

Original Research Article

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Assessment Studies on Genetic Variability Traits in Different Lines of Tomato (*Solanum lycopersicum* L.)

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ABSTRACT

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Thirty eight genotypes of tomato were evaluated for yield and yield-contributing characters. Various genetic parameters like variability, heritability and genetic advance were calculated. The test genotypes exhibited a wide range of variability for traits under consideration. These traits were also found to be highly heritable. Phenotypic coefficient of variation (PCV) was found to be greater than genotypic coefficient of variation (GCV) for all the characters studied. High heritability was observed for fruit yield plant⁻¹ followed by average fruit weight while number of primary branches plant⁻¹ exhibited lowest heritability results. High heritability combined with high genetic advance was shown by average fruit weight suggesting, additive gene action plays a major role in governing these traits. Thus, these traits are amenable to simple selection for their improvement.

Introduction

Tomato (*Solanum lycopersicum* L.) is an important vegetable crop grown throughout the world. Botanically, tomato is a diploid with somatic chromosome number 24 (2n) belonging to family *Solanaceae*, the nightshade family of common vegetables, which also includes chilli, peppers, potato, etc. Tomato ranks third in priority after Potato and Onion in India but ranks second after potato in the world. India ranks second in the area as well as in production of tomato after China (FAOSTAT, 2013). It has become more popular all over the world because of a good

source of vitamins A and C, solids content, good taste and fruit set even at high temperature. It's direct use as raw vegetable to prepare in curries, stews, salads, fresh tomato juices as well as its soups and to make *chutney* have made it very popular (Bose *et al.*, 2002). Tomato fruit provides 3-4% sugar and total soluble solid in tomato ranges from 4 to 7%. It contains 15-30 mg/100 g ascorbic acid, 7.5-10 mg/100 mL titrable acids, 20-50 mg/100 g lycopene per fruit weight (Handbook of Horticulture, 2009). In addition, tomato has become a popular vegetable on account of its short duration, high yield potential, high profitability and economic viability (Bhandari

et al., 2017). Its fruits are consumed more as a fruit rather than as a vegetable. Tomato is a premier vegetable crop of round the year and but the production and productivity of this crop in India is far below as compared to global scenario. Considering the potentiality of this crop, there is a need for improvement and to develop varieties suited to specific agro-ecological conditions and also for specific end use. Efforts are being made to increase its productivity and quality by developing improved varieties. A thorough knowledge regarding the amount of genetic variability existing for various characters is essential for initiating the crop improvement programme. With limited variability much cannot be achieved and the breeder will have to enrich the germplasm or resort to create greater variability through hybridization, mutation and polyploidy breeding.

Germplasm is a prerequisite for any breeding programme; serves as a valuable source material as it provides scope for building of genetic variability. Progress in any breeding programme depends upon the magnitude of useful variability present in the population and the extent to which the desirable characters are heritable. Study of variability, heritability and genetic advance in the germplasm will help to ascertain the real potential value of the genotypes. Lack of variability has been thought to be one of the important reasons for the slow progress witnessed in making any significant gains by crop breeding. Though there is considerable variability in the crop, much of it may be in the form of conserved variability in view of their autogamous nature providing little scope for the shuffling of genes. The study and utilization of existing variability becomes highly essential.

Materials and Methods

The present investigation was carried out during *rabi* season of 2012-2013 at Vegetable

Research Farm, Institute of Agricultural Sciences, Banaras Hindu University, Varanasi (UP), India. The experimental material consist of 38 genotypes / cultivars (both exotic lines and indigenous lines) of tomato received from various sources, including Indian Institute of Vegetable Research, Varanasi and National Bureau of Plant Genetic Resources, New Delhi, at Department of Genetics and Plant Breeding, Institute of Agricultural Sciences, Banaras Hindu University given in Table 1. The experiment was laid out in randomized block design with three replications. Nursery was planted in second week of August and about 4 week old seedlings were transplanted during second week of September with row-to-row x plant-to-plant spacing maintained at 60 cm x 45 cm. Each plot consists of 10 plants and represents a single entry in each replication. Standard agronomic practices were followed to raise a good crop. Observations on days to first flowering and days to 50% fruiting were taken on plot basis. Five plants, excluding border plants, were randomly selected for recording of data on various yield traits such as primary branches, secondary branches, plant height (cm), clusters plant⁻¹, fruitscluster⁻¹, fruit plant⁻¹, locule number, fruit shape index, pericarp thickness (mm), average fruit weight (g), fruit yieldplant⁻¹ (kg). Analysis of variance, genotypic coefficient of variation (GCV), phenotypic coefficient of variation (PCV), heritability (broad sense), genetic advance (GA), genetic advance as per cent of mean, was done using Windostat[®] ver. 8.5 software for statistical data analysis.

Results and Discussion

The extent of variability with respect to thirteen quantitative characters in thirty-eight germplasm measured in term of phenotypic coefficient of variation (PCV), genotypic coefficient of variation (GCV), heritability, genetic advance and genetic advance as

percent of mean are given in Table 3. Analysis of variance revealed significant differences among germplasm for all the traits studies indicating presence of significant variability in the materials which can be exploited through selection Table 2. Similar results were noticed by Basavaraj *et al.*, (2010), Singh and Cheema (2005), Kaushik *et al.*, (2011), Dar and

Sharma (2011), Patel *et al.*, (2013). The research was carried out with the aim to develop high yielding varieties than the existing ones. The available genetic variability in the gene pool provides the opportunity for selecting superior types. The identification of the potential parents is pre-requisite for taking up an efficient breeding programme.

Table.1 List of the genotypes / cultivars used in the experiment

S. No.	Genotypes / Cultivars	Source
1.	PusaSadabahar	IARI, New Delhi
2.	DVRT-1-2	IIVR, Varanasi
3.	H-88-7-4	IIVR, Varanasi
4.	Floradale	IIVR, Varanasi
5.	DT-2	IIVR, Varanasi
6.	H-24	IIVR, Varanasi
7.	NDT- 3	NDUAT, Faizabad
8.	VR - 20	IIVR, Varanasi
9.	HT - 4	IIVR, Varanasi
10.	SwarnaLalima	IIVR, Varanasi
11.	TLC - 1	IIVR, Varanasi
12.	GT-20	IIVR, Varanasi
13.	FLA - 7171	IIVR, Varanasi
14.	NDTVR - 60	NDUAT, Faizabad
15.	Flawery	IIVR, Varanasi
16.	Feb - 04	IIVR, Varanasi
17.	BT - 120	IIVR, Varanasi
18.	NF - 315	IIVR, Varanasi
19.	PS - 1	IIVR, Varanasi
20.	Columbia	IIVR, Varanasi
21.	Cholnak - k	IIVR, Varanasi
22.	T Local	IIVR, Varanasi
23.	EC - 521069	NBPGR, New Delhi
24.	EC-521086	NBPGR, New Delhi
25.	EC-521087	NBPGR, New Delhi
26.	EC-528374	NBPGR, New Delhi
27.	EC-531803	NBPGR, New Delhi
28.	B-S-18-7	IIVR, Varanasi
29.	B-S-2-5	IIVR, Varanasi
30.	B-S-31-3	IIVR, Varanasi
31.	B-S-24-2	IIVR, Varanasi
32.	EC-520061	IIVR, Varanasi
33.	EC -538434	IIVR, Varanasi
34.	EC -538440	IIVR, Varanasi
35.	EC -538405	IIVR, Varanasi
36.	EC -539450	IIVR, Varanasi
37.	EC -538156	IIVR, Varanasi
38.	EC -538155	IIVR, Varanasi

Table.2 Analysis of variance for yield traits in 38 genotypes of tomato

S. No.	Characters	Mean Sum of Squares		
		Replication	Treatment	Error
1	Days to first flowering	14.798*	86.404**	3.069
2	Days to 50 per cent fruiting	6.342	71.598**	2.937
3	Primary branches per plant	4.312**	1.401**	0.292
4	Secondary branches per plant	2.168*	17.835**	0.657
5	Plant height	239.092**	633.236**	36.496
6	Number of clusters per plant	0.699	3.947**	0.358
7	Number of fruits per cluster	0.489	1.273**	0.162
8	Number of fruits per plant	10.87	707.255**	7.742
9	Pericarp thickness	0.038**	0.024**	0.004
10	Number of locules per fruit	0.24	2.382**	0.33
11	Average fruit weight	26.289	936.063**	19.037
12	Fruit yield per plant	0.017	4.792**	0.035
13	Fruit shape index	0.036*	0.104**	0.009

Table.3 Estimates of variability parameters for 13 yield attributes in tomato

S. No	Traits/Parameters	PCV	GCV	h ² (BS)	GA (5%)	GA as % of Mean 5%
1.	DFE	14.15	13.43	90.10	10.30	26.25
2.	D50Fr	8.50	8.00	88.60	9.28	15.51
3.	PB	21.82	16.31	55.90	0.94	25.12
4.	SB	37.61	35.62	89.70	4.67	69.50
5.	PH	20.01	18.40	84.50	26.71	34.83
6.	CI/ P	24.03	21.08	76.90	1.98	38.09
7.	Fr/ CI	20.16	16.81	69.60	1.05	28.89
8.	Fr/ P	32.61	32.08	96.80	30.95	65.01
9.	PT	21.41	16.61	60.20	0.13	26.54
10.	LN	28.00	23.00	67.50	1.40	38.92
11.	FSI	22.46	19.78	77.58	0.323	35.88
12.	AFW	45.44	44.09	94.10	34.94	88.12
13.	FY/ P	52.27	51.71	97.80	2.57	105.36

A wide range and significant variability was observed for most of the traits studied. Among all the thirteen traits studied, average fruit weight showed maximum range. Highest phenotypic and genotypic coefficient of variation (PCV and GCV) was observed for fruit yield plant⁻¹ (52.27 and 51.71) followed by average fruit weight (45.44 and 44.09) and fruitsplant⁻¹ (32.61 and 32.08). Rani and Anitha (2011), Manna and Paul (2012), Patel

et al., (2013) and Reddy *et al.*, (2013) also reported similar reports of higher PCV and GCV for fruits per plant, average fruit weight and fruit yield per plant. The lowest value for PCV and GCV was shown by days to 50% flowering (8.50 and 8.00). Environmental influence was very meagre on expression of these characters as it was evident by narrow gap between genotypic and phenotypic coefficient of variation. Selection among the

genotypes showing higher values of PCV and GCV will be beneficial for improvement of the trait on account to the variation present for the trait between the genotypes. In general, the estimates of broad sense heritability revealed highest value for fruit yield per plant (97.80) followed by number of fruits plant⁻¹ (96.80) and average fruit weight (94.10) where similar reports were made by Bai and Devi (1991), Kumari and Subramanian (1994) and Nwosu *et al.*, (2014). Estimates of genetic advance are more useful in selection than the heritability.

Average fruit weight followed by fruits plant⁻¹ and plant height exhibited high heritability coupled with high genetic advance indicating that these characters are governed by additive gene effect. Pujari *et al.*, (1995) also reported high estimates of heritability coupled with high genetic advance for number of fruits plant⁻¹ and average fruit weight suggesting that additive gene action would be responsible and selection for these traits could be used developing high yielding varieties. It may also be concluded that selection on the basis of these traits will be more useful for the improvement and towards attaining higher yield. Low genetic advance and heritability for pericarp thickness and primary branches indicates that these traits are chiefly controlled by non-additive gene action.

In the findings of Brar *et al.*, (2000), the number of fruit plant⁻¹, average fruit weight and total yield plant⁻¹ had low or moderate estimates of phenotypic and genotypic coefficients of variation, heritability and genetic advance which were in contrast to the present investigation.

A thorough knowledge regarding the amount of genetic variability, heritability and genetic advance existing for various characters is essential for initiating the crop improvement programme for development of new varieties.

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