i>	AGGGA-AATGTAACATATATT-CTTGCCC-ATAAAGAGGAC-CCTACTAC
<i>Snigros6567</i>	??
<i>Tleuco6627</i>	??
<i>Tspic4846</i>	AGGGA-AATT-AACGGTTT--CTTCGTTTC-ATGAGATGATC-CTACTG-C
<i>Ttria6555</i>	AGGGA-A-CTGTCT-GTAT--CTTCGTTTCAT-GAGATGATC-CTACTA-C
<i>Zeadiplo</i>	??
<i>Zea</i>	GTAGA-AAATCTCT-AGAG-TAAGGATAAATC---TCTAC-ACACTT-C
<i>Sacharumofficinatum</i>	-GG-AATACTTTATTG-ATTTTTTAGTCCCTTTAATTGACA--TAGATGC
<i>Icylindrical</i>	-GG-AATACTTTATTG-ATTTTTTAGTCCCTTTAATTGACA--TAGATGC
<i>Dichanthiumannulatum</i>	??

<i>Aadscen6713</i>	????????????????
<i>Aameth6679</i>	GCGTCCCTTTCCA
<i>Aameth6685</i>	GCGACCTTTGTCA
<i>Aappen6543</i>	????????????????
<i>Aappen6946</i>	GTGTTCTTTGCCA
<i>Adist6684</i>	GTGTCCTTTGCCA
<i>Aeuco5869</i>	GTGTTCTTTCCCA
<i>Arundinellaneapalensis</i>	????????????????
<i>Aschiren6581</i>	GTGTTCTTTCCCA
<i>Bbladh</i>	ATGATCTTTGTCA
<i>Binsc5681</i>	GCGTCTTTGTCA
<i>BBrachiaria</i>	????????????????
<i>Bradi4784</i>	????????????????
<i>Bserra5074</i>	GCGTCTTTGCCA
<i>Bserra6654</i>	????????????????
<i>Bserra6954</i>	GTGTTCTTTGCC
<i>Cdieter6642</i>	GCGTCTTTCCCA
<i>Cexc6577</i>	GCGTCTTTCCCA
<i>Cleistachnesorghoides</i>	????????????????
<i>Cmarg4397</i>	????????????????
<i>Cplur6406</i>	CGGACCTTTGGTCA
<i>Cprolix6682</i>	G-GTCTTTCCCA
<i>Danthoniacalifornica</i>	????????????????
<i>Dfilli5872</i>	GCG-CCTTGGTCA
<i>Eragrostisdielsii</i>	????????????????
<i>Evill5085</i>	????????????????
<i>Hanam4807</i>	GTGTCCTTTGCCA
<i>Hcon6942</i>	GTGTTCTTTCCCC
<i>Hdiss5145a</i>	GCA-CCTTTGTCA
<i>Hemarthria9319</i>	GTGTTCTTTCTCA
<i>Hfilli5144</i>	GTAACCTATGTCC
<i>Hhirt6650</i>	????????????????



<i>Hhirt6656f</i>	GTGTCCTTTGCCA
<i>Hlongi6560</i>	?????????????
<i>Hpilg4603</i>	GCGTCCTTTGCCN
<i>Hpilg4738</i>	GTGTCCTTTGCCA
<i>Hquar5088</i>	GTGTCCTTTCCCA
<i>Hrud6576</i>	GTGTCCTTTGTCA
<i>Htamb6585</i>	GTGTCCTTTGTCA
<i>Icylin5114a</i>	GCGTCCTTTGCCA
<i>Icylin6632a</i>	GCGACCTTTGTCA
<i>Mcape6655</i>	G-GTCCTTTCCCC
<i>Monocymbium</i>	?????????????
<i>Pdilatum6617</i>	AGTACTCTACTAA
<i>Pentaschistisaristifol</i>	?????????????
<i>Pfrank4705</i>	GTGTCCTTTGCCA
<i>Pschinz6652</i>	?????????????
<i>Sbic6588</i>	?????????????
<i>Sbic6588</i>	GTGTCCTTTCCCA
<i>Shalep9318</i>	GGGTCCTTTCTCC
<i>Snigros6567</i>	?????????????
<i>Tleuco6627</i>	?????????????
<i>Tspic4846</i>	GTGTCCTTTGCCA
<i>Ttria6555</i>	GTGTCCTTTGCCA
<i>Zeadiplo</i>	?????????????
<i>Zea</i>	TTTTCTTTTAAA
<i>Sacharumofficinarum</i>	AAATACTTTCTAA
<i>Icylindrical</i>	AAATACTTTCTAA
<i>Dichanthiumannulatum</i>	?????????????