

Genetic analysis in ridge gourd [*Luffa acutangula* (Roxb.) L.] under hot arid conditions

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Abstract

The present investigation was conducted to determine variability, heritability, genetic advance and correlation of different yield and yield contributing characters of ridge gourd. The range of variation was higher in fruit length (3.66-26.93cm). The highest GCV (29.40) and PCV (32.29) was recorded in node at which first female flower appeared. The heritability estimate values in broad sense were higher for all the traits except number of branches/ plant, days to first female flower and days to first fruit harvest. The marketable fruit yield per plant had positive and significant correlation with fruit weight (0.786) and fruit length (0.448) at phenotypic level.

Key words: Ridge gourd, variability, GCV, PCV, heritability, genetic advance, correlation.

Introduction

Ridge gourd [*Luffa acutangula* (Roxb.) L.] is one of the important warm season vegetable crop which belong to cucurbitaceae family and grown in different parts of India. The immature fruits are consumed as vegetable and used in the preparation of chutneys and curries (Chandra, 1995). The fruits contain good amount of calcium, phosphorus, ascorbic acid, iron and fibre content (Aykroyd, 1963). Being a warm season vegetable crop, it has the ability to tolerate high temperature which ensures its adaptability for widespread cultivation throughout the tropics. Due to monoecious condition, it is a highly cross pollinated and had wide variability in quantitative characters (Richharia, 1948). Yield is polygenic which is highly influenced by environment and thus complicate the selection processes therefore, the knowledge of correlation is necessary for effective selection (Chowdhury *et al.*, 2002 and Choudhary *et al.*, 2008). The existence of wide genetic diversity in ridge gourd in arid region provides ample scope to screen the best genotypes for specific traits. Keeping in view, the present investigation was carried out with the objectives to determine variability, heritability, genetic advance and correlation among growth and yield traits in

ridge gourd.

Material and Methods

The experimental material consisted of 20 diverse genotypes of ridge gourd and evaluated in Randomized Block Design with three replications at Research Farm, Central Institute for Arid Horticulture, Bikaner, Rajasthan during summer season of 2011 and 2012. The spacing maintained between rows was 2.5m and between plants 0.60m. Fertilization, drip irrigation, other cultural practices and need based plant protection measures were followed as per recommendation. The data were recorded on 12 qualitative characters on five randomly selected plants from each replication. The recorded data were pooled and statistically analyzed using INDOSTAT statistical package (Indostat Services, Hyderabad).

Results and Discussion

The analysis of variance showed highly significant mean sum of square estimates for all quantitative traits under study which indicate a wide range of variability exists in genotypes (Table 1).

Table 1: Analysis of variance (mean sum of squares) for growth, flowering and yield traits of ridge gourd

S. No.	Characters	d.f.	Replication	Treatment	Error
1.	Number of branches/ plant	2	19	38	
2.	Vine length (m)	1.62	1.66**	0.64	
3.	Internodal length (cm)	0.63	1.09**	0.26	
4.	Days to 1 st female flower	13.91	17.87**	4.35	
5.	Node at which 1 st female flower appeared	32.08	29.35*	12.65	
6.	Ovary length (cm)	9.51	58.99**	3.79	
7.	Fruit length (cm)	0.42	4.43**	0.87	
8.	Fruit diameter (cm)	10.83	71.90**	4.41	
9.	Days to 1 st fruit harvest	0.63	3.58**	0.47	
10.	Marketable fruits/ plant	6.85	34.16**	11.77	
11.	Fruit weight (g)	0.69	95.65**	6.50	
12.	Marketable yield/ plant (kg)	148.38	1198.34**	46.74	
		0.10	0.42**	0.06	

*and ** indicate significant at 5 and 1 per cent probability level, respectively.

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The extent of variability among 20 genotypes was measured in terms of mean, range, genotypic coefficient of variation (GCV), phenotypic coefficient of variation (PCV), heritability in broad sense (h^2), genetic advance (GA) and genetic advance as per cent of mean (Table 2). The range of variation was higher in fruit length (3.66-26.93cm) followed by node at which first female flower appeared (2.93-20.27) and fruit weight (18.25-112.16g). The minimum range of variability was recorded in days to first female flower (42.27-53.60). The highest GCV was recorded in node at which first female flower appeared (29.40) followed by number of marketable fruits per plant (25.48) and PCV was highest for node at which first female flower appeared (32.29) followed by ovary length (29.04). High magnitude of GCV is an indication of high genetic variability among the genotypes which provides ample

scope for improvement of these characters through simple selection. These findings are in close agreement with the results of Varalakshmi *et al.* (1995), Singh *et al.* (2002) and Choudhary *et al.* (2008). The heritability estimates in broad sense were higher for all the traits except number of branches/ plant, days to first female flower and days to first fruit harvest indicating substantial improvement can be made using standard selection criteria. Similar findings have also been reported by Varalakshmi *et al.* (1995), Karuppaiah *et al.* (2002) and Singh *et al.* (2002). High heritability along with high GA and high GA as per cent of mean was observed for node at which first female flower appeared, fruit length, marketable fruits/ plant and fruit weight, indicating the presence of additive gene action thus these characters could be used in selection on the basis of phenotypic performance.

Table 2: Mean, range, variability, heritability and genetic advance for growth, flowering and yield traits of ridge gourd

Characters	Mean	Range	GCV	PCV	h^2	Genetic advance (GA)	GA as % of mean
No. of Branches/ plant	4.53	3.47-6.33	12.88	21.86	34.71	0.71	15.63
Vine length (m)	3.50	2.16-4.69	15.05	20.87	51.98	0.78	22.35
Internodal length (cm)	14.08	6.47-16.85	15.08	21.13	50.91	3.12	22.16
Days to 1 st female flower	48.60	42.27-53.60	4.85	8.78	30.56	2.69	5.53
Node at which 1 st female flower appeared	14.59	2.93-20.27	29.40	32.29	82.91	8.04	55.14
Ovary length (cm)	4.92	1.53-8.27	22.20	29.04	58.43	1.72	34.95
Fruit length (cm)	20.85	3.67-26.93	22.74	24.88	83.60	8.93	42.84
Fruit diameter (cm)	4.70	3.47-7.10	21.67	26.13	68.78	1.74	37.02
Days to 1 st fruit harvest	54.60	48.27-61.13	5.00	8.03	38.81	3.51	6.42
Marketable fruits/ plant	21.40	16.67-42.80	25.48	28.12	82.05	10.17	47.54
Fruit weight (g)	84.16	18.25-112.16	23.28	24.67	89.09	38.12	45.27
Marketable yield/ plant (kg)	1.72	0.77-2.21	19.94	24.76	64.86	0.57	33.09

In general the magnitude of genotypic correlation coefficients was higher than respective phenotypic correlation coefficients (Table 3) which establish inherent relationship among the characters as also shown by the findings of Chowdhury *et al.* (2002), Prasanna *et al.* (2002) and Hanumegowda *et al.* (2011). The marketable fruit yield per plant had positive and significant correlation with fruit weight (0.786) and fruit length (0.448) at phenotypic level. Days to first fruit harvest exhibited positive and highly significant correlation with days to first female

flower opening (0.850) and node at which first female flower appeared (0.469) at phenotypic level.

The traits like node at which first female flower appeared, fruit length, marketable fruits/ plant and fruit weight exhibited high heritability along with high GA and high GA as per cent of mean which could be improved through selection. Fruit weight and fruit length had positive and significant effect on yield per plant thus these traits could be utilized in selection of high yielding genotypes of ridge gourd.

Table 3: Genotypic (G) and phenotypic (P) correlation among growth, flowering and yield traits in ridge gourd

Trait	1.	2.	3.	4.	5.	6.	7.	8.	9.	10.	11.	12.
1. G	-	0.328**	0.402**	0.016	0.539**	-0.260*	0.105	0.011	0.237	-0.184	0.084	0.051
2. P	-	-	-0.100	0.074	0.150	-0.029	0.024	0.050	0.152	-0.048	0.069	0.092
3. G	-	-	-	-0.504**	-0.055	-0.309*	-0.228	0.403**	-0.268*	0.195	-0.348**	-0.299*
4. P	-	-	-0.036	-0.196	-0.041	-0.094	-0.076	0.214	-0.175	0.046	-0.219	-0.221
5. G	-	-	-	0.330**	0.450**	0.432**	0.525**	0.570**	0.408**	-0.645**	0.559**	0.617**
6. P	-	-	-	0.026	0.426**	0.090	0.419**	0.233	0.042	-0.360**	0.439**	0.487**
7. G	-	-	-	-	0.826**	0.214	0.434**	-0.348**	0.943**	-0.188	0.495**	0.586**
8. P	-	-	-	-	0.381**	0.216	0.254*	-0.071	0.850**	-0.040	0.187	0.248
9. G	-	-	-	-	-	0.541**	0.731**	0.101	0.795**	-0.620**	0.730**	0.606**
10. P	-	-	-	-	-	0.252*	0.630**	0.010	0.469**	-0.511**	0.606**	0.422**
11. G	-	-	-	-	-	-	0.828**	-0.108	0.494**	-0.814**	0.887**	0.611**
12. P	-	-	-	-	-	-	0.555**	0.063	0.250*	-0.567**	0.665**	0.412**
1. G	-	-	-	-	-	-	-	0.209	0.510**	-0.853**	0.824**	0.617**
2. P	-	-	-	-	-	-	-	0.156	0.288*	-0.704**	0.697**	0.443**
3. G	-	-	-	-	-	-	-	-	-0.289*	0.029	0.055	0.236
4. P	-	-	-	-	-	-	-	-	-0.173	0.067	0.013	0.194
5. G	-	-	-	-	-	-	-	-	-	-0.536**	0.613**	0.655**
6. P	-	-	-	-	-	-	-	-	-	-0.155	0.257*	0.226
7. G	-	-	-	-	-	-	-	-	-	-	-0.781**	-0.447**
8. P	-	-	-	-	-	-	-	-	-	-	-0.646**	-0.139
9. G	-	-	-	-	-	-	-	-	-	-	-	0.856**
10. P	-	-	-	-	-	-	-	-	-	-	-	0.786**
11. G	-	-	-	-	-	-	-	-	-	-	-	-
12. P	-	-	-	-	-	-	-	-	-	-	-	-

Critical r value=0.3 25 at 1 per cent and 0.250 at 5 per cent.

*and ** indicate significant at 5 and 1 per cent probability level, respectively.

1. Number of branches/ plant	5. Node at which 1 st female appeared	9. Days to 1 st fruit harvest
2. Vine length (m)	6. Ovary length (cm)	10. Marketable fruits/ plant
3. Internodal length (cm)	7. Fruit length (cm)	11. Fruit weight (g)
4. Days to 1 st female flower	8. Fruit diameter (cm)	12. Marketable yield/ plant (kg)

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