



## RESEARCH ARTICLE

### AGROMORPHOLOGICAL CHARACTERIZATION OF ACCESSIONS OF TRADITIONAL COTTON (GOSSYPIUM SPP) COLLECTED IN THE AGROECOLOGICAL ZONES PRODUCING COTTON IN BENIN

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#### ABSTRACT

Sixty-six traditional cotton accessions (*Gossypium* spp), collected in Benin, and were characterized by a set of 31 agromorphological traits established by UPOV and IPGRI. The evaluation was carried out during the 2011-2012 rainy season through a randomized complete block design at Cana CPE. Statistical analysis showed very highly significant differences between accessions for the 15 qualitative characteristics observed. The accessions Q111, Q122, Q95, Q93, Q135, Q92, Q71 and Q90 have characteristics of higher qualities. Analysis of the Pearson correlation matrix showed strong positive and negative correlations. The first two axes of the PCA absorb 73% of the total variability of the quantitative characters (54% for the vegetative development and 19% for the precocity). The first two factors of the CFL of qualitative traits account for 82% of inertia (70% for petal color, macula, pollen color and leaf shape, 12% for hairiness in lower part of The leaf, the denticulation of the bracts, the size of the capsules and the shape of the plant). The ascending hierarchical classification based on the Euclidean distance divided the accessions into six distinct groups comprising respectively 2, 14, 8, 11, 2 and 29 accessions. The results of this study provide information that can be used for the conservation and rational use of biodiversity for the improvement of commercial varieties.

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## INTRODUCTION

In Benin and most cotton-producing countries in West Africa, cotton is the main cash crop and plays an important social and economic role. The cotton *Gossypium*. Spp was cultivated in 2015 on an area of 33.9 million hectares (ha) with a production of 21.02 million tons of fiber, of which 8.6% of the area and 5.4% of the production of Fibers in French-speaking Africa (CICC, 2016). But there is a stagnation of seed cotton production around 300 000 tonnes in Benin over the last four cotton seasons with a low average yield of seed cotton. Genetic traits that make it possible to produce more for a lower cost are, first of all, those that remove the abiotic and biotic constraints that limit production (Mendez del Villar et al., 2006).

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Similarly, it is necessary to improve the agronomic, morphological and technological characteristics of the existing varieties which can meet the requirements of the actors of the cotton sector, based on the old varieties of cultivated plants, often referred to as "traditional", "Local" or "country" (Marchenay et al., 1987). Traditional varieties are also the basic material for today's biologists and breeders: they find the genes they will need tomorrow to better understand the nature of diversity and to use it more effectively (Marchenay et al., 1987). Traditional varieties must be maintained: so that genetics and breeding can continue to create new varieties that will allow agriculture to adapt to an uncertain future (Marchenay et al., 1987). Traditional varieties have long been modeled in each region by farmers in order to adapt them to their needs, as well as to selection pressures related to cultivation techniques (Sié, 1998, Harlan, 1987). These varieties thus constitute an important reserve of variability for breeders, which justify their systematic collection (Cauderon, 1986) and their evaluation (Sié, 1998).

The anthropogenic pressure exerted on the different parts of the traditional cotton tree, especially the roots and the new cultivation techniques practiced by the producers, can lead in time to the loss of the variability existing in these accessions. Indeed, if the genetic variability is the raw material of the breeder, it can be kept efficiently only after a precise knowledge of it. Not only does this knowledge allow good preservation; but also the diversity available for improvement and the genetic structure of the forms preserved in situ. There are studies of genetic variability in India, China, the United States and Brazil, where traditional cotton is widely used, but under very different agro-climatic and socio-economic conditions. In West and Central Africa, some studies have characterized and valued the genetic potential of local cotton farmers (Lhuillier, 1950, Franquin, 1954, Fournier and Gurknecht, 1991, Seignobos and Schwendiman, 1991). In Benin, on the other hand, there is no collection of traditional cotton genotypes that would allow conservation and rational use. In the absence of such a collection, it is difficult to design a cotton plant breeding program that can meet the needs of a country subjected to harsh abiotic and biotic conditions. In Benin, no studies on the characterization of traditional cotton have been carried out. This is why any contribution to the maintenance and valorization of genetic resources (Franckel *et al.*, 1995) must precede a study of its genetic diversity. This is the starting point of any program to improve and produce new varieties of cotton. Therefore, the objectives of the present investigation are to assess the structure and agromorphological variability of traditional cotton accessions collected in Benin by using morphological markers.

## MATERIALS AND METHODS

### Vegetable material

Traditional cotton samples from collections in the Guineo-Congolese, Guineo-Sudanese and Sudanian zones (Sinha *et al.*, 2016) have been used. Accessions of traditional non-germinated cotton or without sufficient seeds have been eliminated. Sixty-six (66) accessions collected in farmers' fields, in houses, in forests and along roadsides were characterized.

### Description of the experimental site and technical itinerary

Agromorphological characterization was carried out at the Permanent Center for Experimentation (CPE) in Cana (7 ° 6'11 " East, 2 ° 5'17 " North, and 89 m altitude) during the 2011 - 2012 rainy season. The CPE is characterized by a ferrallitic soil, denatured, developed on clay-sandy sediments of the continental terminal (bar land) and a climate of Guinean type with a bimodal regime with two rainy seasons intercalated by two dry seasons (mean annual rainfall was 1,300 mm in 2012 against 1,043 mm in 2011). Cotton accessions were characterized in a randomized complete block design without replication. Plots were single rows, 5m in length and 1m apart with 0.5m plant spacing. Each accession was sown 10 plants per plot. The seeds were soaked in water 24 hours before sowing. Soil preparation was manual (0.20 to 0.25 m depth). The accessions were numbered from Q61 to Q140. Technical routes have been monitored according to research recommendations, but other cropping factors are optimized, particularly mineral fertilization, weed control and insect pest control. The mineral fertilizer was based on 250 kg of

N14P23K14S5B1 brought on the 15th day after sowing (das) followed by a 50 kg / ha supply of urea and an additional 50 kg of KCl at the 40th das. Six weedings and a hoeing in the 50th das enabled the effective control of the weeds. Insecticide protection reinforced by systematic weekly sprays starting from the 50th das to the harvest in order to effectively control the parasitic pressure of the insects; The first four insecticide sprays were based on Thian; The following sprays combined a pyrethroid product with an acaricid or aphicid organophosphorus product depending on whether it was at mid-bloom or at the end of bloom. The marking of 4 plants per accession was carried out at random in the 90th das, with woolen strings and numbered from 1 to 4 on each row.

### Quantitative agromorphological variables observed

Fifteen (15) quantitative agromorphological characters selected from the list of cotton descriptors (UPOV, 2001) and illustrated by Harem *et al.* (2012) made it possible to characterize the various accessions. The following quantitative agromorphological variables were measured: mean date of first flower appearance (MDFFA), average opening date of the first boll (AODFB), ginning outturn (GO), weight of 100 seeds (g) or Seed Index (SI), height of the main stem (HT), insertion height of the first fruiting branch (IHFFB), length of the longest vegetative branch (LVB), length of the longest fruiting branch (LFB), number of nodes on the main stem (NN), number of insertion nodes of the first fruiting branch (NNFB), number of fruiting branches (NFB), number of nodes on vegetative branches (NVB), percentage of bolls on fruiting branches (% BFB) and percentage of bolls on vegetative branches (% BVF).

### Statistical analysis of data

The data obtained was processed using the Microsoft Excel 2010 spreadsheet and analyzed for variance analysis. Minimums, maxima, averages, standard deviations and coefficients of variation were then determined. The assessment of the structuring of agromorphological diversity was carried out by a Principal Component Analysis (PCA), an Ascending Hierarchical Classification (HAC) and a Discriminant Factorial Analysis (AFD) using the software STATISTICA version 6.1. Heritability ( $h^2$ ) of morphological characters was also estimated in order to assess the proportion of the phenotypic variability of genetic origin which is heritable and fixable in whole or in part.

## RESULTS AND DISCUSSION

### Variability of quantitative characteristics

The analytical characteristics of the variance such as minima, maxima, averages, standard deviations, coefficients of variation and Fisher's F-meanings are presented in Table 1. The analysis in the table highlights differences very significant between the various accessions for all the characters ( $P < 0.0001$ ). Significant differences were found between the minimum and maximum values of the 15 quantitative variables. The total height of the plant (HT) ranged from 105 to  $305 \pm 20.4$  cm. The insertion height of the first fruiting branch (IHFFB) ranged from 0 to 98 cm. The length of the vegetative branches (LVB) ranged from 32 to 160 cm. The length of the fruiting branches (LFB) ranged from 0 to 90 cm.

**Table 1. Results of the variance analysis realized on the measured variables**

Variables	Mean	Min	Max	Stan.Dev	CV%	h <sup>2</sup>	F	P<F
HT	198	105	305	20,4	14,6	>0,80	12,7	***
HPBF	24	0	130	4,2	24,1	-	105,8	***
LBV	96	32	160	17,6	25,9	>0,80	4,0	***
LBF	27	0	90	6,2	32,0	-	44,9	***
HNR	5	3	12	1,3	34,5	0,55	2,3	***
NN	40	23	67	4,6	16,4	>0,80	8,6	***
NPBF	7	0	25	1,0	20,2	-	122,6	***
NBF	12	0	38	2,4	28,5	-	52,9	***
NBV	19	1	43	2,5	18,8	>0,80	46,0	***
CBF	36	0	94	8,5	33,0	-	36,4	***
CBV	64	6	100	8,5	18,9	-	36,5	***
RE	25	18	43	2,2	12,0	>0,80	19,4	***
SI	7	5	11	0,6	12,3	>0,80	10,2	***
PFM	112	65	143	3,1	4,0	>0,80	99,9	***
OPCM	157	112	193	2,7	2,4	>0,80	146,7	***

P:\*\*\* Very highly significant ( $p < 0.001$ ); CV%: Coefficient of variation expressed as a percentage; Min: minimum; Max: maximum; HT (cm): height of a seedling, LBV (cm): length of vegetative branch, LBF (cm): fruit-bearing length of branch, NN: numbers nodes, HNR (cm): height of internode, HPBF (cm): height of insertion of the first fruit-bearing branch, NPBF: node of insertion of the first fruit-bearing branch, NBF: numbers fruit-bearing branches, NBV: numbers vegetative branches, % CBF: percentage of bolls collected on the fruit-bearing branches, %CBV: percentage of bolls collected on the vegetative branches, RE: Output with shelling, IF: seed index, PFM: goes back to appearance average of the flowers, OPCM: goes back to average opening of the bolls

**Table 2. Stamp correlation of Pearson of the qualitative characters of cotton plant**

Variables	HT	HPBF	LBV	LBF	HNR	NN	NPBF	NBF	NBV	%CBF	%CBV	RE	SI	PFM	OPCM
HT	1														
HPBF	-0,35	1													
LBV	0,39	-0,06	1												
LBF	-0,49	0,48	-0,03	1											
HNR	0,40	-0,34	0,12	-0,26	1										
NN	0,59	0,12	0,24	-0,14	-0,33	1									
NPBF	-0,43	0,94	-0,08	0,57	-0,38	0,09	1								
NBF	-0,42	0,62	-0,05	0,85	-0,34	0,05	0,70	1							
NBV	0,69	-0,49	0,21	-0,82	0,28	0,34	-0,58	-0,80	1						
%CBF	-0,55	0,61	-0,16	0,81	-0,33	-0,13	0,70	0,85	-0,84	1					
%CBV	0,55	-0,61	0,16	-0,81	0,33	0,13	-0,70	-0,85	0,84	-1,00	1				
RE	-0,44	0,05	-0,16	0,49	-0,05	-0,44	0,06	0,31	-0,54	0,40	-0,40	1			
SI	-0,40	0,06	-0,17	0,46	-0,07	-0,37	0,12	0,37	-0,48	0,46	-0,46	0,38	1		
PFM	0,47	-0,05	-0,03	-0,72	0,06	0,40	-0,15	-0,58	0,73	-0,62	0,62	-0,59	-0,50	1	
OPCM	0,46	0,02	0,00	-0,64	0,03	0,42	-0,09	-0,49	0,65	-0,54	0,54	-0,54	-0,43	0,87	1

Values in fat are significantly different from 0 on a level of significance  $\alpha=0,01$  (Pearson correlation), HT (cm): height of a seedling, LBV (cm): length of vegetative branch, LBF (cm): fruit-bearing length of branch, NN: numbers nodes, HNR (cm): height of internode, HPBF (cm): height of insertion of the first fruit-bearing branch, NPBF: node of insertion of the first fruit-bearing branch, NBF: numbers fruit-bearing branches, NBV: numbers vegetative branches, % CBF: percentage of bolls collected on the fruit-bearing branches, %CBV: percentage of bolls collected on the vegetative branches, RE: Output with shelling, IF: seed index, PFM: goes back to appearance average of the flowers, OPCM: goes back to average opening of the bolls

**Table 3. Eigen values and contribution of the characters to axes 1 and 2 of the CPA**

	PC1	PC2
explicative variance (%)	54	19
HT	-0,27	-0,12
HPBF	0,23	-0,37
LBV	-0,08	-0,06
LBF	0,32	-0,02
HNR	-0,18	0,34
NN	-0,12	-0,45
NPBF	0,24	-0,38
NBF	0,31	-0,18
NBV	-0,34	-0,03
%CBF	0,34	-0,10
%CBV	-0,34	0,10
RE (%)	0,21	0,32
SI (g)	0,22	0,25
PFM (jas)	-0,27	-0,28
OPCM (jas)	-0,24	-0,31

HT (cm): height of a seedling, LBV (cm): length of vegetative branch, LBF (cm): fruit-bearing length of branch, NN: numbers nodes, HNR (cm): height of internode, HPBF (cm): height of insertion of the first fruit-bearing branch, NPBF: node of insertion of the first fruit-bearing branch, NBF: numbers fruit-bearing branches, NBV: numbers vegetative branches, % CBF: percentage of bolls collected on the fruit-bearing branches, %CBV: percentage of bolls collected on the vegetative branches, RE: Output with shelling, IF: seed index, PFM: goes back to appearance average of the flowers, OPCM: goes back to average opening of the bolls

The height of the internodes per plant (HIP) ranged from 3 to 12 cm. The total number of nodes (NN) ranged from 23 to 67. The number of insertion nodes of the first fruiting branch (NINFB) varies from 0 to 25. The number of fruiting branches (NFB) ranged from 0 to 38 fruiting branches. The number of vegetative branches (NVB) ranged from 1 to 43 vegetative branches. The percentage of capsules per plant harvested on the fruiting branches ranged from 0 to 94% and finally the percentage of bolls per plant harvested on the vegetative branches ranged from 6 to 100%. Coefficients of variation (CV%) indicate small variations ranged from 2.4 to 4% between accessions respectively for the average opening date of the first boll (AODFB) and the date of appearance of the first flower average (DAFFA). Conversely, there is a very large variation between the accessions for the height of the internodes on the main stem (HIMS), the percentage of the boll on the fruiting branches (%BFB), the length of the longest fruiting branch (LFB), the number of fruiting branches (NFB) and the length of the vegetative branches (LVB) (35%, 33%, 32%, 29% and 26% respectively).

branches (NBV) have very high heritability ( $h^2 > 0,80$ ) Height to node ratio (HNR) is average ( $h^2 = 0.55$ ). This would indicate that genotypic variances are superior to those of the environment. The variability observed in these characteristics between accessions is therefore due to a large part of the genotype. According to Copur, (2006), significant variability in morphological characteristics can be attributed to differences in the genetic make-up of accessions. Similarly, Mandal *et al.* (2008) note that a high value of heritability is due to additive effects that ensure that progeny selection would be effective for the trait (s). This makes it possible to say that the selection of cotton accessions studied could be effective from these variables. However, this variability could also be explained by the diverse geographical origin of accessions (Aghaee *et al.*, 2010) and by farmer practices in seed management (Adoukonou-Sagbadja *et al.*, 2007). Similarly, several authors have shown that farmers' practices in seed management, including the exchange of varieties among farmers, are the source of a significant diversity between

**Table 4. Characteristics of the 6 groups of the accessions of traditional cotton plants resulting from the CAH**

Variables	G I	G II	G III	G IV	G V	G VI	F	P
N		19	11	12	9	8	7	
HT	257,05±3,8 <sup>a</sup>	213,81±5 <sup>b</sup>	200,7±4,7 <sup>b</sup>	144,44±5,5 <sup>c</sup>	131,87±5,87 <sup>c</sup>	153,28±6,27 <sup>c</sup>	103,8	<0,000
HPBF	0±1,63 <sup>d</sup>	48,54±2,15 <sup>b</sup>	0±2,06 <sup>d</sup>	65,55±2,37 <sup>a</sup>	20,5±2,51 <sup>c</sup>	28,28±2,69 <sup>c</sup>	191,5	<0,000
LBV	110,3±3,2 <sup>a</sup>	103,45±4,3 <sup>ab</sup>	84,25±4,1 <sup>c</sup>	86,66±4,79 <sup>bc</sup>	64,12±5,08 <sup>d</sup>	116,85±5,43 <sup>a</sup>	17,51	<0,000
LBF	0±1,54 <sup>c</sup>	57,81±2,03 <sup>a</sup>	0±1,94 <sup>c</sup>	32,55±2,24 <sup>b</sup>	59±2,38 <sup>a</sup>	58,57±2,54 <sup>a</sup>	213	<0,000
HNR	6,26±0,21 <sup>a</sup>	4,36±0,28 <sup>cd</sup>	5,66±0,27 <sup>ab</sup>	3,88±0,31 <sup>d</sup>	4,87±0,33 <sup>bcd</sup>	5,42±0,35 <sup>abc</sup>	11,54	<0,000
NN	44±0,94 <sup>c</sup>	52±1,23 <sup>c</sup>	38,58±1,18 <sup>c</sup>	37,66±1,36 <sup>c</sup>	28±1,44 <sup>c</sup>	30,28±1,54 <sup>c</sup>	45,49	<0,000
NPBF	0±0,31 <sup>d</sup>	15,27±0,41 <sup>b</sup>	0±0,39 <sup>d</sup>	19,33±0,45 <sup>a</sup>	7,25±0,48 <sup>c</sup>	9,28±0,51 <sup>c</sup>	391,6	<0,000
NBF	0±0,51 <sup>d</sup>	28,54±0,67 <sup>a</sup>	0±0,64 <sup>d</sup>	15,44±0,73 <sup>c</sup>	20,87±0,78 <sup>b</sup>	21,85±0,83 <sup>b</sup>	363,1	<0,000
NBV	31,89±0,57 <sup>a</sup>	11,45±0,76 <sup>c</sup>	27,58±0,72 <sup>b</sup>	12,66±0,83 <sup>c</sup>	3,5±0,89 <sup>d</sup>	5,42±0,95 <sup>d</sup>	262	<0,000
CBF	0±1,55 <sup>c</sup>	75,09±2,04 <sup>a</sup>	0±1,96 <sup>c</sup>	55±2,26 <sup>b</sup>	80,37±2,40 <sup>a</sup>	63±2,56 <sup>b</sup>	351,5	<0,000
CBV	100±1,55 <sup>a</sup>	25,27±2,04 <sup>c</sup>	100±1,96 <sup>a</sup>	45,55±2,26 <sup>b</sup>	20±2,40 <sup>c</sup>	37,42±2,56 <sup>b</sup>	350	<0,000
RE	21,78±0,64 <sup>c</sup>	24,27±0,84 <sup>c</sup>	23,5±0,80 <sup>c</sup>	24,77±0,93 <sup>c</sup>	36,5±0,99 <sup>a</sup>	29,71±1,06 <sup>b</sup>	36,48	<0,000
SI	6,21±0,14 <sup>b</sup>	6,90±0,18 <sup>b</sup>	6,25±0,18 <sup>b</sup>	6,66±0,21 <sup>b</sup>	8,5±0,22 <sup>a</sup>	7,64±0,24 <sup>a</sup>	21,71	<0,000
PFM	124,5±1,3 <sup>ab</sup>	109,18±1,83 <sup>c</sup>	129,5±1,75 <sup>a</sup>	120,33±0,21 <sup>b</sup>	80,87±0,22 <sup>d</sup>	77,14±2,29 <sup>d</sup>	128,2	<0,000
OPCM	168,47±1,5 <sup>a</sup>	157,81±1,9 <sup>b</sup>	172,66±1,8 <sup>a</sup>	167,77±2,1 <sup>a</sup>	125±2,31 <sup>c</sup>	121,85±2,47 <sup>c</sup>	109	<0,000

G I:Group I;G II:Group II;G III: Group III; G IV:Group IV; G V: Group V; G VI:Group VI; N: number of individuals,  $p < 0.0001$ :very highly significant;the averages followed by the same letter, in the same line are not significantly different according to the test from Newman-Keuls (Comparison joint committee); HT (cm): height of a seedling, LBV (cm): length of vegetative branch, LBF (cm): fruit-bearing length of branch, NN: numbers nodes, HNR (cm): height of internode, HPBF (cm): height of insertion of the first fruit-bearing branch, NPBF: node of insertion of the first fruit-bearing branch, NBF: numbers fruit-bearing branches, NBV: numbers vegetative branches, % CBF: percentage of bolls collected on the fruit-bearing branches, %CBV: percentage of bolls collected on the vegetative branches, RE: Output with shelling, IF: seed index, PFM: goes back to appearance average of the flowers, OPCM: goes back to average opening of the bolls

**Table 5. Discriminating analysis based on the agromorphological characters**

Variables	$\lambda$ Wilk	Partiel $\lambda$	F	P	R	(R) <sup>2</sup>
HT	0,028	0,786	13,639	< 0,001	0,429	0,571
LBV	0,026	0,837	9,754	< 0,001	0,842	0,158
HNR	0,023	0,947	2,771	0,019	0,403	0,597
NN	0,029	0,756	16,164	< 0,001	0,342	0,658
NBV	0,047	0,462	58,213	< 0,001	0,782	0,218
RE	0,024	0,906	5,173	< 0,001	0,885	0,115
SI	0,023	0,962	1,957	0,086	0,920	0,080
PFM	0,024	0,910	4,958	< 0,001	0,558	0,442
OPCM	0,025	0,874	7,181	< 0,001	0,660	0,340

HT (cm): height of a seedling, LBV (cm): length of vegetative branch, LBF (cm): fruit-bearing length of branch, NN: numbers nodes, HNR (cm): height of internode, HPBF (cm): height of insertion of the first fruit-bearing branch, NPBF: node of insertion of the first fruit-bearing branch, NBF: numbers fruit-bearing branches, NBV: numbers vegetative branches, % CBF: percentage of bolls collected on the fruit-bearing branches, %CBV: percentage of bolls collected on the vegetative branches, RE: Output with shelling, IF: seed index, PFM: goes back to appearance average of the flowers, OPCM: goes back to average opening of the bolls

This indicates a very strong heterogeneity between the accessions for these five characters. Moreover, some characteristics such as mean date of first flower appearance (PFM), average opening date of the first capsule (OPCM), Seed index (SI), ginning yield (RE), height of main stem (HT), length of the longest vegetative branch (LBV), number of nodes on the main stem (NN) and number of vegetative

crop populations (Mckeye *et al.*, 2001 Delaunay *et al.*, 2008, Missihoun *et al.*, 2012). These agromorphological differences between accessions are often conceived and implemented by various evolutionary processes. Agroecosystems are likely to exert selection pressures on genotypes (Sadiki and Jarvis, 2005). This is also the case for anthropogenic pressures (Robert *et al.*, 2005).



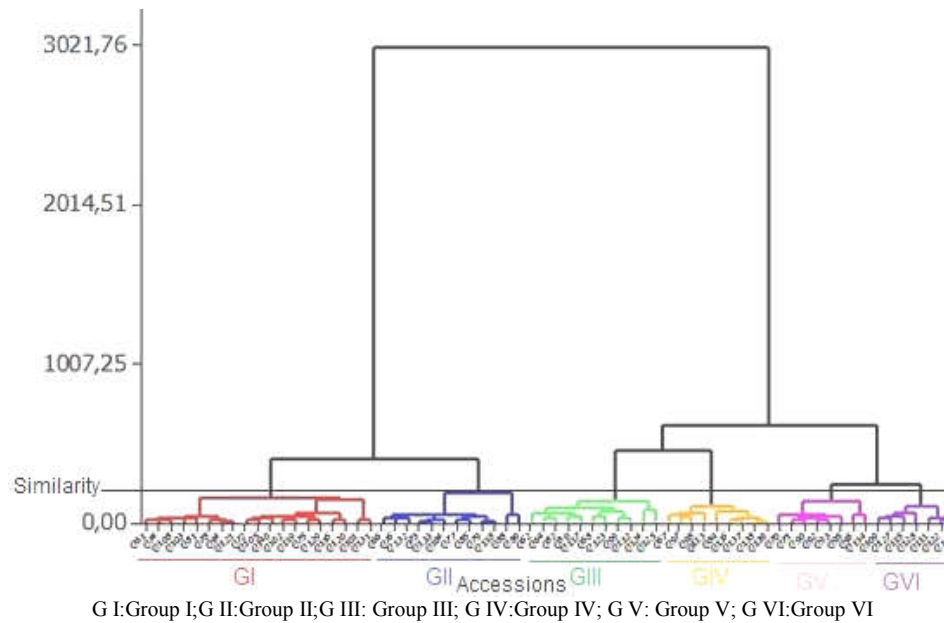


Figure 1. Ascending hierarchical clustering (CAH) of the 66 accessions of cotton plant by the Ward method

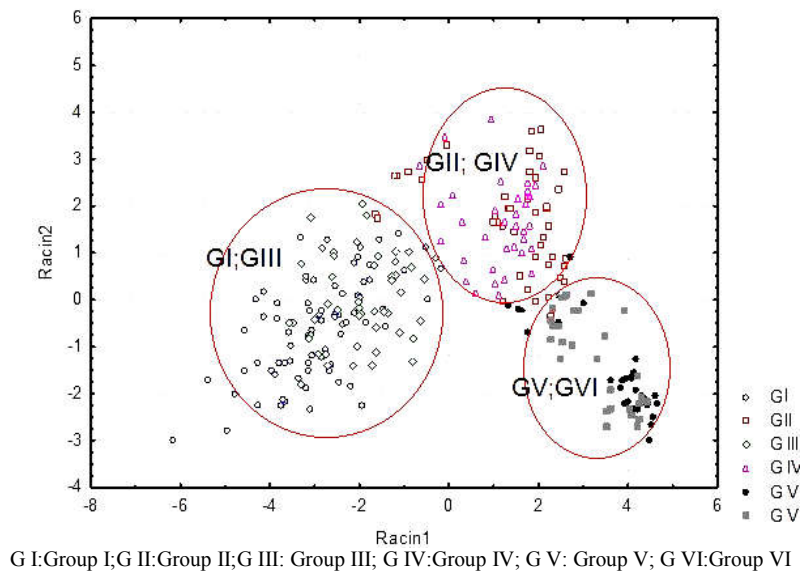


Figure 2. Representation of the various groups in the canonical plan discriminating formed by the canonical components 1 and 2

cotton plants (PFM  $77 \pm 2.29$  jas and OPCM  $122 \pm 2.47$  jas) of large size ( $153 \pm 6.27$  Cm), with very long vegetative branches ( $117 \pm 5.43$  cm) and very long fruiting branches ( $59 \pm 2.54$  cm). They have capsules on both the fruiting branches ( $63 \pm 2.56\%$ ) and the vegetative branches ( $37 \pm 2.56\%$ ). They have a large number of fruiting branches ( $22 \pm 0.83$ ), with very few vegetative branches ( $5 \pm 0.95$ ) for a high ginning yield ( $30 \pm 1.06\%$ ) and medium sized seeds ( $8 \pm 0.24$  g). Results showed that most of cotton accessions are considered vegetative and considered indeterminate to long cycle. Nevertheless, those of groups V and VI are very early. Cotton earliness is a quantitative characteristic that is mainly affected by the environment and crop genotype (Kassianenko *et al.*, 2003). Therefore, in any cropping system, selection of early cultivars is the main factor (Nichols *et al.*, 2004). The advantage of earliness of cotton varieties has been reported in Pakistan and other countries for the timely rotation of other crops for the wheat-cotton-wheat cropping system (Ali *et al.*, 2003). Precociousness is a trait sought by breeders and growers to escape unfavorable weather conditions, to avoid yield losses from disease and pest complexes (Singh, 2004). According to

José Martin (2001), with an equal productivity, an early cultivar has lower requirements for seasonal length but increased during the season for water and mineral feeding and protection. Since production is more concentrated over time, nutrient flows must be higher, and any biotic or abiotic stress occurring in a given period will have a greater impact. Moreover, the accessions of groups I, II, III and IV being very vegetative and late, would have advantages when the abiotic conditions (compaction of soil, salinity, water stress) are unfavorable and disadvantages in case of excessive irrigation Or of density too high according to Bassett and Kerby (1996). Thus, Hearn (1995) recalls that long-cycle varieties (indeterminate) are more suited to drought (different water stress) than the varieties determined. Then, long cycle accessions can be used for areas of water deficit. According to Lançon *et al.* (2000), these criteria all have sufficient variability to be combined in specific selection strategies of the type sought. This makes these accessions a material of choice in cotton improvement and extension programs in Benin and in Africa.

Table 6. Coefficient of canonical function

Axes	F 1	F 2
Eigen value	6,594	1,472
Percentage of cumulated inertia	0,713	0,872
HT	-0,450	-0,522
LBV	0,056	0,266
HNR	0,134	0,411
NN	0,349	0,869
NBV	-0,746	-0,562
RE	0,050	-0,161
SI	0,132	-0,084
PFM	-0,121	0,427
OPCM	-0,221	0,460

HT (cm): height of a seedling, LBV (cm): length of vegetative branch, LBF (cm): fruit-bearing length of branch, NN: numbers nodes, HNR (cm): height of internode, HPBF (cm): height of insertion of the first fruit-bearing branch, NPBF: node of insertion of the first fruit-bearing branch, NBF: numbers fruit-bearing branches, NBV: numbers vegetative branches, % CBF: percentage of bolls collected on the fruit-bearing branches, %CBV: percentage of bolls collected on the vegetative branches, RE: Output with shelling, IF: seed index, PFM: goes back to appearance average of the flowers, OPCM: goes back to average opening of the bolls

### Discriminant analysis

The discriminant analysis was carried out by using as a categorical variable the 6 groups obtained in the ascending hierarchical classification (CHA) to find the most discriminating variables with respect to the determined groups. The variables most contributing to the observed morphological variability are the height of the main stem (HT), length of the longest vegetative branch (LVB), number of nodes on the main stem (NN), number of vegetative branches (NVB), ginning outturn (GO), average date of the first flower (MDFFA) Opening of the first boll (AODFB). The Centered-Reduced Coefficients of the Canonical Discriminant Function are presented in Table 6. The first canonical function is the most important and explains more than 71% of the total variance and the second explains 5.9% of the total variance (Table 6). The analysis of this table also showed that the number of vegetative branches (NVB) is relatively well correlated on one side as on the other of axis 1. The characters height of the main stem (HT), height (HIP), number of nodes on the main stem (NN), average date of appearance of the first flower (MDFFA) and average opening date of the first boll (AODFB) Axis 2. Moreover, the graphical representation of discriminating canonical analysis (Figure 2) presents three major groups. It shows a grouping between the accessions of groups I and III, II and IV then V and VI.

### Conclusion

The results on the diversity and agromorphological structure of the 66 cotton accessions in Benin on the basis of the 16 characters clearly showed that the accessions analyzed present an important variation for all the characters used, in particular those related to the fruiting branch, The vegetative branch, the insertion height of the first fruiting branch, the internode height and the ginning yield. This variability observed between accessions constitutes a real reservoir of genetic diversity, an argument of choice for the conservation of these cotton accessions. Characterization based on agromorphological criteria is certainly of great value for an efficient valorization, but should be supplemented by other more in-depth studies, in particular molecular techniques such as microsatellites, which will make it possible to better characterize the accessions inside Different groups. Finally, the introduction of these accessions into genetic reserves and their maintenance in situ

will contribute to the preservation of genetic integrity and the adaptability of the material in order to avoid a scientific catastrophe.

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