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Discriminant Function Method of Selection in Vegetable Cowpea Genotypes

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ABSTRACT

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Twenty two diverse genotypes of vegetable cowpea (*Vigna unguiculata* subsp. *Unguiculata* [L.] Verde.) were evaluated for yield and a few related characters in a field experiment in randomized block design with three replications during 2011-12 at College of Agriculture, Vellayani, Trivandrum, Kerala. Discriminant function technique was adopted for the construction of selection index for yield using fruit yield per plant (X_7) and the component characters viz., vine length (X_1), days to first flowering (X_2), pod length (X_3), pod girth (X_4), pod width (X_5) and number of pods per plant (X_6). These component characters showed relatively stronger association with yield and could form a valuable selection index for yield in this crop. Among Twenty two genotypes, VU 7 (1475.31) followed by VU 1 (1468.72) and VU14 (1426.92) recorded high selection index values and VU 3 (908.28), recorded low selection index value.

Introduction

Vegetable cowpea (*Vigna unguiculata* L. Walp) is one of the important food legumes and a valuable component of the traditional cropping systems in the semiarid tropics (Singh *et al.*, 2002; Ayisi *et al.*, 2000). The crop is adaptable to harsh environments and withstands extreme temperatures, water limiting conditions and poor soil fertility. It yields well in harsh environments where other food legumes do not thrive. Due to adaptation versatility, ability to fix atmospheric nitrogen and considerable level of seed protein, minerals and vitamin contents cowpea should significantly contribute as viable and alternative crop in low input farming systems.

The economic worth of a plant depends upon several characters so while selecting a desirable plant from a segregating population the plant breeder has to give due consideration to characters of economic importance. Selection index is one such method of selecting plants for crop improvement based on several characters of importance. This method was proposed by Smith (1947) using discriminant function of Fisher (1936). Smith suggested that a better way of exploiting genetic correlation with several traits having high heritability is to construct an index, called selection index, which combines information on all the

characters associated with the dependent variable like yield. Thus selection index refers to a linear combination of characters associated with yield. The best known selection indices involve discriminant functions based on the relative economic importance of various characters. The discriminant function analysis measures the efficiency of various character combinations in selection. Selection index leads to simultaneous manipulation of several characters for genetic improvement of economic yield. This technique provides information on yield components and thus aids in indirect selection for the genetic improvement.

Materials and Methods

The experiment was carried out at Department of Olericulture, College of Agriculture, Vellayani (8° 5' N latitude and 77° 1' E longitude at an altitude of 29 m above mean sea level) during 2011-2012 with 22 vegetable cowpea genotypes (Table 1). Soil type is a red loam belonging to Vellayani series, texturally classified as sandy clay loam. The trial was laid out in a randomized block design with three replications. Seeds were sown at a spacing of 45 X 30 cm. Timely crop management practices were adopted as recommended by Kerala Agricultural University (KAU, 2007) to raise healthy crop. Observations on growth, yield as well as yield contributing characters was noted on five randomly selected plants in each plot at different stages of the crop.

The selection index developed by Smith (1937) using discriminate function of Fisher (1936) was used to discriminate the genotypes based on all the characters.

The selection index is described by the function, $I = b_1 x_1 + b_2 x_2 + \dots + b_k x_k$ and the merit of a plant is described by the function, $H = a_1 G_1 + a_2 G_2 + \dots + a_k G_k$

where x_1, x_2, \dots, x_k are the phenotypic values and G_1, G_2, \dots, G_k are the genotypic values of the plants with respect to characters, x_1, x_2, \dots, x_k and H is the genetic worth of the plant. It is assumed that the economic weight assigned to each character is equal to unity i. e., $a_1, a_2, \dots, a_k = 1$.

The regression coefficients (b) are determined such that the correlation between H and I is maximum. The procedure will reduce to an equation of the form, $b = P^{-1}Ga$ where, P is the phenotypic variance-covariance matrix and G is the genotypic variance-covariance matrix x .

Results and Discussion

Discriminant function technique was adopted for the construction of selection index for yield using fruit yield per plant (X_7) and the component characters *viz.*, vine length (X_1), days to first flowering (X_2), pod length (X_3), pod girth (X_4), pod width (X_5) and number of pods per plant (X_6). These component characters showed relatively stronger association with yield and could form a valuable selection index for yield in this crop.

The selection index, worked out in the present study is given below.

$$I = 0.9566 X_1 + 0.9798 X_2 + 1.1151 X_3 + 1.9063 X_4 + 0.4786 X_5 + 0.6706 X_6 + 0.9003 X_7$$

The selection index value for each genotype was determined and they were ranked accordingly (Table 2). Five genotypes *viz.*, VU 7 (1475.31), VU 1 (1468.72), VU14 (1426.92), VU 16 (1371.64) and VU 6 (1351.56) recorded high selection index values. Five genotypes *viz.*, VU 3 (908.28), VU 24(939.50), VU 11 (952.97), VU 21 (1007.05), VU 10 (1148.04) recorded low selection index values.

Identification of superior genotypes of vegetable cowpea based on discriminant function analysis was also done some workers.

In yard long bean, Resmi (1998) worked out the selection indices using thirteen characters and found that the genotype VS 6 had the

maximum index value followed by VS 11. Superior genotypes were identified by constructing selection indices using the characters namely vine length, primary branches, petiole length, length and breadth of lateral leaflets, days to flowering, pod length, pod girth, pod weight, pods per inflorescence, pods per kilogram, pods per plant and yield.

Table.1 List of bush cowpea accessions used for the evaluation

Sl. No.	Genotype Number	Genotype Name	Source
1	VU 1	Anaswara	College of Horticulture, Vellanikkara
2	VU 2	Local	Kollam
3	VU 3	Bhagyalakshmi	College of Agriculture, Vellayani
4	VU 4	Local	Haritha Agrofarm, Trivandrum
5	VU 5	Arka Suman	IIHR, Bangalore
6	VU 6	Arka Garima	IIHR, Bangalore
7	VU 7	Local	Thodupuzha
8	VU 8	Pusa Phalguni	IARI, New Delhi
9	VU 9	Co-26	TNAU, Coimbatore
10	VU 10	Kanakamony	RARS, Pattambi, Palakkad
11	VU 11	Local	Kottamangalam, Ernakulam
12	VU 13	Local	College of Agriculture, Vellayani
13	VU 14	Local	College of Agriculture, Vellayani
14	VU 15	GC-9732	RARS, Pattombi, Palakkad
15	VU 16	CO-2	TNAU, Coimbatore
16	VU 17	Local	Brahmamangalam, Kottayam
17	VU 18	Local	College of Agriculture, Vellayani
18	VU 19	Local	Thrippunithura, Ernakulam
19	VU 20	Local	College of Agriculture, Vellayani
20	VU 21	GC-3	College of Agriculture, Vellayani
21	VU 22	Local	Pilicode, Kasargode
22	VU 24	Local	College of Horticulture, Vellanikkara

Table.2 Vegetable cowpea genotypes ranked according to selection index

Accession	Index	Ranks in ascending order
VU 7	1475.31	1
VU 1	1468.72	2
VU 14	1426.92	3
VU 16	1371.64	4
VU 6	1351.56	5
VU 22	1350.27	6
VU 17	1333.83	7
VU 2	1290.76	8
VU 8	1265.53	9
VU 4	1257.67	10
VU 18	1241.98	11
VU 13	1207.84	12
VU 19	1198.24	13
VU 5	1196.97	14
VU 9	1184.4	15
VU 15	1158.91	16
VU 20	1152.57	17
VU 10	1148.04	18
VU 21	1007.05	19
VU 11	952.97	20
VU 24	939.5	21
VU 3	908.28	22

(Based on discriminant function analysis)

Philip (2004) worked out selection indices for 50 genotypes of cowpea on the basis of pods per plant, number of inflorescence per plant, pods per inflorescence, pod length, seeds per pod and 100 seed weight.

Five superior genotypes were selected for hybridization programme as female parents to develop F₁ hybrids.

Selection index for the genotype was computed based on the nine characters having significant genotypic correlation coefficients namely pods per cluster, pods per plant, pod yield per plant, pod weight, pod length, pod breadth, seeds per pod, length of harvest period and main stem length. The maximum selection index value was obtained for VS 41,

while the least value was for VS 7 by Lovely (2005).

Manju (2006) observed selection indices involving the characters, peduncle length, pod length, pod girth, pod weight, pods per plant, seeds per pod, 100 seed weight, number of harvests, pod protein and yield per plant. Based on selection index, VS 27 ranked first followed by VS 8 and VS 19.

Discriminant function technique involves development of selection criteria on a combination of various characters and aids the breeder in indirect selection for genetic improvement in yield. In plant breeding, selection index refers to a linear combination of characters associated with yield.

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