

## Selection of Elephant-Grass Genotypes for Forage Production

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Received: May 30, 2018

Accepted: August 12, 2018

Online Published: November 15, 2018

doi:10.5539/jas.v10n12p148

URL: <https://doi.org/10.5539/jas.v10n12p148>

### Abstract

The objective of this study was to evaluate the agronomic traits of 80 accessions of elephant grass under the soil and weather conditions of Campos dos Goytacazes/RJ, Brazil. The experimental design was set as randomized blocks with 2 replicates. The experiment continued from March 2012 to May 2013, with 5 harvests made in the dry and rainy seasons. The following traits were assessed: percentage of dry matter (%DM), dry matter yield (DMY), number of tillers per meter (NT), plant height (HGT), stem diameter (SD), leaf blade width (LBW) and leaf blade length (LBL). Data from each harvest were subjected to analysis of variance and to the Scott-Knott test ( $P < 0.05$ ). Tocher's optimization method, Mahalanobis distance, and canonical variables were utilized for the multiple traits, and the importance of the characters in the canonical variables. Genotypes with high yield were Elefante da Colômbia, Taiwan A-25, Albano, Hib. Gigante da Colômbia, Elefante de Pinda, Taiwan A-121, P241 Piracicaba, Guaçu/I.Z.2, CPAC, EMPASC 309, EMPASC 307, Australiano, and Pasto Panamá. Stem diameter (rainy season) and LBW (dry season) were the most important variables to differentiate between genotypes. There was wide phenotypic variation between genotypes, which could be divided into 15 groups by Tocher's optimization method.

**Keywords:** canonical variables, genetic divergence, *Pennisetum purpureum*

### 1. Introduction

Elephant grass (*Pennisetum purpureum* Schum.) is a tropical, perennial species with a high photosynthetic capacity. Because of its large genetic variability, it is able to adapt to the predominant climatic conditions throughout Brazil (Daher et al., 1997; Lista, 2008).

Introducing elephant grass germplasm has proved efficient in the identification of genotypes with desirable features (Shimoya et al., 2002; Lima et al., 2011) for future breeding programs. This demonstrates the importance of establishing and maintaining germplasm banks so as to guarantee the availability of the genetic variability necessary for breeding programs.

The elephant grass is a species of forage that has been highlighted by the nutritive balance and the healthy forage production per unit area (Ferreira et al., 2010). According to Santos et al. (2001) the protein contents in the elephant grass genotype, greater than 7%. Levels lower than 7% are limiting to animal production due to digestibility coefficients and negative nitrogen balance. The chemical composition of elephantgrass is not limiting the digestion of non-ruminal fiber, which ends up reflecting not the animal's weight gain (Garcia et al., 2011).

Thus, genetic breeding programs for elephant grass aim to select material superior to cultivars used at present and adapted to environmental conditions in the different regions of the country, followed by post-breeding studies such as research on diversity based on molecular markers (Azevedo et al., 2012; Lima et al., 2011) and based on discriminating morphological traits. These studies guide the crossings, with a view to maximizing yield and quality and also enable the indication of genotypes for cultivation in the regions of interest.

The objective of this study was to estimate the diversity among 80 accessions of elephant grass through morphological traits and to identify the elite genotypes, for possible use in future breeding programs.

## 2. Materials and Methods

### 2.1 Establishment and Development of the Experiment

The experiment was conducted on the Experimental Field of the State Center for Research on Agro-energy and Waste Use (Centro Estadual de Pesquisa em Agroenergia e Aproveitamento de Resíduos, Pesagro-RJ), located in Campos dos Goytacazes/RJ, Brazil ( $21^{\circ}19'23''$  S,  $41^{\circ}19'40''$  W; 13 m asl.). According to the Köppen classification, the climate in the region is an Aw type (hot, wet tropical), with annual precipitation around 1,150 mm. The soil was classified as a "Distrophic Argisol" (Embrapa, 2006), displaying the following properties: pH 5.5; phosphorus ( $\text{mg dm}^{-3}$ ) 18; potassium ( $\text{mg dm}^{-3}$ ) 83; Ca ( $\text{cmol}_c \text{dm}^{-3}$ ) 4.6; Mg ( $\text{cmol}_c \text{dm}^{-3}$ ) 3.0; Al ( $\text{cmol}_c \text{dm}^{-3}$ ) 0.1; H + Al ( $\text{cmol}_c \text{dm}^{-3}$ ) 4.5; and %C 1.6.

Eighty genotypes that belonged to the Active Germplasm Bank of Elephant Grass of Universidade Estadual do Norte Fluminense (UENF) were evaluated as shown in Table 1.

Table 1. Identification of 80 elephant grass genotypes (Campos dos Goytacazes, RJ, 2012/2013)

Identification	Genotypes	Identification	Genotypes
1	Elefante da Colômbia	41	Gramafante
2	Mercker	42	Roxo
3	Três Rios	43	Guaçu/I.Z.2
4	Napier Volta Grande	44	Cuba-115
5	Mercker Santa Rita	45	Cuba-116
6	Pusa Napier N 2	46	Cuba-169
7	Gigante de Pinda	47	King Grass
8	Napier N 2	48	Roxo Botucatu
9	Mercker S. E. A	49	Mineirão IPEACO
10	Taiwan A-148	50	Vruckwona Africano
11	Porto Rico 534-B	51	Cameroon
12	Taiwan A-25	52	CPAC
13	Albano	53	Guaçu
14	Hib. Gigante Colômbia	54	Napierzinho
15	Pusa Gigante Napier	55	IJ 7125 cv EMPASC 308
16	Elefante Hibrido 534-A	56	IJ 7126 cv EMPASC 310
17	Costa Rica	57	IJ 7127 cv EMPASC 309
18	Cubano Pinda	58	IJ 7136 cv EMPASC 307
19	Mercker Pinda	59	IJ 7139
20	Mercker Pinda México	60	IJ 7141 cv EMPASC 306
21	Mercker 86 México	61	Goiano
22	Taiwan A-144	62	CAC-262
23	Napier S.E.A.	63	Ibitinema
24	Pusa Napier N 1	64	903-77 ou Australiano
25	Elefante de Pinda	65	13 AD
26	Mole de Volta Grande	66	10 AD IRI
27	Napier	67	07 AD IRI
28	Mercker Comum	68	Pasto Panamá
29	Teresópolis	69	BAG-92
30	Taiwan A-46	70	09 AD IRI
31	Duro de Volta Grande	71	11 AD IRI
32	Turrialba	72	06 AD IRI
33	Taiwan A-146	73	01 AD IRI
34	Cameroon - Piracicaba	74	04 AD IRI
35	Taiwan A-121	75	13 AD IRI
36	Vruckwona	76	03 AD IRI
37	P241 Piracicaba	77	02 AD IRI
38	IAC-Campinas	78	08 AD IRI
39	Elefante C. Itapemirim	79	BAG UENF 79
40	Capim Cana D'África	80	BAG UENF 80

Planting took place in February 2011, and 2 standardization cuts were made: in December 2011 and March 2012. After the second cut, 5 harvests were made during the evaluation phase: 2 in the dry season (June and August 2012), and 3 in the rainy season (October 2012, February and May 2013). A randomized-block experimental design was used with 2 replications. Each plot was  $5.5 \times 2$  m, totaling 11 m<sup>2</sup>, with a floor area of 2 m<sup>2</sup>.

## 2.2 Evaluated Traits

At each harvest, the following morpho-agronomic traits were measured: plant height (HGT); stem diameter (SD); leaf blade width (LBW); and number of tillers (NT) per linear meter of row. To estimate dry matter yield (DMY) 2 plants from each plot were cut, placed in 5-kg paper bags and dried in a forced-ventilation oven at 65 °C for 72 h until constant weight (air-dry). The dried material was subsequently ground in a Wiley mill with 1 mm screen, and then 2 g of ground material were weighed and dried in an oven at 105 °C for 24 h (oven-dried sample). With the DMY data from the dry (Dry) and rainy (Rainy) seasons, it was possible to obtain the total dry matter yield (Total) and the percentage of total production from the dry season was calculated (%Dry).

## 2.3 Statistical Analyses

For the univariate analyses, an analysis of variance (ANOVA) was performed for each variable in each evaluation, according to the following model:  $Y_{ij} = M + G_i + B_j + e_{ij}$ , where,

$Y_{ij}$  represents the observation of the i-th group in the j-th block;

$M$  represents an overall constant associated with this random variable;

$G_i$  represents the effect of the i-th genotype;

$B_j$  represents the effect of the j-th block; and

$e_{ij}$  represents the experimental error associated with observation Y.

Subsequently, the means of the genotypes were grouped for each variable within each evaluation, using Scott and Knott's clustering. For the multivariate analyses, the canonical variables, relative importance of the traits in the canonical variables, generalized Mahalanobis distance (D2), and Tocher's clustering methods were employed using the GENES computer software (Cruz, 2013).

In the present study we addressed clustering through Tocher's optimization method by Mahalanobis distance (D2), which adopts the criterion that the average of the dissimilarity measures within each group should be lower than the distances between any groups. By the dissimilarity matrix we identified the pairs of most similar accessions, which formed the initial group, in which the possibility of inclusion of new accessions could be analyzed (Vasconcelos et al., 2011).

## 3. Results and Discussion

### 3.1 Dry Matter Yield and Percentage of Dry Matter

The overall mean for DMY in the 5 harvests was 23.92 t/ha, with 45.6% produced during the dry season, and the remaining 54.4% produced during the rainy season (Table 2). Working with 132 hybrids from crossings involving 12 cultivars of millet and 11 cultivars of elephant grass, Souza Sobrinho et al. (2005) obtained an average DMY of 7.45 t/ha in 8 harvests. In a study in the region of Campos dos Goytacazes/RJ (Brazil), Lista (2008) evaluated the potential of 10 genotypes and 2 cultivars of elephant grass under irrigation and found an average of 15 t/ha in 4 harvests, while Leão et al. (2012) studied the performance of 13 genotypes of elephant grass and obtained an average of 4.86 t/ha in 2 harvests, value below that found in the present study.

Based on high DMY in wet and dry seasons and overall, the genotypes identified as being in the elite group were (1) Elefante da Colômbia, (12) Taiwan A-25, (13) Albano, (14) Hib. Gigante Colômbia, (25) Elefante de Pinda, (35) Taiwan A-121, (37) P241 Piracicaba, (43) Guaçu/I.Z.2, (52) CPAC, (57) EMPASC 309, (58) EMPASC 307, (64) Australiano and (68) Pasto Panamá. However, other genotypes were equally good in either wet or dry season and overall so they should not be dismissed.

The genotypes (3) Três Rios, (7) Gigante de Pinda, (22) Taiwan A-144, (26) Mole de Volta Grande, (27) Napier, (28) Mercker Comum, (31) Duro de Volta Grande, (33) Taiwan A-146, (37) P241 Piracicaba, (38) IAC-Campinas, (39) Elefante C. Itapemirim, (41) Gramafante, (42) Roxo, (44) Cuba-115, (45) Cuba-116, (46) Cuba-169, (47) King Grass, (49) Mineirão IPEACO, (53) Guaçu, (54) Napierzinho, (55) IJ 7125 cv EMPASC 308, (59) IJ 7139, (61) Goiano, (62) CAC-262, (65) 13 AD, (67) 07 AD IRI, (69) BAG – 92, (70) 09 AD IRI, (71) 11 AD IRI, (73) 01 AD IRI, (74) 04 AD IRI, (77) 02 AD IRI, (78) 08 AD IRI showed higher DMY in the dry season, however, lower DMY in the rainy season compared to the other evaluated genotypes. According to Cunha et al. (2013) it is important to identify clones of elephant grass that show superiority in productive

performance and greater stability in forage production throughout the year, which in this case are part of the group of elite genotypes.

Table 2. Dry matter yields (DMY), in t/ha, of 80 genotypes of elephant grass in the dry and rainy seasons and total, plus percentage produced during the dry season (%Dry), in five harvests (Campos dos Goytacazes/RJ, Brazil, 2012/2013)

Genotype	Dry	Rainy	Total	%Dry	G <sup>T</sup>	Dry	Rainy	Total	%Dry
1	11.79 a <sup>1</sup>	19.61 a	31.40 a	37.59 b	41	16.89 a	12.26 b	29.15 a	58.27 a
2	8.93 b	11.26 b	20.19 b	43.46 b	42	11.53 a	8.52 b	20.06 b	57.50 a
3	12.28 a	9.76 b	22.04 b	55.87 a	43	14.66 a	19.05 a	33.71 a	45.55 a
4	8.22 b	7.99 b	16.22 b	50.34 a	44	15.20 a	11.52 b	26.73 a	56.68 a
5	7.31 b	17.46 a	24.77 a	30.35 b	45	10.86 a	10.76 b	21.62 b	50.92 a
6	6.58 b	9.00 b	15.58 b	42.99 b	46	12.63 a	11.44 b	24.06 a	53.22 a
7	14.93 a	12.38 b	27.32 a	52.65 a	47	11.80 a	13.97 b	25.78 a	47.65 a
8	62.17 b	12.95 b	19.17 b	32.45 b	48	7.96 b	10.90 b	18.86 b	42.08 b
9	8.01 b	13.43 b	21.44 b	37.08 b	49	12.05 a	12.79 b	24.83 a	48.16 a
10	8.61 b	17.53 a	26.15 a	32.75 b	50	9.91 b	17.76 a	27.67 a	35.22 b
11	8.59 b	7.80 b	16.39 b	52.53 a	51	9.39 b	9.87 b	19.26 b	4.932 a
12	11.28 a	14.73 a	26.01 a	42.06 b	52	12.53 a	14.93 a	27.46 a	45.43 a
13	10.87 a	14.91 a	25.78 a	42.06 b	53	11.68 a	10.60 b	22.29 b	51.72 a
14	11.19 a	17.18 a	28.37 a	40.33 b	54	11.23 a	10.37 b	21.60 b	52.60 a
15	9.69 b	12.37 b	22.06 b	43.75 a	55	11.60 a	14.03 b	25.63 a	45.31 a
16	9.65 b	11.86 b	21.51 b	45.07 a	56	8.06 b	15.15 a	23.21 b	35.16 b
17	7.55 b	12.37 b	19.92 b	37.76 b	57	14.95 a	16.44 a	31.39 a	47.69 a
18	6.01 b	6.50 b	12.51 b	48.05 a	58	16.64 a	18.67 a	35.31 a	47.05 a
19	5.90 b	11.83 b	17.74 b	33.29 b	59	12.95 a	11.61 b	24.56 a	53.17 a
20	5.58 b	10.54 b	16.12 b	34.10 b	60	9.41 b	10.93 b	20.34 b	46.16 a
21	8.661 b	13.93 b	22.59 b	38.17 b	61	11.63 a	13.04 b	24.68 a	47.64 a
22	11.77 a	13.63 b	25.40 a	46.61 a	62	13.81 a	13.61 b	27.42 a	50.35 a
23	6.35 b	16.45 a	22.80 b	28.20 b	63	9.65 b	13.89 b	23.55 b	41.48 b
24	9.29 b	12.35 b	21.64 b	44.41 a	64	14.89 a	14.58 a	29.48 a	50.20 a
25	14.95 a	16.00 a	30.95 a	47.42 a	65	12.55 a	12.64 b	25.18 a	43.07 b
26	12.29 a	12.21 b	24.49 a	48.44 a	66	10.02 b	12.84 b	22.86 b	52.91 a
27	10.96 a	10.07 b	21.03 b	49.79 a	67	12.19 a	11.02 b	23.21 b	47.58 a
28	11.40 a	9.07 b	20.47 b	55.45 a	68	20.05 a	22.14 a	42.19 a	56.57 a
29	8.66 b	15.74 a	24.40 a	34.99 b	69	14.03 a	10.67 b	24.70 a	50.44 a
30	8.46 b	10.33 b	18.79 b	46.04 a	70	13.75 a	134.73b	27.22 a	48.61 a
31	13.17 a	13.87 b	27.05 a	50.21 a	71	11.42 a	120.87b	23.51 b	30.78 b
32	10.59 b	14.38 a	24.99 a	42.29 b	72	5.85 b	13.11 b	18.96 b	55.72 a
33	12.89 a	11.56 b	24.46 a	52.39 a	73	12.04 a	96.93 b	21.73 b	48.65 a
34	8.77 b	10.30 b	19.07 b	45.98 a	74	12.06 a	12.75 b	24.81 a	39.81 b
35	11.79 a	14.71 a	26.49 a	44.57 a	75	8.71 b	13.02 b	21.73 b	29.33 b
36	7.56 b	7.84 b	15.39 b	48.94 a	76	7.77 b	18.88 a	26.66 a	51.03 a
37	12.78 a	17.92 b	30.70 a	41.13 b	77	15.68 a	13.37 b	29.05 a	52.86 a
38	13.34 a	12.70 b	26.05 a	51.27 a	78	15.03 a	12.39 b	27.41 a	41.13 b
39	13.33 a	13.25 b	26.58 a	49.14 a	79	7.84 b	11.31 b	19.15 b	44.25 a
40	9.58 b	10.63 b	20.22 b	47.40 a	80	9.98 b	12.46 b	22.44 b	50.79 a
Mean	10.96	12.96	23.92	45.62					
CV(%)	23.21	21.37	17.47	15.83					

Note. <sup>1</sup>Values within columns followed by the same letter do not differ by Scott-Knott's test at 5% probability.

The number of tillers per linear meter of row (NT) varied between 22.0 and 91.3 for genotypes Capim Cana D'África and BAG-92, respectively, and averaged 45.1 in the dry season. In the rainy season, NT varied from 26.1 to 72.6 for genotypes Goiano and EMPASC 309, respectively, averaging 46.0 (Table 2); the aforementioned values are higher than the 21.6 tillers found by Daher et al. (1997). Silva (2011) observed an average of 47.6 and Cunha (2012) obtained average values varying from 27.7 to 46.8 tillers. However, Daher et al. (2004) stressed that

this variable was the to the most useful for predicting dry matter yield potential, being positively correlated with this parameter.

Plant height averaged 1.43 and 1.22 m in the dry and rainy seasons, respectively (Table 3). Lower HGT values are found in studies with shorter cutting intervals (Cavalcante & Lira, 2010). Leão et al. (2012) found an average HGT for the genotypes of 1.57 m, which exceeds the value obtained in the current study.

Table 3. Total means for number of tillers (NT) per linear meter and plant height (HGT, m), of 80 genotypes of elephant grass in the dry and rainy seasons, in five harvests (Campos dos Goytacazes/RJ, Brazil, 2012/2013)

G <sup>1</sup>	NT		HGT		G <sup>1</sup>	NT		HGT	
	Dry	Rainy	Dry	Rainy		Dry	Rainy	Dry	Rainy
1	52.57	a <sup>1</sup>	52.10	a	1.57	a	1.41	a	41
2	36.32	b	39.42	a	1.30	b	1.22	a	42
3	51.61	a	52.25	a	1.21	b	1.13	b	43
4	51.25	a	38.65	a	1.13	b	1.03	b	44
5	43.25	b	60.83	a	1.26	b	1.26	a	45
6	44.67	b	35.39	a	1.22	b	1.22	a	46
7	75.45	a	45.45	a	1.30	b	1.20	b	47
8	35.25	b	50.19	a	1.22	b	1.26	a	48
9	36.00	b	47.29	a	1.26	b	1.25	a	49
10	42.75	b	47.17	a	1.44	a	1.28	a	50
11	45.17	b	48.00	a	1.48	a	1.13	b	51
12	47.09	b	46.06	a	1.33	b	1.25	a	52
13	25.31	b	28.97	a	1.61	a	1.39	a	53
14	45.20	b	50.14	a	1.50	a	1.32	a	54
15	37.75	b	43.46	a	1.39	b	1.39	a	55
16	28.14	b	33.56	a	1.30	b	1.30	a	56
17	26.50	b	45.37	a	1.17	b	1.21	a	57
18	24.71	b	37.97	a	1.27	b	1.29	a	58
19	43.75	b	40.06	a	1.11	b	1.22	a	59
20	32.25	b	33.79	a	1.22	b	1.17	b	60
21	41.00	b	41.05	a	1.22	b	1.17	b	61
22	42.25	b	45.33	a	1.52	a	1.25	a	62
23	40.25	b	49.24	a	1.43	a	1.25	a	63
24	36.75	b	31.89	a	1.34	b	1.21	b	64
25	57.48	a	40.56	a	1.37	b	1.29	a	65
26	60.75	a	69.45	a	1.25	b	1.23	a	66
27	50.51	a	56.56	a	1.17	b	1.15	b	67
28	51.59	a	56.22	a	1.17	b	1.05	b	68
29	44.00	b	62.83	a	1.58	a	1.16	b	69
30	42.25	b	47.53	a	1.60	a	1.23	a	70
31	45.13	b	49.88	a	1.52	a	1.17	b	71
32	33.44	b	35.62	a	1.53	a	1.32	a	72
33	63.51	a	43.47	a	1.30	b	1.14	b	73
34	49.35	a	40.14	a	1.50	a	1.31	a	74
35	53.28	a	38.83	a	1.06	b	1.10	b	75
36	31.31	b	27.79	a	1.33	b	1.28	a	76
37	32.07	b	62.00	a	1.31	b	1.04	b	77
38.	34.47	b	48.00	a	1.40	b	1.23	a	78
39	52.25	a	58.67	a	1.72	a	1.27	a	79
40	22.00	b	31.24	a	1.77	a	1.46	a	80
Mean Dry	45.09		1.43		Mean Rainy	46.02		1.22	

Note. <sup>1</sup>Values within columns followed by the same letter do not differ by Scott-Knott's test at 5% probability.

Stem diameter in the dry season averaged 1.06 cm and ranged from 0.70 cm for Mercker Comum to 1.45 cm for Pusa Napier and 1.43 cm for Albano. In the rainy season, the average stem diameter was 1.09 cm and again ranged from 0.67 cm for Mercker Comum to 1.45 cm for Albano and 1.43 cm for Pusa Napier (Table 4). Studying dwarf elephant grass, Silva et al. (2009) obtained an average of 1.05 cm; similarly Silva (2011) obtained 1.66 cm, and Cunha (2012) found higher stem diameters: 1.03 and 1.64 cm, demonstrating the large variability in this trait among the evaluated accessions. Rezende et al. (2008) evaluated Cameroon grass in the dry season, and obtained stem diameters varying from 0.54 to 1.51 cm, and 1.59 to 2.03 cm in the rainy season.

The average LBWs were 2.64 and 2.69 cm in the dry and rainy seasons, respectively (Table 4). Zhang et al. (2010) obtained an average of 3.33 cm, and Silva (2011) obtained 3.11 cm, which are higher values than those found in the present study. However, Silva et al. (2009) found an average value lower than 1.96 cm.

Table 4. Total means for stem diameter (SD, cm) and leaf blade width (LBW, cm) of 80 genotypes of elephant grass in the dry and rainy seasons, in five harvests (Campos dos Goytacazes/RJ, Brazil, 2012/2013)

G <sup>1</sup>	SD		LBW		G <sup>1</sup>	SD		LBW				
	Dry	Rainy	Dry	Rainy		Dry	Rainy	Dry	Rainy			
1	1.13	a	1.12	c	2.38	b	3.07	a	2.78	a	3.01	a
2	1.25	a	1.30	b	2.79	a	3.18	a	2.46	b	2.76	b
3	1.13	a	1.18	c	2.27	b	2.87	a	3.03	a	3.05	a
4	1.13	a	1.15	c	3.13	a	3.37	a	2.53	b	2.93	a
5	1.31	a	1.25	b	2.63	b	3.01	a	2.43	b	2.79	b
6	1.22	a	1.14	c	2.53	b	3.19	a	2.99	a	3.09	a
7	1.01	b	1.12	c	2.69	b	2.85	a	2.51	b	2.76	b
8	1.29	a	1.30	b	2.48	b	2.93	a	2.56	b	3.01	a
9	1.12	a	1.13	c	2.50	b	2.98	a	2.56	b	2.77	b
10	1.07	b	1.03	d	2.48	b	2.71	b	2.63	b	3.02	a
11	0.80	b	0.87	e	2.27	b	2.44	b	3.12	a	3.62	a
12	1.00	b	1.04	d	2.16	b	2.74	b	3.06	a	3.36	a
13	1.43	a	1.45	a	2.75	a	3.22	a	2.97	a	3.30	a
14	1.07	b	1.22	c	2.38	b	2.96	a	2.45	b	2.98	a
15	1.06	b	1.09	c	2.36	b	2.69	b	2.79	a	2.75	b
16	1.08	b	1.27	b	2.82	a	2.85	a	2.15	b	2.79	b
17	0.91	b	1.20	c	3.10	a	2.93	a	2.47	b	2.57	b
18	1.16	a	1.19	c	2.63	b	3.10	a	2.97	a	3.22	a
19	1.19	a	1.15	c	1.59	b	2.52	b	3.19	a	3.61	a
20	1.23	a	1.19	c	1.84	b	2.62	b	2.81	a	2.94	a
21	1.07	b	1.04	d	2.08	b	2.75	b	2.85	a	3.29	a
22	1.24	a	1.17	c	2.43	b	2.80	b	2.99	a	2.92	a
23	1.05	b	1.04	d	2.59	b	2.74	b	2.68	b	2.48	b
24	1.45	a	1.43	a	2.44	b	2.94	a	2.91	a	2.97	a
25	1.40	a	1.30	b	2.35	b	2.59	b	2.13	b	2.30	c
26	1.02	b	1.03	d	2.24	b	2.44	b	3.38	a	2.76	b
27	1.05	b	1.13	c	2.34	b	2.57	b	2.29	b	2.58	b
28	0.70	b	0.67	f	2.11	b	1.86	c	1.17	a	3.16	a
29	0.83	b	0.85	e	2.33	b	2.06	c	0.71	b	2.21	c
30	1.18	a	0.99	d	2.18	b	2.16	c	0.85	b	2.53	b
31	1.25	a	1.30	b	2.49	b	1.98	c	0.99	b	2.61	b
32	1.04	b	0.85	e	2.64	b	2.53	b	0.80	b	2.54	b
33	0.95	b	1.00	d	2.12	b	2.52	b	1.08	b	2.91	a
34	1.01	b	1.14	c	3.13	a	2.87	a	1.06	b	2.97	a
35	0.89	b	0.94	d	2.49	b	2.87	a	0.95	b	2.61	b
36	0.99	b	1.14	c	2.64	b	3.04	a	0.97	b	2.68	b
37	1.11	a	1.24	b	3.05	a	3.23	a	1.21	a	2.77	b
38.	1.03	b	0.97	d	3.14	a	3.27	a	0.96	b	2.55	b
39	0.95	b	0.90	e	2.68	b	2.70	b	0.87	b	2.85	a
40	1.05	b	1.11	c	3.20	a	3.22	a	3.35	a	3.08	a
Mean Dry	1.06		2.64		Mean Rainy	1.09		2	2.69			

Note. <sup>1</sup>Values followed by the same letter within columns do not differ by Scott-Knott's test at 5% probability.

### 3.2 Multivariate Analysis

When a set of traits represents qualitatively different variables and there is no correlation between them, univariate analysis is the most appropriate procedure to apply; however, when a dataset displays variables that are correlated, multinormality should be assumed, and a multivariate analysis of variance should be performed (Freitas et al., 2000).

For forage plants, a multivariate analysis is recommended, since traits usually display an appreciable level of correlation, being measured in the same tussock (Freitas et al., 2000).

When this analysis was performed on our data, the variance accumulated by the first two canonical variables was 66.8% of the total variance (Table 5), which made it possible to adopt this technique in the study of scattering of genotypes on a 2-dimensional graph. Evaluating the phenotypic divergence among 99 genotypes of elephant grass in Coronel Pacheco/MG (Brazil), Shimoya et al. (2002) observed that the first 2 canonical variables explained only 50.0% of the total variance. However, Daher et al. (1997) obtained a total of 80.7% of the variation in the first 2 canonical variables using 17 genotypes of elephant grass in Campos dos Goytacazes/RJ, and inferred that there was genetic divergence among the genotypes.

Table 5. Estimates of variances (eigenvalues,  $\lambda_j$ ) associated with the canonical variables, and respective weight coefficients (eigenvectors) of 8 variables assessed in 80 genotypes of elephant grass in Campos dos Goytacazes/RJ, Brazil

$\lambda_j$	Accumulated variance (%)	Variables								
		(1)	(2)	(3)	(4)	(5)	(6)	(7)	(8)	
7.358	45.488		-0.141	-0.039	-0.088	0.072	0.373	0.733	0.456	0.284
3.457	66.860		0.007	0.006	0.152	-0.221	-0.445	-0.313	0.644	0.466
1.752	77.693		-0.575	-0.007	0.533	0.148	-0.010	0.014	-0.397	0.454
1.263	85.504		0.176	0.079	0.620	0.452	-0.260	0.215	0.251	-0.443
0.946	91.349		0.404	-0.371	0.004	0.588	0.348	-0.329	0.026	0.351
0.604	95.085		-0.380	0.510	-0.444	0.576	-0.135	-0.167	0.139	-0.017
0.429	97.734		0.489	0.750	0.173	-0.099	0.159	0.069	-0.171	0.313
0.366	100.000		0.265	-0.178	-0.272	0.179	-0.656	0.419	-0.327	0.280

Note. 1 = NT in Dry, 2 = NT in Rainy; 3 = HGT in Dry, 4 = HGT in Rainy, in m; 5 = SD in Dry, 6 = SD in Rainy, in mm; 7 = LBW in Dry; and 8 = LBW in Rainy, in cm.

In the clustering analysis of the 80 genotypes, 15 Groups were formed with Group 1 containing 39 genotypes, Group 2 having 14 Genotypes, Group 6 having 7 genotypes and 7 of the remaining groups having only a single genotype (Table 6; Figure 1). The genotypes that comprised the elite group based on dry matter yields, described previously, were contained in Groups 1, 2, 3, 4 and 10.

Table 6. Clustering of 80 genotypes of elephant grass into groups by Tocher's optimization method based on generalized Mahalanobis distance ( $D^2$ ) (Campos dos Goytacazes/RJ, Brazil, 2012/2013)

Group	Genotypes
1	10 12 23 47 49 55 22 15 01 42 50 44 09 14 60 75 76 41 35 57 71 61 62 58 64 78 74 73 27 26 67 63 06 79 21 45 56 33 66
2	5 8 2 18 53 37 46 43 52 16 77 51 36 40
3	19 20 25
4	13 24
5	38 59
6	11 29 65 70 54 32 39
7	4 34 73
8	17 72
9	80
10	68
11	31
12	48
13	30
14	69
15	28

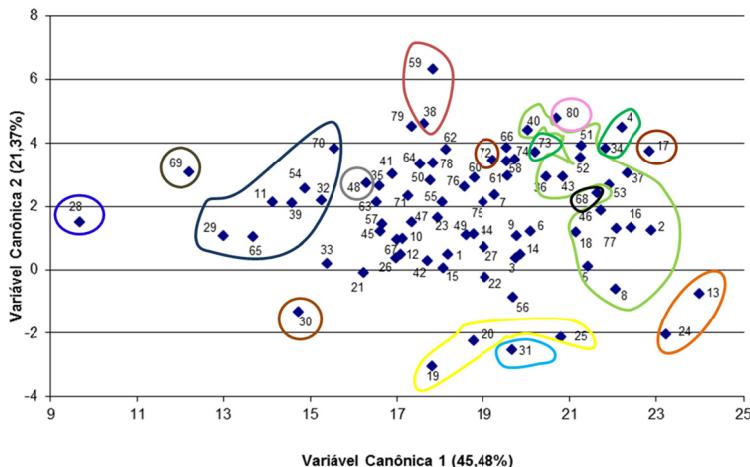


Figure 1. Scattering diagram of 80 genotypes of elephant grass, obtained by analysis of canonical variables.  
 Legend: G1 = no color; G2 = light green; G3 = yellow; G4 = orange; G5 = dark red; G6 = dark blue; G7 = dark green; G8 = dark brown; G9 = pink; G10 = black; G11 = light blue; G12 = light gray; G13 = light brown; G14 = dark gray; G15 = royal blue

#### 4. Conclusion

The genotypes that comprised the elite group were Elefante da Colômbia, Taiwan A-25, Albano, Híbrido Gigante da Colômbia, Elefante de Pinda, Taiwan A-121, P241 Piracicaba, Guaçu/I.Z.2, CPAC, EMPASC 309, EMPASC 307, Australiano, and Pasto Panamá. Future evaluations of the bromatological quality of these genotypes should be carried out.

Clustering analysis provided direction for crossings involving five groups; stem diameter (rainy season) and leaf blade width (dry season) were the most important variables to explain the scattering of genotypes.

#### Acknowledgments

The authors thank Fundação Carlos Chagas Filho de Amparo à Pesquisa do Estado do Rio de Janeiro (FAPERJ) for financing this research.

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