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Divya Arti
Department of Vegetable
Science, Dr. YS Parmar
University of Horticulture and
Forestry, Nauni, Solan,
Himachal Pradesh, India

AK Sharma
Department of Vegetable
Science, Dr. YS Parmar
University of Horticulture and
Forestry, Nauni, Solan,
Himachal Pradesh, India

Genetic variability studies for yield and quality parameters in Brinjal (*Solanum melongena* L.)

Divya Arti and AK Sharma

Abstract

The estimates of variability viz. coefficients of variability (phenotypic and genotypic), heritability (in broad sense) and genetic advance as percent of mean (genetic gain) were worked out for selection of various characters. The phenotypic and genotypic coefficients of variability were high for incidence of fruit rot, incidence of fruit borer, number of marketable fruits per plant, fruit weight, yield per plant, yield per plot, yield per hectare, fruit length, number of branches per plant and fruit breadth. The high estimates of heritability (broad sense) were found for all the characters studied viz. incidence of fruit borer (99.39%), incidence of fruit rot (99.23%), ascorbic acid content (97.39%), plant height (96.83%), number of branches per plant (96.65%), number of marketable fruits per plant (96.44%), fruit breadth (94.80%), fruit length (94.42%), fruit weight (93.75%), total harvest duration (90.43%), total soluble solids (90.33%), days to 50 percent flowering (85.10%), days to first harvest (84.82%), yield per plant (83.87%), yield per plot (84.72%), yield per hectare (84.72%).

Keywords: Brinjal, variability, heritability, genetic advance, genetic gain

Introduction

Brinjal, eggplant or aubergine (*Solanum melongena* L.) is the most popular and widely cultivated warm season vegetable crop in the central, southern and south-east Asia and in some African countries. Based upon its highest production potential and availability of the produce to consumers, it is also termed as poor man's vegetable. The genetic coefficient of variation alone gives the real picture of variability. The extent of genetic variability existing in a crop is of great importance, since greater the genetic diversity, wider is the scope for selection. The phenotypic variability among a collection of germplasm gives an indication of potential genotypic variability; however, the quantitative characters are greatly influenced by the environment. Therefore, selection of the important traits for a sound breeding programme should be based on the extent of variability along with heritability and genetic advance.

Materials and Methods

Fifty diverse genotypes of brinjal were used for the investigations. The list of genotypes along with their sources is presented in Table 1. The investigation was carried out at the Experimental Farm of the Department of Vegetable Science, Dr YS Parmar University of Horticulture and Forestry, Nauni, Solan, (HP) during summer – rainy season of 2017. The Experimental Farm is situated at 30°51' N latitude and 77°11' E longitude at an elevation of 1270 m (amsl). Agro – climatically, the farm falls in the mid hill zone of Himachal Pradesh and is characterized by sub – temperate to sub – tropical climate with moderate rainfall (1000 – 1300 mm). Parameters of variability were estimated as per formula given by Burton and De vane (1953) ^[4]. Heritability in broad sense was calculated as per formula given by Burton and De vane (1953) ^[4] and Allard (1960) ^[2]. The expected genetic advance resulting from selection of five percent superior individuals was calculated as per Allard (1960) ^[2]. Genetic gain expressed as percent ratio of genetic advance and population mean was calculated by the method given by Johnson *et al.* (1955) ^[5].

Correspondence

Divya Arti
Department of Vegetable
Science, Dr. YS Parmar
University of Horticulture and
Forestry, Nauni, Solan,
Himachal Pradesh, India

Table 1: List of genotypes along with their sources

SN	Source	Genotype
1.	YSP UHF, Nauni, HP	UHF BRL – 1, UHF BRL – 2, UHF BRL – 3, UHF BRL – 4, UHF BRL – 5, UHF BRL – 6, UHF BRL – 7, UHF BRL – 8
2.	ASPEE College of Horticulture & Forestry, Navsari	IC-074224-1, H-295-3
3.	ICAR - IARI, New Delhi	DB-110, DBR-128, DB-144, DB-143, DBL-139, DB-109, DB-181, DBR-134, DB-30, Pusa Kranti
4.	ICAR - IARI, RRS, Katrain	PPC, PPL
5.	ICAR - IIHR, Bangluru	Arka Nidhi Arka Keshav, Arka Shirish, Arka Kusumkar, Arka Neelkanth
6.	GBPUA&T, Pantnagar	Pant Samrat, PB-4, PB-6, BARI, Swarn Anubhav
7.	PAU, Ludhiana	SR-301, SR-305, SR-312, PBH-3, SR-303, SR-333, BH-2, SR-321, BR 101, BR-322-2, BR-16, BR-123, PBHR-41, PBHR-42, Punjab Nagina, Punjab Sadabahar, Punjab Barsati

Results and discussion

Amongst the various parameters of variability, high phenotypic and genotypic coefficients of variability were found in incidence of fruit rot, incidence of fruit borer, number of marketable fruits per plant, yield per plant, yield per plot, yield per hectare, fruit weight, fruit length, number of branches per plant and fruit breadth indicating wide range of variations and offered better scope for improvement through selection. Moderate phenotypic as well as genotypic coefficients of variability were recorded in traits; ascorbic acid content, plant height, days to 50 percent flowering, total soluble solids and total harvest duration whereas, low phenotypic and genotypic coefficients of variability were found in days to first harvest. In the earlier studies also, the high estimates of phenotypic (PCV) and genotypic (GCV) coefficients of variation have been observed for fruit weight, number of fruits per plant, fruit length, fruit girth, yield per plant by Prasad and Singh (2003) [13], for number of fruits per plant and yield per plant by Negi *et al.* (2000) [10] and Ram *et al.* (2007) [14] and, for number of fruits per plant, fruit length and fruit yield per plant by Naik (2006) [9], while moderate PCV and GCV were observed for plant height by Patel *et al.* (2017) [11]; ascorbic acid content by Ravali *et al.* (2017) [15]; total soluble solids by Tirkey *et al.* (2018) [19]; plant height by Muniappan *et al.* (2010) [8], Ansari *et al.* (2011) [3], Lokesh *et al.* (2013) [7], Yadav *et al.* (2014) [21], Vidhya and Kumar (2015) [20] and Sujin *et al.* (2017) [18].

The estimates of heritability (broad sense) were found to be high for all the characters studied *viz.* incidence of fruit borer (99.39%), incidence of fruit rot (99.23%), ascorbic acid content (97.39%), plant height (96.83%), number of branches per plant (96.65%), number of marketable fruits per plant (96.44%), fruit breadth (94.80%), fruit length (94.42%), fruit weight (93.75%), total harvest duration (90.43%), total soluble solids (90.33%), days to 50 percent flowering (85.10%), days to first harvest (84.82%), yield per plant (83.87%), yield per plot (84.72%), yield per hectare (84.72%). High heritability for different traits indicated that large proportion of phenotypic variance was attributed to genotypic variance and therefore, reliable selection could be made for these traits on the basis of phenotypic expression. Similarly, Shekar *et al.* (2012) also reported high heritability in fruit length (99.00%), fruit diameter (97.00%), days to first flowering (97.00%), days to first picking (97.00%), number of fruits per plant (89.00%), fruit yield per plant (83.00%), fruit yield per plot (83.00%), fruit yield per hectare (83.00%) and number of branches per plant (81.00%). Akpan *et al.* (2016) [1] reported high broad sense heritability estimates for fruit circumference (98.37%), fruit diameter (97.44%), days to 50 percent flowering (96.69%), number of fruits per plant (83.27%) in brinjal grown in early season, while the late season planting experiment showed that fruit yield per hectare (98.08%), number of fruits per plant (99.48%), fruit circumference (99.16%) and fruit diameter (98.50%) were among traits with high broad sense heritability.

Table 2: Estimates of phenotypic and genotypic coefficients of variability, heritability, genetic advance and genetic gain of different characters in brinjal

Characters	Mean	Range	Coefficient of variation (%)		Heritability (%)	Genetic advance	Genetic gain
			PCV	GCV			
Plant height (cm)	91.11	52.13-130.47	21.15	20.81	96.83	38.43	42.18
Number of branches per plant	5.31	3.02-11.56	32.32	31.77	96.65	3.42	64.35
Days to 50 percent flowering	40.90	29.33 - 61.67	18.72	17.27	85.10	13.42	32.82
Days to first harvest	73.30	57.00-96.17	13.74	12.65	84.82	17.60	24.01
Total harvest duration	90.59	55.00-113.33	15.94	15.16	90.43	34.47	29.69
Fruit length (cm)	10.66	5.80-19.94	33.80	32.85	94.42	7.01	65.75
Fruit breadth (cm)	4.40	1.91-7.55	31.75	30.92	94.80	2.73	62.01
Fruit weight (g)	83.43	23.01-219.13	49.01	47.45	93.75	78.96	94.65
Number of marketable fruits per plant	12.60	1.39-29.86	62.46	61.34	96.44	15.64	159.03
Yield/plant (kg)	0.59	0.153-1.180	50.42	46.18	83.87	0.51	87.12
Yield /plot (kg)	9.12	2.39-18.25	50.61	46.59	84.72	8.06	88.33
Yield/hectare (q)	179.50	46.96-359.02	50.61	46.59	84.72	158.55	88.33
Total soluble solids (°Brix)	4.13	3.10-5.40	16.38	15.57	90.33	1.26	39.07
Ascorbic acid (mg/100 g)	12.35	5.29-18.89	27.51	27.14	97.39	6.82	70.72
Incidence of fruit rot (%)	4.26	1.17-19.81	85.69	85.36	99.23	7.47	224.48
Incidence of fruit borer (%)	9.02	1.14-23.22	64.19	63.99	99.39	11.85	168.42

An inquisition of data in Table 2 revealed that genetic gain (expressed as percent of population mean) was low to high in nature and ranged from 24.01 to 224.48 percent for different

characters under study. It was found high for the traits *viz.* incidence of fruit rot (224.48%), incidence of fruit borer (168.42%), number of marketable fruit per plant (159.03%),

fruit weight (94.65%), yield per hectare (88.33%), yield per plot (88.33%), yield per plant (87.12%), ascorbic acid content (70.72%), fruit length (65.75%), number of branches per plant (64.35%) and fruit breadth (62.01%). Moderate genetic gain was observed for plant height (42.18%), total soluble solids (39.07%) and days to 50 percent flowering (32.82%) while, low genetic gain was observed for days to first harvest (24.01%) and total harvest duration (29.69%).

The expression of the traits with high heritability and genetic advance is predominantly governed by additive gene effects and therefore, selection based on phenotypic performance will be useful to improve these characters in future as suggested by Prasad *et al.* (2004) ^[12]. High heritability coupled with moderate genetic advance as percent of mean was observed for harvesting period, fruit circumference, fruit length indicated that these characters were also controlled by additive gene and selection would be effective for improvement as reported by Karak *et al.* (2012) ^[6] and Solaimana *et al.* (2015) ^[17]. In our investigations, high heritability with low genetic gain was reported for the traits; days to first harvest and total harvest duration which implied that there was high influence of genotype x environment interaction and these traits were regulated by non additive gene action as suggested by Vidhya and Kumar (2015) ^[20] and accordingly, the improvement in these traits would be effective through recombinant breeding.

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