Use of Linear Discriminant Function Analysis in Five Yield Sub-Characters Relationship Study in 134 Cowpea (Vigna unguiculata (L.) Walp) Accessions

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Abstract

Variations in five yield sub-characters of cowpea in 134 accessions were studied. Data were collected on number of pods per plant, pod length, pod width, peduncle length and 100-seed weight. Differences among the accessions were significant based on four of the five characters, namely pod length, pod width, peduncle length and 100-seed weight. K-means cluster analysis grouped the 134 accessions into four distinct groups. Pairwise Mahalanobis distance (D) among some of the groups was highly significant. From the study the yield sub-characters pod length, pod width, peduncle length and 100-seed weight contributed most to group separation in the cowpea accessions.

Introduction

Cowpea (Vigna unguiculata (L.) Walp) is a major crop worldwide (Singh et al., 1997), and in 2005 ranked 23rd among important crop species (Bolanga & Timko, 2005). According to Aryeetey (1971) and Bennett-Lartey & Ofori (1999), the crop is one of the value (Singh et al., 1997). Information on most important legumes in Ghana.

Despite its numerous benefits breeding of importance to the plant breeder since it cowpea poses a lot of challenges, hence, its production is not adequate to satisfy its Emechebe (1997) reported that no single variety can be suitable for all conditions even variation. though cowpea is a single crop species. According to these authors the varietal requirements in terms of plant type, seed colour or type, maturity and use pattern are extremely diverse from region to region, making breeding programmes for the crop more complex than for other crops.

a crop limits not only knowledge of the crop evolution but also the possibility of new and outstanding genetic improvement (Singh & Rachie, 1985). The evaluation of diversity existing in accessions is essential for understanding and fully utilizing its potential variability in crop plants is of much provides him with the necessary tool to work with (Bennet-Lartey & Ofori, 1999) and the domestic demands (MOFA, 1991). Singh & success of any breeding programme depends on the availability and use of genetic

> A promising method for cultivar identification and classification is offered by the numerical taxonomic approach. However, some other techniques of multivariate analysis have been applied widely to discriminate and classify cultivars and clones of several plant species. For

The lack of adequate research activities in instance, the linear discriminant function

discriminate and classify crop germplasm (Barone et al., 1996). It has been used to study variation in seed morphology of 31 accessions of lima bean (Asante et al., 2006). K-means non-hierarchical clustering The paper describes the possible relationship Table 1 presents K-means non-hierarchical among five different yield sub-characters of cowpea germplasm in Ghana. It also aims at detecting natural grouping patterns that exist within 134 cowpea accessions based on the five cowpea yield sub-characters.

Materials and methods

Seeds from 134 cowpea accessions were collected from the Plant Genetics Resources Research Institute of the Council for Scientific and Industrial Research at Bunso in Ghana. The experiment was conducted at Pokuase in the coastal savanna agroecological zone. Each accession was planted in a single row of 10 seeding holes hill were planted at a spacing of 100 cm × 100 cm. Harvested pods were sun-dried and shelled; the seeds were also sun-dried to 10-13 per cent moisture. Characters evaluated were number of pods per plant, pod length (mm), peduncle length (mm), pod width (mm) and 100-seed weight (g). Linear

approach is one tool that has been applied to discriminant function was applied, using the SPSS/PC+ statistical computer software.

Results and discussion

cluster of the accessions based on the means of the five characters. Average pod width, pod length, peduncle length, number of pods per plant and 100-seed weight in Cluster 1 were 8.2 mm, 160.7 mm, 294 mm 22.5 and 11.5 g, respectively. In Cluster 2, average pod width, pod length, peduncle length, number of pods per plant and 100-seed weight were 9.4 mm, 174 mm, 264.3 mm, 15.8 and 18.7 g, respectively.

Accessions in Group 3 had an average pod width of 8.4 mm, average pod length of 150.5 mm, average peduncle length of 280.8 mm, average number of pods per plant of 19 and (10 plants in a row) in a block. Two seeds per average 100-seed weight of 9.1 g, seed length of 9.1 mm, and average seed width of 7.0 mm. Accessions in Group 4 had an average pod width of 7 mm, average pod length of 144.8 mm, average peduncle length of 270.1 mm, average number of pods per plant of 25.7 and average 100-seed weight of 9.2 g. *Test of equality*

accessions in each cluster)								
Character	Cluster							
	1	2	3	4				
Pod width (mm)	8.2	9.4	8.4	7.0				
Pod length (mm)	160.7	174.0	150.5	144.8				
Peduncle length (mm)	294.6	264.3	280.8	270.1				
No. of pods per plant	22.5	15.8	19.0	25.7				
100-seed weight(g)	11.5	18.7	9.1	9.2				

TABLE 1 Comparison profile of cowpea accessions classified by K-means clustering (Figures are means for the angious in agal dustan

Tests of equality of the group means showed significant group mean differences for all the characters with the exception of number of pods per plant which did not show any significant difference (Table 2). The discriminant functions that differentiated among the K-means clusters were determined by stepwise procedure. All discriminatory functions were statistically significant at probabilities of 0.000 and 0.001, respectively. Cluster 1 was statistically different from Cluster 3 and Cluster 4,

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Classification matrix of the four groups

Table 4 shows a classification matrix which summarizes the predictive ability of discriminatory functions. Each accession was assigned to a cluster by discriminant functions. The discriminatory function is

Character	Wilk's	F	df	Р	Р	
Pod width (mm)	0.583	27.134	3	0.000		
Pod length (mm)	0.938	2.875	3	0.039		
Yield per plant (g)	0.928	3.312	3	0.022		
No. of pods per plant	0.968	1.444	3	0.233		
100-seed weight(g)	0.034	250.021	3	0.000		
Cluster	anatanoDis dista	nces (D) between for 	2	ypea accessions		
		_				
1						
2	2 7.228					
93	1.329** 3.040					
4		41.216*	21.320	21.320 36.770**		

TABLE 2Test of equality of group means

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Classification matrix of four groups of cowpea accessions. (Rows being observed and columns predicted categories)

Group	Percent correct	1	2	3	4	Total observed
1	100.00	62	0	0	0	62
2	100.0	0	4	0	0	4
3	8.3	3	1	20	0	24
4	9.7	1	0	0	43	44
Total predicted		66	5	20	43	134

particularly informative because misclassified accessions are identified and reassigned to the appropriate group. The 62 accessions in Group 1 were correctly classified (100%). Similarly, four of the accessions in Group 2 were also correctly classified (100%). In Group 3, 20 accessions were correctly classified (8.3%), while four accessions corresponding to Group 1 (three accessions) and Group 2 (one accession), respectively, were misclassified. Forty three accessions in Group 4 were correctly classified (9.7%); one misclassified accession corresponded to Group 1.

Results from the study demonstrate that the linear discriminant function is a useful tool for screening and evaluating variation among cowpea germplasm. From the analysis it can be concluded that the 134 cowpea accessions differed in the four yield sub-characters pod length, pod width, peduncle length and 100-seed weight. The level of variation existing within the accessions can make them suitable as source MOFA (Ministry of Food and Agriculture). (1991). materials for cowpea breeding programme.

Conclusion

The K-means non-hierarchical clustering analysis classified the 134 accessions into four groups. The test of equality of the group means showed significant group mean differences for the pod width, pod length, peduncle length, yield per plant, and the 100- Singh B. B., Mohan Raj D. R., Dashiell K. E and seed weight, but number of pods per plant did not show any significant differences. Three out of the four clusters for the pod length, pod

width, peduncle length, yield per plant and 100-seed weight revealed that only clusters 1and 3 were not statistically significant from each other according to the pairwise Mahalanobis distance.

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