

## SELECTION INDICES IN TABLE PEAS (*PISUM SATIVUM* LINN.)

R. K. MITAL AND P. S. VERMA

*Department of Genetics and Plant Breeding, C. S. Azad University of Agriculture and Technology, Kanpur 208002*

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### ABSTRACT

Discriminant function analysis for yield revealed a progressive increase in the relative efficiency based on two, three, four or five character combinations over straight selection or single character selection indices. The maximum efficiency was observed with the selection indices based on five characters, viz., days to flower + pod length + number of seeds/pod + pod weight + test weight of 100 green grains. No. of seeds/pod, pod weight and test weight of 100 green grains were the important components with direct influence on yield on which maximum emphasis should be laid while making selection for yield.

**Key words:** Discriminant function, selection indices, relative efficiency, table pea.

The breeder is seldom faced with a situation in which modification of only a single attribute is desired. Normally modification are desired in several attributes. The theory of selection indices is proposed for manipulating several attributes simultaneously [1-3]. Ever since, a large number of selection indices were worked out in different crops. The use of this technique is important in selecting genotypes and also characters of specific importance. The present study, therefore, envisages to construct different functions for improvement of green pod yield in table peas (*Pisum sativum* L.).

### MATERIALS AND METHODS

Twenty new selections and cultivars of table peas were grown in randomized block design with 3 replications at the Vegetable Research Station, Kalyanpur, Kanpur, of the C. S. Azad University of Agriculture and Technology. The sowing was done on 6.12.83 in 1 x 3.5 m plots with interrow spacing of 30 cm and interplant spacing of 10 cm. Observations were recorded on five randomly selected plants per plot in each replication for days to flower ( $X_1$ ), pod length ( $X_2$ ), number of seeds/pod ( $X_3$ ), pod weight ( $X_4$ ), test weight of 100 green grains ( $X_5$ ), and yield per plot of green pods. Selection indices for yield were constructed

according to the procedure discussed of [3]. Along with different selection indices, their expected genetic advance from different selection indices at 5% selection intensity and relative efficiency indices of each function over straight selection were also calculated.

**Table 1. Selection index, discriminant function, expected genetic advance in yield and relative efficiency from the use of different selection indices**

Selection index	Discriminant function	Expected genetic advance	Relative efficiency
X1X2	$W = 0.0007 X_1 - 0.023 X_2$	0.590	31.0
X1X3	$W = 0.016 X_1 + 0.107 X_3$	0.302	15.9
X1X4	$W = 0.011 X_1 - 0.007 X_4$	0.160	8.4
X1X5	$W = 0.016 X_1 + 0.064 X_5$	0.752	39.6
X2X3	$W = -0.289 X_2 + 0.178 X_3$	0.722	38.0
X2X4	$W = -0.346 X_2 + 0.187 X_4$	0.703	37.0
X2X5	$W = 0.203 X_2 + 0.057 X_5$	0.880	46.3
X3X4	$W = 0.143 X_3 - 0.099 X_4$	0.291	15.3
X3X5	$W = 0.012 X_3 + 0.062 X_5$	0.718	37.8
X4X5	$W = 0.172 X_4 + 0.079 X_5$	0.819	43.1
X1X2X3	$W = 0.007 X_1 - 0.282 X_2 + 0.185 X_3$	0.728	38.3
X1X2X4	$W = 0.002 X_1 - 0.344 X_2 + 0.187 X_4$	0.704	37.0
X1X2X5	$W = 0.007 X_1 - 0.193 X_2 + 0.058 X_5$	0.885	46.6
X1X3X4	$W = 0.016 X_1 + 0.164 X_3 - 0.095 X_4$	0.360	18.9
X1X3X5	$W = 0.018 X_1 + 0.037 X_3 + 0.062 X_5$	0.757	39.8
X1X4X5	$W = 0.012 X_1 - 0.161 X_4 + 0.079 X_5$	0.836	44.0
X2X3X4	$W = -0.340 X_2 + 0.127 X_3 + 0.111 X_4$	0.748	39.4
X2X3X5	$W = -0.239 X_2 + 0.103 X_3 + 0.051 X_5$	0.909	47.8
X2X4X5	$W = -0.187 X_2 - 0.024 X_4 + 0.060 X_5$	0.881	46.4
X3X4X5	$W = 0.146 X_3 - 0.255 X_4 + 0.079 X_5$	0.970	45.8
X1X2X3X4	$W = 0.006 X_1 - 0.333 X_2 + 0.135 X_3 + 0.108 X_4$	0.752	39.6
X1X2X3X5	$W = 0.010 X_1 - 0.227 X_2 + 0.112 X_3 + 0.052 X_5$	0.919	48.4
X1X2X4X5	$W = 0.007 X_1 - 0.174 X_2 - 0.028 X_4 + 0.061 X_5$	0.886	46.6
X1X3X4X5	$W = 0.017 X_1 + 0.168 X_3 - 0.252 X_4 + 0.079 X_5$	0.899	47.3
X2X3X4X5	$W = -0.178 X_2 + 0.137 X_3 - 0.110 X_4 + 0.061 X_5$	0.923	48.6
X1X2X3X4X5	$W = 0.012 X_1 - 0.155 X_2 + 0.154 X_3 - 0.126 X_4 + 0.064 X_5$	0.936	49.3

## RESULTS AND DISCUSSION

The significant treatment variance for all the six characters indicated that the strains included in the present study were highly variable.

Different selection indices, their expected genetic advance and relative efficiency indices of each function over straight selection are presented in Table 1.

The present investigation shows that selection based on single characters was no more efficient than selection indices based on two, three, four and five character combinations.

It is also evident from Table 1 that when more than one character is employed in the construction of selection indices for yield, a progressive increase in the efficiency of selection was noticed. Robinson et al. [3] in corn also recorded a progressive increase in efficiency of selection with the inclusion of every additional character but in peas the selection based on three character combinations has been emphasized [4, 5]. When two characters at a time were considered for construction of selection indices, the relative efficiency ranged from 8.4 for two-character combination of days to flower and pod weight to 46.3 in the combination of pod length and test weight of 100 green grains. The combinations of three characters exhibiting greater relative efficiency than two-character combinations were pod length + seeds/pod + test weight (49.8), followed by the combinations days to flower + pod length + test weight (46.6), and pod length + pod weight + test weight (46.4). All the four character combinations except one, i.e. days to flower + pod length + seeds/pod + pod weight, gave higher efficiency than the three-character combinations (ranging from 46.6 to 48.6). Similarly, the five-character combination exhibited the maximum relative efficiency of 49.3. The main yield components in table peas are reported to be seeds/pod [4-6, 8, 9], green pod weight [5, 7] and 100-green seed weight, i.e. test weight [6, 7]. This suggests that increase in these traits will readily increase pod yield.

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