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Genetic Merit Based Genotype Selection for Physical Fruit Traits in Mango

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ABSTRACT

Selection indices based on their genetic merit of fruit characters were constructed for fruit yield/tree in 48 genotypes in mango (*Mangifera indica* L.). Genotypes, SBM 01-35, SBM 01-26, SBM 01-9, SBM 01-10 and SBM 01-26 in on year while, SBM 01-12, SBM 01-36, SBM 01-26, SBM 01-28 and SBM 01-9 in off year recorded highest fruit yield /tree. Genotypes SBM 01-10, SBM 01-36, SBM 01-09, SBM 01-30 and SBM 01-6 exhibited comparatively high estimates of selection indices during on year while, SBM 01-36 followed by SBM 01-12, Totapari, SBM 01-17 and SBM 01-29 showed the maximum estimates of selection index for fruit yield/ tree during off year. Alphonso followed by SBM 01-5, SBM 01-13, Langra and SBM 01-14 during on year and Dashehari, Safeda, SBM 01-3 and SBM 01-39 in off year exhibited the minimum estimates of varietal indices. Genotypes differed considerably in their ranking pattern based on selection indices. Genotypes SBM 01-9, SBM 01-10, SBM 01-30, SBM 01-6 and SBM 01-36 showed maximum varietal indices and phenotypic performance in both the years thus, appeared promising for use as parent in mango improvement programme.

Introduction

Mango (Mangifera indica L.) is the king of tropical fruits. The important mango growing countries are India, China, Thailand, Mexico, Pakistan, Philippines, Indonesia, Brazil, Nigeria and Egypt. India is the largest producer in the world. The productivity of this crop is 8.43m ton/ha which is low and needs improvement. The selection of genotypes based on phenotypic performance or yield per se is not much effective due to presence of genotype-environment interaction in phenotypic expression of the genes. On the other hand, selection based on genetic merits of several characters in the form of selection indices using discriminant function analysis of Fisher (1936) has been found a powerful tool in selection breeding programme. This analysis discriminate the economic genotypes from a population based on their genetic merits. However, the practical utility of this technique has not been tested in mango. In the present study, an attempt was therefore, made to select out the promising genotypes for fruit yield/tree based on genetic merits of fruit characters in mango (Mangifera indica L.).

Materials and Methods

The material of the present study comprised forty eight genotypes comprising 31 landraces from Bhopal division of Madhya Pradesh, 15 improved varieties and

02 hybrids, which were selected based on popular fruit characters like fruit size, suitability for pickles, sucking and table purposes, peel colour, firmness of pulp, flavor and storability. These genotypes were evaluated for physical characters of fruit in randomized complete block design with three replications in two subsequent on and off years. Two trees per genotype were randomly selected in each replication after fruit set. The fruits of selected trees were collected from each genotypes in each replications at full maturity stage and subjected to ripening for recording observations on fruits/ tree, fruit yield/tree (kg), weight per fruit (g), length and width of fruit (cm), peel thickness (cm), length and width of stone (cm), percentage weight of peel, pulp and stone and ratio indices of length-width, peel-pulp and pulp-stone. The procedure given by Smith (1936) and outlined by Singh and Choudhary (1985) was used for calculation of discriminate function coefficients for various characters. The mean values of each character of individual genotype were multiplied by respective discriminate coefficients and the sum was taken as selection index for genotype. Finally, the genotypes were arranged in order of their merit to select the best variety for further improvement through breeding techniques.

Table 1 Varietal indices for fruit yield/tree based on physical characters of fruits in mango

S.No.	Genotypes	uit yield/tree based on physical characters of for Selection indices for fruit yield/ tree		Per se	
		On year	Off year	On year	Off year
1	Alphonso	1015.84 I	11.59	60.57 V	27.57
2	Amrapali	1218.37	26.35	45.54 I	33.73
3	B Green	1078.66	17.35	80.54	19.57
4	Chousa	1257.37	24.16	112.07	27.87
5	Dahiyar	1324.06	8.07	114.97	19.27
6	Dashehari	1174.45	-0.68 II	72.08	14.80 IV
7	Fazli	1201.52	25.68	72.00	13.00 III
8	Gajaria	1106.66	10.43	70.26	15.23 V
9	Gulabkhas	1078.94	13.62	63.24	10.70 I
10	Langra	1054.94 IV	14.42	87.24	16.67
11	Mallika	1141.21	38.75	51.43 II	38.03
12	Neelum	1319.27	17.50	60.41 IV	35.00
13	Safeda	1396.58	0.52 III	67.30	17.83
14	Sinduria	1419.00	-5.01 I	59.89 III	11.17 II
15	Totapari	1251.47	49.50 III	111.44	57.80
16	Sehroli	1284.43	19.64	84.07	25.43
17	Suvaranrekha	1279.03	28.05	61.48	31.40
18	SBM 01-1	1172.09	40.91	184.95	42.17
19	SBM 01-2	1208.28	17.53	91.55	20.00
20	SBM 01-2	1514.04	7.58 V	94.01	21.93
21	SBM 01-4	1517.83	13.27	92.30	20.70
22	SBM 01-5	1026.94 II	11.71	102.48	23.10
23	SBM 01-6	2349.19 V	32.91	266.80	48.80
24	SBM 01-9	2421.64 III	38.82	332.43 III	55.20 IV
25	SBM 01-10	2668.86 I	34.91	307.64 IV	50.93
26	SBM 01-10	2146.73	20.90	226.08	22.67
27	SBM 01-11 SBM 01-12	1939.36	58.85 II	168.95	97.07 I
28	SBM 01-12	1035.06 III	26.83	90.26	41.80
29	SBM 01-14	1059.32 V	29.18	92.64	46.87
30	SBM 01-15	1902.71	24.66	118.00	24.03
31	SBM 01-17	1478.81	46.77 IV	110.51	52.73
32	SBM 01-19	1905.24	33.36	96.37	28.77
33	SBM 01-19	1603.17	35.38	185.76	46.87
34	SBM 01-22	1657.42	34.36	160.66	40.03
35	SBM 01-22	1614.64	37.19	215.42	37.17
36	SBM 01-24	1614.45	32.61	195.30	44.23
37	SBM 01-25	1178.83	22.15	158.18	36.63
38	SBM 01-26	1408.59	41.97	362.63 II	68.07 III
39	SBM 01-20	1451.48	21.11	123.39	29.50
40	SBM 01-28	1273.50	40.39	191.77	54.53 V
41	SBM 01-29	1525.14	45.05 V	201.32	48.60
42	SBM 01-30	2397.54 IV	44.02	263.96 V	51.07
43	SBM 01-30	1653.99	43.35	110.68	44.40
44	SBM 01-35	1253.67	62.04 I	388.92 I	96.43 II
45	SBM 01-35	2442.57 II	42.89	225.88	40.07
46	SBM 01-30	2011.83	15.37	142.88	22.47
47	SBM 01-37 SBM 01-38	2311.94	29.82	212.35	37.90
48	SBM 01-38	1802.80	6.98 IV	90.51	17.23

Results and Discussion

Significant differences among the genotypes were recorded for fruit yield/tree and all the physical characters of fruits including ratio indices. It indicates the existence of considerable variability for all the studied characters thus, offers good scope for the selection of desirable genotypes. The mean squares due to years was also significant for fruit yield/tree, peel thickness, percentage weight of peel and pulp, pulp to peel ratio and pulp to stone ratio. It revealed that these traits were influenced by the ambient climatic conditions prevailed in two years. The genotype and environment interaction was also significant for some of the fruit quality traits. Shrivastava

et al. (1987), Kashyap and Jyotishi (1969), Samad and Faruque (1976), Kapse et al. (1989). Yadav et al. (1995), Singh (2002) and Dwivedi and Mitra (2003) have also reported significant variability for physical fruit appearance in fruit crops confirming the present findings.

The varietal indices for fruit yield/ tree based on linear combination of physical fruit characters was in between 1015.84 to 2668.86 and -0.01 to 62.04 during on and off years, respectively (Table 1). SBM 01-10 followed by SBM 01-36, SBM 01-9, SBM 01-30 and SBM 01-6 exhibited maximum estimates of selection indices during the on year. Alphonso followed by SBM 01-5, SBM 01-

13, Langra and SBM 01-14 showed minimum estimates of selection indices in this year. During off year, SBM 01-36 recorded the maximum estimates of selection indices, followed by SBM 01-12, Totapari, SBM 01-17 and SBM 01-29 while, Sinduria followed by Dashehari, Safeda, SBM 01-39 and SBM 01-3 showed the minimum estimates for fruit yield/tree. The ranking pattern of the genotypes based on selection indices was different in two years, which may be due to genetic capability of alternate bearing in mango. Genotypes SBM 01-9, SBM 01-10, SBM 01-30, SBM 01-6 and SBM 01-36 showed maximum varietal indices and phenotypic performance in both the years thus, appeared promising for use in breeding programme aimed at genetic improvement in fruit yield/ tree in mango.

The present study thus conclude that landraces SBM 01-9, SBM 01-10, SBM 01-6 and SBM 01-13 appeared as higher yielder based on yield *per se* and genetic merits of fruit characters hence, these genotypes can be used in breeding programmes for genetic amelioration in fruit yield/ tree in mango. However, the study also suggests the testing of large number genotypes over years and environment for selection of genetically diverse genotypes in mango. It further suggests that promising landraces of Bhopal division of Madhya Pradesh be maintained in a natural park or mango garden to conserve the biodiversity in order to check the genetic erosion in mango.

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