

Genetic variability for yield and quality traits in local germplasm of rice of Himachal Pradesh

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Rice (*Oryza sativa* L. 2n: 2x: 24) is the most important cereal crop cultivated widely in many parts of the world. Rice is grown under various agro-climatic conditions thus having greater variability in the local germplasm so, there is need to evaluate local germplasm adapted to H.P. for further use in breeding programme. Sufficient genetic variability, if present, can be exploited for developing superior cultivars. According to Fisher (1918), continuous variation exhibited by a quantitative characters, includes the heritable and non-heritable component. Wright (1921) reported that heredity component comprised of additive and non-additive portion and former responds to the selection.

Forty five diverse genotypes of rice from different geographical origin in H.P. were transplanted in 3 replications with in RBD design at Rice and Wheat Research Centre, Malan, HP during Kharif, 2018. In each replication, single seedling was transplanted per hill in 4 rows with 20 x 15 cm spacing.

The observations were recorded on five randomly taken plants from each plot for days to 50 per cent flowering, days to 75 per cent maturity, plant height at maturity, total tillers/plant at maturity, effective tillers/plant at maturity, grain yield/plant, spikelets/panicle, grains/panicle, panicle length, 1000-grain weight, spikelet fertility, kernel elongation, grain length (L), grain breadth (B), length breadth ratio (L:B), gel consistency (GC), gelatinization temperature (GT) rating, protein content and amylose content. Analysis of variance was carried out as suggested by Panse and Sukhatme (1978) GCV and PCV were carried out as per the methods suggested by Burton (1952) Heritability (BS) and Genetic Advance were estimated by using the formula suggested by Burton and Devane (1953).

Analysis of variance revealed that significant differences existed among the genotypes for all the characters studied. (Table 1&2). A perusal of genetic parameters revealed that phenotypic and genotypic coefficients of variation were high for GC. Moderate for plant height, total tillers/plant, effective tillers/plant, grains/panicle, spikelets/panicle and grain yield/plant and low PCV and GCV was recorded for rest of traits. Results were in accordance with Shrivastava *et al.*, (2015), Panwar (2005), Seyoum *et al.*, (2012), Devi (2015) Shashidhara (2017).

High heritability (Table 3) coupled with high genetic advance was observed for GC. High heritability coupled with moderate genetic advance was observed by days to 50% flowering, plant height, grains/panicle, spikelets/panicle, L:B ratio, and GT. Depicting these traits were controlled by additive gene effects thus selection for these traits will be effective. Grain yield showed moderate heritability coupled with low genetic advance. High heritability with low genetic advance was depicted by characters panicle length, 1000-grain weight, amylose content and protein content indicating that character is governed by non-additive genes and heterosis breeding may be useful here. Low heritability coupled with low genetic advance was shown by spikelet fertility. These findings were in confirmation with the results of Panwar (2005), Lingaiah *et al.*, (2018), Rukmani *et al.*, (2016), Singh *et al.*, (2006), Abebe *et al.*, (2017), Prabhu *et al.*, (2017), Dutta *et al.*, (2016), and Pang *et al.*, (2016).

Analysis of variation showed significant differences among all the genotypes for all the traits suggesting prevalence of wide range of genetic variability and scope of selection for these traits. Estimates of variability revealed that plant height, total tillers/plant, effective tillers/plant, panicle

Table 1: Analysis of variance for yield and morphological traits

Source of Variation	df	Days to 50% flowering	Days to 75% maturity	Plant height (cm)	Total tillers/plant at maturity	Effective tillers/plant at maturity	Panicle length (cm)	Grains /panicle (Nos.)	Spikelets /panicle (Nos.)	Grain yield/plant(g)	1000-grain weight (g)
Replication	2	1.43	10.14	18.53	1.62	2.02	3.77	96.83	93.96	31.39	92.99
Treatment	44	78.34*	11.35*	1208.24*	2.72*	2.68*	31.77*	2364.39*	2723.30*	23.38*	45.08*
Error	88	1.53	1.43	19.19	0.26	0.32	0.89	66.57	134.93	3.60	3.73

*Significant at 5% level

Table2: Analysis of variance for quality traits

Source of Variation	df	Grain length (mm)	Grain breadth (mm)	Length Breadth \ ratio (L:B)	Kernel elongation (mm)	Spikelet fertility (%)	Gel consistency (mm)	Gelatinization temperature rating	Amylose (%)	Protein (%)
Replication	2	0.113	0.020	0.024	0.24	9.65	4.58	0.59	0.02	0.12
Treatment	44	1.244*	0.137*	0.582*	1.74*	72.76*	734.85*	4.22*	16.42*	1.91*
Error	88	0.109	0.013	0.027	0.13	16.36	0.99	0.28	0.10	0.02

*Significant at 5% level

Table 3: Estimates of parameters of variability for different traits in rice genotypes

Traits	Mean	Range (min-max)	PCV (%)	GCV (%)	Heritability broad sense h ² bs (%)	GA as percentage of mean
Days to 50% flowering	90.70	78.00-100.33	5.74	5.58	94.37	11.16
Days to 75% maturity	123.37	120.33-129.00	1.76	1.47	69.84	2.54
Plant height (cm)	129.00	89.67-159.47	15.80	15.43	95.39	31.05
Total tillers/plant	5.10	3.73-7.60	20.34	17.75	76.13	31.90
Effective tillers/plant	4.99	3.40-7.53	21.10	17.78	71.01	30.87
Panicle length (cm)	25.80	20.77-31.77	12.96	12.43	92.03	24.57
Grains /panicle	134.08	79.33-195.10	21.52	20.64	92.00	40.79
Spikelets/panicle	149.91	82.27-211.60	21.07	19.59	86.48	37.54
Grain yield/plant (g)	19.00	13.30-26.43	16.81	13.52	64.67	22.39
1000- grain weight (g)	33.17	24.83-40.87	12.62	11.19	78.68	20.45
Grain length (mm)	6.68	5.43-7.73	10.44	9.21	77.70	16.72
Grain breadth (mm)	2.29	1.77-2.70	10.17	8.83	75.37	15.79
L:B ratio	2.95	2.03-4.07	15.62	14.58	87.09	28.02
Kernel elongation (mm)	8.96	7.30-10.60	9.11	8.20	80.95	15.19
Spikelet fertility (%)	89.20	78.33-96.63	6.65	4.86	53.46	7.32
Gel consistency (mm)	45.43	23.33-86.33	34.50	34.43	99.60	70.78
Gelatinization temperature (rating)	4.48	2.33-7.67	28.18	25.57	82.31	47.78
Amylose (%)	22.54	18.30-28.90	10.44	10.35	98.29	21.14
Protein (%)	8.39	6.53-10.27	9.63	9.46	96.54	19.15

length, grains/panicle, spikelets/panicle, L:B ratio ,GC and GT exhibited high to moderate heritability coupled with high to moderate genetic advance thus depicting these traits were controlled by additive gene effects thus selection for these traits will be effective. Grain yield showed moderate heritability coupled with low PCV, GCV and genetic advance. Moderate PCV and GCV was observed for days to grains/panicle, spikelets/panicle and L:B ratio, which indicated the presence of additive gene action and sufficient variability scope for improvement through selection.

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