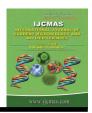


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Genetic Divergence Studies for Yield and Quality Traits in Onion (Allium cepa L.)

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ABSTRACT

Keywords

Onion, Genetic divergence, Cluster analysis, Yield, Quality

Article Info

Accepted: 21 May 2020 Available Online: 10 June 2020 Genetic divergence study, using Mahalanobis D² statistics, was conducted in twenty-three onion genotypes for fourteen characters. Among the characters studied; total yield, phenol, pyruvic acid, vitamin C content contributed maximum towards divergence. The maximum number of genotypes was grouped into cluster VI while the minimum was in cluster VIII. The intra-cluster distance varied from 0.00 (cluster VIII) to 14.618 (cluster IV). The maximum inter-cluster distance (32.763) was found between cluster V and VIII, indicating that hybridization can be taken up between the genotypes of these two clusters for obtaining superior recombinants or transgressive segregants. The genotype of cluster VIII showed superior performance for the plant height at 60 DAT, polar and equatorial diameter, number of scales per bulb, total sugar and dry matter percentage, TSS, vitamin C, phenol and pyruvic acid content and total yield. The genotypes of cluster III were found to be superior for the number of days to maturity. Thus, based on the diversity pattern and mean performances, the genotypes viz., Sukhsagar from cluster VIII; Arka Bheem and Arka Niketan from cluster VI; Bhima Raj from cluster II and HO-3 from cluster V would be the best choice as parents for future hybridization programs.

Introduction

Onion (*Allium cepa* L.) is one of the most important vegetables cum spice crops throughout the globe. It is commercially grown in temperate, sub-tropical as well as in tropical regions throughout the world (Brewster, 1990; Corgan and Kedar, 1990;

Currah and Proctor, 1990). Onion had its primary centre of origin in Southwest Asian gene centre, whereas, the Mediterranean gene centre is regarded as its secondary centre of origin (Vavilov, 1926). In India, majority of the harvest is obtained from winter (*rabi*) season. Onion is purely photosensitive and, on this basis, varieties are precisely classified

into short- and long-day types. Short day types generally have low yielding potential but have better shelf life whereas long day types have high yield potential with poor shelf life. The severity of purple blotch (caused by Alternaria porri) and Stemphylium blight (caused by Stemphylium vesicarium) disease shutters down the productivity during rabi season (Bal et al., 2019). Despite its global importance, there is a sharp lacuna in genetic improvement of onion when compared to other vegetable crops (McCallum et al., 2008). This may probably be due to lack of proper evaluation and data classification for various parameters including yield. Moreover, under Indian conditions, it is very challenging to breed short-day onion varieties for high yield potential coupled with resistance to different biotic and abiotic stresses. So, for success in any genetic improvement programme involving hybridization, selection of genetically diverse parents is of utmost importance (Arunachalam, 1981; Singh et al., 2013; Rivera et al., 2016). Moreover, genetic diversity studies provide an insight into the genetic distance among the accessions and thus increases the probability of obtaining transgressive segregants in F₂ and subsequent generations (Pal et al., 2017). Keeping these facts in mind, the present study was attempted to estimate the genetic divergence for different morphological, yield and quality traits in twenty-three genotypes through Mahalanobis D^2 analysis.

Materials and Methods

The present experiment was conducted at 'C' Block Farm of Bidhan Chandra Krishi Viswavidyalaya, Kalyani, Nadia, West Bengal. The experimental plot is located at 23.50° N latitude and 89° E longitude at a mean sea level of 9.75 m. The soil of the experimental farm was fine sandy-loam in texture, neutral in reaction with good water capacity. meteorological holding The observations taken during the experimental

period of 2017-18 and 2018-19 have been presented in Figures 1 and 2.

The experimental material was comprised of twenty-three onion genotypes collected from different states of the country (Table 1). The experiment was laid out in Randomized Complete Block Design with three replications. The seeds were sown in the nursery beds to raise the seedlings. Subsequently, 50 days old seedlings were transplanted in the main field during the rabi season of 2017-18 and 2018-19. A spacing of 15×10 cm was followed between rows and plants, respectively, in a plot size of $2 \text{ m} \times 1$ m, accommodating 133 plants per plot. The standard package of practices was followed thereafter to raise a healthy crop. The observations were recorded for plant height and number of leaves at 60 days after transplanting (DAT; cm), days to maturity, equatorial and polar diameter (mm), average weight of ten bulbs (kg), number of scales per bulb, total sugar and dry matter percentage, TSS (^oBrix), vitamin C (mg/g), phenol (mg/g), pyruvic acid (µmole/g) and total yield (kg/plot). The data were analyzed using SPAR 1.0 software packages.

Results and Discussion

To estimate the genetic distance among different genotypes for fourteen different traits, the genetic divergence with clustering pattern was worked out in the present study. The analysis of variance inferred highly significant differences among the genotypes for all the characters under study, indicating the existence of ample genetic divergence among them. Information on genetic diversity was also used to identify promising diverse genotypes, which may further be used in breeding programmes.

Analysis based on D^2 statistics revealed that the total yield per plot contributed maximum towards divergence *i.e.*, 57.312% (Table 2).

Meheta *et al.* (2008) also reported that the gross and marketable yield were major contributors towards genetic divergence in onion. The traits viz., phenol (27.272), pyruvic acid (6.719), vitamin C (6.324) content, number of days to maturity (0.791), TSS (0.400), dry matter percentage (0.400), number of scales per bulb (0.400), average weight of 10 bulbs (0.400) cumulatively contributed to 42.70% of the total divergence. The contribution of plant height and number of leaves at 60 DAT, equatorial diameter, total sugar content was found to be negligible.

Based on their genetic divergence, clustering pattern of twenty-three onion genotypes have been presented in Table 3. All the genotypes were grouped into eight clusters. The maximum number of genotypes had entered into cluster VI(n=5) followed by cluster I (n=4), cluster IV (n=4), cluster V (n=3), cluster II (n=2), cluster III (n=2) cluster VII (n=2) and cluster VIII (n=1). It was also evident from the grouping pattern that the collected from genotypes the same geographical area were clustered into different clusters, indicating that the genetic diversity is independent of the place of collection which might have resulted from the frequent exchange of germplasm between different geographical regions.

Table.1 List of the genotypes and along with their sources

Sl. No.	Name of the genotype	Sources				
1	Arka Bheem	ICAR-IIHR, Bangalore				
2	Arka Kirtiman	ICAR-IIHR, Bangalore				
3	Arka Lalima	ICAR-IIHR, Bangalore				
4	Arka Niketan	ICAR-IIHR, Bangalore				
5	Akola Safed	PDKV, Akola, Maharashtra				
6	Agrifound Light Red	NHRDF, Nasik, Maharashtra				
7	Bhima Dark Red	DOGR, Rajgurunagar				
8	Bhima Kiran	DOGR, Rajgurunagar				
9	Bhima Raj	DOGR, Rajgurunagar				
10	Bhima Red	DOGR, Rajgurunagar				
11	Bhima Safed	DOGR, Rajgurunagar				
12	Bhima Shakti	DOGR, Rajgurunagar				
13	Bhima Subhra	DOGR, Rajgurunagar				
14	Bhima Super	DOGR, Rajgurunagar				
15	Bhima Sweta	DOGR, Rajgurunagar				
16	Hisar-2	HAU, Hisar				
17	Hisar-4	HAU, Hisar				
18	Kalyanpur Red Round	CSAUAT, Kanpur				
19	L-28	NHRDF, Nasik, Maharashtra				
20	Onion HO-3	HAU, Hisar				
21	PRO-6	PAU,Ludhiana, Punjab				
22	Punjab Naroya	PAU,Ludhiana, Punjab				
23	Sukhsagar	Local market				

Table.2 Contribution of fourteen morphometric, yield-contributing and quality traits genetic divergence

Sl. No.	Characters	Contribution towards genetic divergence (%)				
1	Plant height at 60DAT (cm)	0.000				
2	Number of leaves at 60 DAT	0.000				
3	Number of days to maturity	0.791				
4	Polar diameter(mm)	0.000				
5	Equatorial diameter (mm)	0.000				
6	Average weight of 10 bulbs (kg)	0.400				
7	Number of scales per bulb	0.400				
8	Total sugar (%)	0.000				
9	Dry matter (%)	0.400				
10	TSS (°Brix)	0.400				
11	Vitamin C (mg/g)	6.324				
12	Phenol (mg/g)	27.272				
13	Pyruvic acid (µmole/g)	6.719				
14	Total yield (kg/plot)	57.312				
	Total	100				

Table.3 Clustering pattern of twenty-three onion genotypes based on genetic divergence

Cluster	Number of genotypes	Name of genotypes
I	4	Bhima Sweta, Hisar 4,L-28, Bhima Subhra
II	2	Bhima Raj, Hisar 2
III	2	PRO-6, Arka Lalima
IV	4	Punjab Naroya, Bhima Dark Red, Kalyanpur Red Round, Bhima Red
V	3	Bhima Super, Bhima Safed, Onion HO-3
VI	5	Akola Safed, Agrifound Light Red, Arka Bheem, Arka Kirtiman, Arka Niketan
VII	2	Bhima Shakti, Bhima Kiran
VIII	1	Sukhsagar

Table.4 Average intra- (Diagonal and underlined) and inter-cluster (Upper half Diagonal) distance

Clusters	I	II	III	IV	V	VI	VII	VIII
I	<u>9.791</u>	10.939	9.433	22.068	9.007	20.153	11.917	30.294
II		<u>5.095</u>	8.934	16.768	12.977	14.774	7.965	22.528
III			<u>5.462</u>	16.974	10.345	15.188	11.124	25.946
IV				<u>14.618</u>	24.460	12.774	19.993	18.441
V					<u>8.708</u>	22.464	13.304	32.763
VI						<u>12.407</u>	17.697	17.036
VII							<u>12.296</u>	24.437
VIII								<u>0.000</u>

Table.5 Cluster means for different characters among twenty-three genotypes of onion

Characters	I	II	III	IV	V	VI	VII	VIII
Plant height at 60DAT(cm)	53.06	48.55	51.33	51.70	51.87	50.78	51.35	59.96
Number of leaves at 60DAT	5.36	5.36	5.000	4.90	5.13	5.30	5.56	6.06
Days to maturity	131.16	130.33	133.50	132.58	130.11	133.06	131.50	132.00
Polar diameter (mm)	51.20	48.06	55.08	55.31	53.24	52.31	51.79	58.50
Equatorial diameter (mm)	50.24	52.61	46.73	50.37	47.00	49.78	53.74	56.95
Average weight of 10 bulbs (kg)	0.57	0.54	0.53	0.57	0.46	0.54	0.67	0.62
Number of scales/bulb	7.08	6.77	7.54	7.52	7.07	9.33	6.83	10.33
Total sugar%	1.96	1.22	1.59	1.33	1.48	1.76	2.17	3.48
Dry matter%	4.65	4.76	12.93	8.26	12.88	14.63	20.08	32.33
TSS(°Brix)	8.39	8.42	7.72	8.79	7.15	12.06	10.20	14.29
Vitamin C (mg/g)	0.24	0.48	0.25	0.34	0.24	0.37	0.48	0.71
Phenol (mg/g)	3.14	4.55	4.25	7.10	2.78	6.44	3.83	8.20
Pyruvic acid (µmole/g)	1.30	1.49	1.68	1.13	1.37	1.36	1.99	3.16
Total yield(kg/plot)	2.86	2.83	2.41	2.45	1.98	2.71	3.67	5.25

Figure.1 Meteorological observations during the experimental period of 2017-18

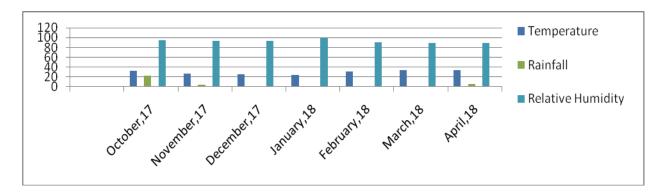
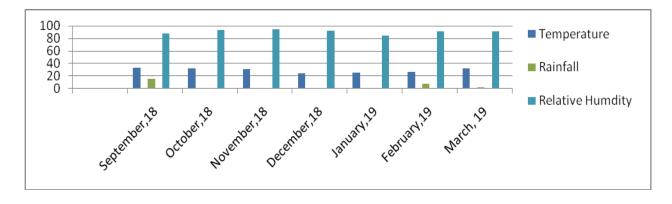


Figure.2 Meteorological observations during the experimental period of 2018-19



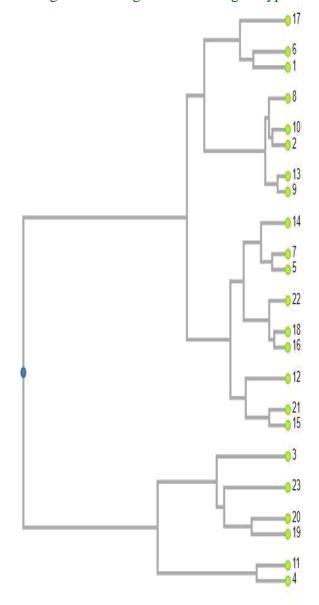


Figure.3 Representation of genetic divergence in onion genotypes through the dendrogram

Where,

1-Bhima Sweta, 2-Hisar-4, 3-Punjab Naroya, 4-Bhima Dark Red, 5-PRO-6, 6-Bhima Super, 7-Arka Lalima, 8-Bhima Safed, 9-L-28, 10-Onion HO-3, 11-Akola Safed, 12-Agrifound Light Red, 13-Bhima Subhra, 14-Arka Bheem, 15-Kalyanpur Red Round, 16-Bhima Raj, 17-Bhima Shakti, 18-Hisar-2, 19-Arka Kirtiman, 20-Arka Niketan, 21-Bhima Red, 22-Bhima Kiran, 23-Sukhsagar

The average inter-cluster and intra-cluster distance (D^2) values have been presented in Table 4. The diagonal figures in the table represent the intra-cluster distance (Fig. 3). The intra-cluster distance varied from cluster VIII (0.000) to cluster IV (14.618). The

minimum intra-cluster distance in cluster VIII was due to the presence of only a single genotype in that. The maximum inter-cluster distance (32.763) was found between cluster V and VIII. This indicates that hybridization can be taken up between the genotypes

belonging to these two clusters for obtaining superior recombinants or transgressive segregants. Further, the inter-cluster distance between cluster V and VIII was followed by cluster I and VIII (30.924), cluster III and VIII (25.946), cluster IV and V (24.460), cluster VII and VIII (24.437). The minimum inter-cluster distance (7.965) was recorded between cluster II and VII, indicating the genetic relatedness of the genotypes grouped into these two clusters. Therefore, breeding schemes involving hybridization should be avoided between the members of cluster II and VII.

The cluster means of different traits under study have been presented in Table 5. The genotypes of cluster VIII showed superior performances for the plant height at 60 DAT (59.96 cm), polar (58.50 cm) and equatorial diameter (56.95 cm), number of scales per bulb (10.33), total sugar percentage (3.48), dry matter percentage (32.33), TSS (14.29 °B), vitamin C (0.71 mg/g), phenol (8.20 mg/g), pyruvic acid (3.16 µmole/g) content and total yield per plot (5.25 kg). The genotypes of cluster III outperformed for the number of days to maturity (133.50).

Based on the diversity pattern and mean performances of the genotypes under study, it is concluded that the genotypes Sukhsagar from cluster VIII; Arka Bheem and Arka Niketan from cluster VI; Bhima Raj from cluster II and HO-3 from cluster V would be the best choice as parents for future hybridization programs for improving different morphometric, yield and quality traits in onion.

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