The study of genetic component in okra [(*Abelmoschus esculentus* L.) Moench]" crop

ABSTRACT

The present study of genetic component in okra okra [*Abelmoschus esculentus* (L.) Moench]" crop was conducted during kharif 2019 to ziad 2020 at the research farm of shri Durgaji post graduate college Chaneshwar Azamgarh U.P. with ten genotypes Azad Bhindi1, Azad Bhindi 2, Azad Bhindi 3, KS 312, VRO5, KS 442, KS 439, BO2, Arka Abhay and Prabhani Kranti along with 11 characters viz days to flowering, height of plant, number of branches per plant /main shoot,number of first fruiting node height of first fruiting node number of nodes per plant ,length of internodes, length of fruit, width of fruits , number of fruits per plant and fruit yield per plant . Genetic component of variance showed significant values for additive component $(\hat{D})$ for all the characters in both the generations except height of plant and number of branches in F_1generation.The dominance components $(\hat{H})$ were also highly significant for all the characters in both the generations except fruit yield per plant in F_2population.The degree of dominance $(\hat{H}_1/D)^{0.5}$ showed over dominance for all the characters in both the generations. The ratio of $(\hat{h}^2/\hat{H}_2)$ showed less than unity for all the characters in both generations.Analysis of variance for combining ability showed highly significant values both for gca and sca variances in based on both the generations. However, the magnitude of gca variance was more than sca variance for all the characters in both F_1 and F_2generations.Additive gene action was observed for all the characters in both the generations.

Introduction

Bhindi [*Abelmoschus esuculentus* (L.) Moench] is an important vegetable crop of India. It belongs to the family Malvaceae and having chromosome number 2n = 130. It behave as often cross-pollinated crop although it is potential self pollinated crop, 8.75 to 9.61 percent out crossing (Purewal and Randhawa, 1947). Okra is believed to have originated in the Hindustani centre of origin. However, Ethiopia is also considered as its native place from where it disseminated into Arabia down Nile Valley and was introduced in to Europe by the Moors and further into Louisiana during the early 17th century by French colonists (Woodruff, 1927). Additional utilization of vegetable in human diet may be ampel nutritive value of food as they are profusely nutritive. It is a better source of vitamin ‘A’, ‘B’, ‘C’, protein and minerals. (Singh *et al.,* 1974) analysed fruits and reported, 6.60 to 10.40 percent crude fibre, 84.60 to 90.50 percent edible portion, 14.40 to 18.60 percent protein and 8.20 to 9.16 percent ash of the total dry weight.

Okra is not only nutritive but has medicinal value also. The Ayurveda considered it as a virilitics. The root powder of okra mixed with sugar is used for treatment of *leucorrhea* which proved to be very effective remedy for this disease in women. People consume it for renal infections (i.e., stones in kidney). Eating fresh raw fruits as early morning nourishes the body and increases the semen content. Yawalkar (1965) reported it as an excellent source of iodine which helps in the control of goiter. Woodruff (1927) reported that mucilage from the stem and roots are used for cleaning of sugar cane juice in jaggery manufacturing in India. Fully ripened fruits and stem containing crude fibre are used in paper making. Its ripe seeds are roasted, ground and used as substitute for coffee in Turky (Mehta, 1959).

Material and Methods

The study of variability heritability and genetic advance in okra okra [*Abelmoschus esculentus*(L.) Moench]" crop was conducted during kharif 2019 to ziad 2020 at the research farm of shriDurgaji post graduate college Chaneshwar Azamgarh U.P. with ten genotypes Azad Bhindi1, Azad Bhindi 2, Azad Bhindi 3, KS 312, VRO5, KS 442, KS 439, BO2, Arka Abhay and Prabhani Kranti along with 11 characters viz days to flowering,height of plant, number of branches per plant /main shoot,number of first fruiting node height of first fruiting node number of nodes per plant ,length of internodes, length of fruit, width of fruits , number of fruits per plant and fruit yield per plant The following statistics were also estimated for drawing the conclusion.

Results:

$(\hat{H}_1 / \hat{D})^{1/2}$ = Mean degree of dominance

$(\hat{H}_2 / 4\hat{H}_1)$ = Proportion of positive and negative effects of genes in the Parents,

$\frac{4(\hat{D}\hat{H}_1)^{1/2} + \hat{F}}{4(\hat{D}\hat{H}_1)^{1/2} - \hat{F}}$ = It is used for F_1 generation and is also represented by KD/KR, which is the proportion of dominant and recessive genes in parents; F being significantly different from zero,

$\frac{[\frac{1}{4}(4\hat{D}\hat{H}_1)]^{\frac{1}{2}} + (\frac{1}{2})\hat{F}}{[\frac{1}{4}(4\hat{D}\hat{H}_1)]^{\frac{1}{2}} - (\frac{1}{2})\hat{F}}$ = It is used for F_2 generations

$(\hat{h}^2 / \hat{H}_2)$ = Gene group governing the character.

The estimated value of additive $(\hat{D})$ component of variation representing additive effect of genes was significant for all the characters in F_1& F_2 generations except height of plant and number of branches per plant in F_1 generation.In F_1 and F_2 generations the value of dominance variance $(\hat{H}_1 \& \hat{H}_2)$ were found highly significant for all the characters except fruit yield per plant in F_2 generation. The value of dominant component was also higher than additive component for all the characters in both generations.The value of $\hat{h}^2$ component was found significant and positive for days to followering, number of branches per plant, length of internode and number of fruits per plant in F_1 generation only. The value of $\hat{h}^2$ was positive and non significant for all remaining characters in both generations.The value of $\hat{F}$ component was found positive for characters number of branches per plant, number of nodes per plant, length of internode and length of fruit in F_1 and F_2 generations, height of plant and number of first fruiting node in F_1 and height of first fruiting node in F_2 generation only.Negative value of $\hat{F}$ component was exhibited for the characters days to flowering, width of fruit, number of fruits per plant and fruit yield per plant in both the generations, height of first fruiting node in F_1 and height of plant and number of first fruiting node in F_2 generation only.The estimated value of expected environmental component $(\hat{E})$ was observed positive and non significant responses for all the characters in both the generations.The value of average degree of dominance $(\hat{H}_1/\hat{D})^{0.5}$ was observed more than unity for all the characters in both the generations. The proportionf positive and negative alleles in parents denoted by ratio $(\hat{H}_2/4\hat{H}_1)$ was less than the theoretical value 0.25 for all the characters in both the generations.The ratio of dominant and recessive genes in parents i.e. $[(4\,\hat{D}\hat{H})^{0.5} + F / (4\,\hat{D}\hat{H})^{0.5} - F]$ was recorded less than unity for all the characters in both the generations except number of branches per plant, number of nodes per plant, length of internode and length of fruit in both the generations, height of plant and number of first fruiting node in F_1 and height of first fruiting node in F_2 generation only. The value of ratio $(\hat{h}^2/\hat{H}_2)$ was found less than unity for all the characters in both the generations.

Discussion and conclusion

Genetic component of variance showed significant values for additive component $(\hat{D})$ for all the characters in both the generations except height of plant and number of branches in F_1generation.The dominance components $(\hat{H})$ were also highly significant for all the characters in both the generations except fruit yield per plant in F_2population.The degree of dominance $(\hat{H}_1 / D)^{0.5}$ showed over dominance for all the characters in both the generations. The ratio of dominance and recessive genes in parents was recorded less than unity for all the characters in both the generations except number of branches per plant, number of nodes per plant, length of internode and length of fruit in both the generations, height of plant and number of first fruiting node in F_1 and height of first fruiting node in F_2 generation only.The ratio of $(\hat{h}^2 / \hat{H}_2)$

showed less than unity for all the characters in both generations. Analysis of variance for combining ability showed highly significant values both for gca and sca variances in based on both the generations. However, the magnitude of gca variance was more than sca variance for all the characters in both F_1 and F_2generations.Additive gene action was observed for all the characters in both the generations.

Table 1: Estimates of genetic components and related statistics for different characters under study in okra [*Abelmoschus esculentus* (L.) Moench]

Components	Generation	Characters										
		Days to flowering	Height of plant	Number of branches per plant/main shoot	Number of first fruiting node	Height of first fruiting node	Number of nodes per plant	Length of internode	Length of fruit	Width of fruit	Number of fruits per plant	Fruit yield per plant
$\hat{D}$	F1	13.27**	76.65	0.50	1.88**	8.57**	6.26**	4.11**	3.51**	0.11**	5.28**	414.50*
	SE±	1.98	45.13	0.06	0.29	0.83	1.99	1.19	1.18	0.03	1.53	188.11
	F2	13.26**	76.73*	0.51*	1.89**	8.53**	6.44**	4.11**	3.57**	0.12*	5.39**	7813.28**
	SE±	2.76	35.49	0.25	0.31	1.81	1.78	1.04	0.89	0.06	2.17	217.37
$\hat{H}_1$	F1	14.36**	275.35**	1.00**	2.78**	7.00**	14.99**	13.19**	8.33**	0.34**	10.46**	2347.17**
	SE±	4.22	96.06	0.13	0.62	1.77	4.23	2.54	2.50	0.06	3.26	400.41

	F2	110.20**	1474.63**	13.97**	16.11**	65.24**	107.74**	57.08**	36.63**	2.05**	74.73**	9601.53**
	SE±	23.53	302.21	2.15	2.63	14.40	15.12	8.86	7.57	0.47	18.48	1850.77
	F1	10.58**	200.85*	0.90**	2.32**	6.24**	9.61**	8.64**	5.44**	0.28**	6.94**	1834.37**
	SE±	3.59	71.64	0.11	0.53	1.50	3.60	2.16	2.13	0.05	2.77	340.31
$\hat{H}_2$	F2	103.86**	1280.70**	11.41**	14.36**	55.19**	103.45**	46.38**	26.59**	1.97**	65.73**	7813.28**
	SE±	20.00	256.84	1.83	2.24	13.09	12.85	7.53	6.44	0.40	15.71	1572.95
	F1	5.43*	-0.14	0.27**	0.07	1.83	-.0.06	3.03*	0.30	0.01	3.82**	138.24
$\hat{h}^2$	SE±	2.140	54.65	0.07	0.35	1.01	2.41	1.44	1.42	0.03	1.85	227.79
	F2	5.23	22.46	0.05	0.50	3.75	0.05	0.25	0.51	0.00	2.71	50.02
	SE±	3.35	42.98	0.31	0.37	2.19	2.15	1.26	1.08	0.07	2.63	263.22
	F1	-6.31	14.62	0.02	1.15	-0.71	5.56	4.61	4.00	-0.02	-3.83	-212.09
	SE±	4.56	104.13	0.14	0.67	1.92	4.49	2.75	2.71	0.06	3.53	434.03
$\hat{F}$	F2	-7.43	-149.49	0.70	-0.83	8.11	0.73	5.17	8.08	-0.07	-6.00	-297.74
	SE±	12.75	163.79	1.16	1.43	8.34	8.20	4.80	4.14	0.26	10.02	1003.08

Contd….

Components	Genera - tion	Days to flowering	Height of plant	Number of branches	Number of first fruiting	Height of first fruiting	Number of nodes	Length of internod	Length of fruit	Width of fruit	Number of fruits per	Fruit yield per plant

				per plant/main shoot	node	node	per plant	e			plant	
$\hat{E}$	F1	0.23	2.15	0.02	0.03	0.15	0.27	0.07	0.13	0.01	0.21	28.21
	SE±	0.60	13.61	0.02	0.09	0.25	0.60	0.36	0.35	0.01	0.16	56.72
	F2	0.31	2.06	0.02	0.02	0.19	0.09	0.07	0.07	0.01	0.09	18.09
	SE±	0.83	10.70	0.08	0.09	0.55	0.54	0.31	0.27	0.02	0.65	65.54
$(\hat{H}_1/\hat{D})^{1/2}$	F1	1.04	1.90	1.40	1.22	0.90	1.55	1.79	1.54	1.73	1.41	2.38
	F2	2.89	4.38	5.26	2.92	2.77	4.09	3.73	3.20	4.19	3.72	4.76
$(\frac{1}{4}\hat{H}_1/\hat{D})^{1/2}$	F1	0.52	0.95	0.70	0.61	0.45	0.77	0.89	0.77	0.86	0.70	1.19
	F2	1.44	2.19	2.63	1.46	1.38	2.04	1.86	1.60	2.09	2.86	2.38
$(\hat{H}_2/4\hat{H}_1)$	F1	0.18	0.18	0.23	0.21	0.22	0.16	0.16	0.16	0.21	0.17	0.20
	F2	0.24	0.22	0.20	0.22	0.21	0.24	0.20	0.18	0.24	0.22	0.20
$\frac{(4\hat{D}\hat{H}_1)^{1/2}+\hat{F}}{(4\hat{D}\hat{H}_1)^{1/2}-\hat{F}}$	F1	0.03	1.10	1.02	1.12	0.91	1.81	1.91	2.17	0.90	0.59	0.81
$\frac{\frac{1}{4}(4\hat{D}\hat{H}_1)^{1/2}+(\frac{1}{2})\hat{F}}{\frac{1}{4}(4\hat{D}\hat{H}_1)^{1/2}-(\frac{1}{2})\hat{F}}$	F2	0.82	0.64	1.30	0.86	1.42	1.03	1.41	2.11	0.87	0.74	0.97

$\hat{h}^2/\hat{H}_2$	F1	0.51	-0.0	0.30	0.03	0.29	-0.01	0.35	0.06	0.04	0.55	0.08
	F2	0.05	0.02	0.01	0.03	0.07	0.00	0.00	0.02	0.00	0.04	0.01
r	F1	-0.39	-0.14	-0.21	-0.58	-0.09	0.16	0.71	0.10	-0.30	0.23	0.39
	F2	0.26	0.24	-0.10	-0.44	0.33	-0.10	-0.21	0.36	0.55	-0.39	0.10

* significant at 5 per cent level

** significant at 1 per cent level

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