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# Genetic parameters and character association studies in rice (*Oryza sativa* L.)

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#### Abstract

The genotypic coefficient of variation (GCV), phenotypic coefficient of variation (PCV), heritability, genetic advance, correlation and direct and indirect effect of different yield and yield attributing traits of 36 rice genotypes were studied. Among the traits number of productive tillers and number of total tillers had high genotypic and phenotypic coefficient of variation. High positive correlation was found for the traits *viz.*, panicle weight, number of grains per panicle and number of filled grains with single plant yield. But grain length, grain breadth and seed weight had high negative correlation with single plant yield. Path coefficient analysis showed that number of productive tillers, days to maturity and grain weight had high direct effect on grain yield per plant. Among the different traits studied, maximum indirect effect on grain yield was found with total tillers per plant followed by days to maturity. Here priority should be given for all these characters during selection for the improvement of grain yield.

Keywords: GCV, PCV, heritability, genetic advance, correlation and path coefficient analysis

#### Introduction

Rice is the most important food crop, since more than half of the world's population depends on rice for food calorie and protein. By the year 2025, the world will need about 760 million tonnes of paddy, in order to meet the growing demand (Duwayri *et al.*, 2000)<sup>[1]</sup>. To feed the ever increasing population, yield potential of rice should be improved through evolution of new breeding lines. Grain yield is the primary targeted trait for improvement of rice at both favourable and unfavourable condition (Tiwari *et al.*, 2011)<sup>[2]</sup>. Knowledge on genetic variability existing in the population and association between the yield and its related characters is very important for the development of high yielding varieties in future.

Estimation of genetic variability in polygenic traits is the key component of breeding programmes for broadening the gene pool of rice.

Association analysis is to determine the direction of selection and number of characters to be considered in improving grain the yield. The existence of correlation may be attributed to the presence of linkage or pleiotropic effect of genes or physiological and development relationship or environmental effect or in combination of all. Path coefficient analysis provides a direct and indirect effect and also observed a critical examination of specific forces to give correlation and measures the relative importance of each factor (Kishore *et al.*, 2015) <sup>[3]</sup>. Therefore, the present study was undertaken to determine the genetic parameters, correlation coefficient and path analysis among quantitative traits for isolation of superior genotypes in mutant lines of Anna (R) 4 derived through EMS treated, gamma ray and electron beam irradiation.

#### **Materials and Methods**

The present investigation was conducted at Agricultural College and Research Institute, Madurai. The experimental material consists of 35 genotypes which includes 12 advanced breeding lines ( $M_7$  generation) derived from Anna (R) 4 EMS induced mutant lines, 7 homozygous lines ( $F_5$  generation) of MDU 6 x *Jaldi Dhan* 6 and one homozygous line ( $F_5$ generation) of TKM 6 x *Jaldi Dhan* 6 and 15 mutant lines ( $M_4$  generation) of Anna (R) 4 irradiated entries derived from gamma irradiation (8 entries) and Electron beam irradiation (7 entries). These materials were evaluated with the check variety Anna (R) 4. These breeding materials are early maturing genotypes with drought tolerance. The crop was raised in *Kharif*, 2019 with a spacing of 25 cm between row to row and 15 cm between plant to plant in Randomized Block Design (RBD) with two replications. Each genotype consists of three rows with 3m length in both the replication. Totally seventeen quantitative characters were recorded at appropriate crop stages based on Standard Evaluation System of rice (IRRI, 2013). From each genotype five plants were randomly selected and the yield and yield attributing traits were recorded. The characters measured were plant height on 50<sup>th</sup> day (cm), days to fifty per cent flowering (days), SPAD meter value, plant height (cm) at mature stage, panicle length (cm), flag leaf length (cm), flag leaf breadth (cm), total tillers, productive tillers, days to maturity (days), panicle weight (g), number of grains per panicle, number of filled grains, grain length (cm), grain breadth (cm), hundred seed weight (g) and single plant yield (g).

Analysis of genotypic coefficient of variation (GCV), phenotypic coefficient of variation (GCV), broad sense heritability ( $h^2$ ), and genetic advance as per cent of mean, genotypic and phenotypic correlation, direct and indirect effect of different characters on yield was done by using the software GENRES, version 7.01.

#### **Result and discussion**

#### Variability, heritability and genetic advance

Genetic variations measured as PCV, GCV, broad sense heritability  $(h^2)$  and genetic advance as per cent of mean designates a wide range of variability among 36 genotypes for 17 yield component traits (Table 1).

The phenotypic coefficient of variation was higher than genotypic coefficient of variation for panicle weight and seed yield. It reveals that environmental influence on expression of that particular character (Mishra *et al.*, 2019)<sup>[4]</sup>. The environmental effect on any particular character depends on range of difference between genotypic coefficient of variation and phenotypic coefficient of variation. Other traits showed minimum difference between genotypic coefficient of variation indicates involvement of environment effect on expression of those traits is small. These similar findings were also reported by Hossain *et al.* (2018)<sup>[5]</sup>.

In this present study, number of total tillers and number of productive tillers were measured as high GCV and PCV values (Table 1 & Fig.1). It point towards the variability observed in 36 genotypes was high and so selection would be more effective for tiller numbers. The moderate genotypic coefficient of variation and phenotypic coefficient of variation values were observed for the traits number of filled grains followed by number of grains per panicle, single plant yield and grain weight. Hossain *et al.* (2018) <sup>[5]</sup> and Aditya and Bhartiya, (2013) <sup>[6]</sup> also reported moderate to high GCV and PCV values of number of grains per panicle and yield per plant. The lowest GCV and PCV values were found with days to maturity, followed by plant height, days to fifty per cent flowering and grain length. The same results also supported by Hossain *et al.* (2018) <sup>[5]</sup>.

Estimation of heritability helps to select the elite genotypes, in which those characters having very high and high heritability are greater than 80% and 60-80% respectively is specifies that character is less influenced by environmental effect. Selection of genotypes based on heritability without genetic advance leads inappropriate selection because heritability includes both the effect of additive and non- additive gene action (Aditya and Bhartiya, (2013)<sup>[6]</sup>. In order to get better selection efficiency, select those traits exhibits high heritability along with high genetic advance will give effective improvement. In this experiment high heritability was observed in seed weight (96.45) followed by plant height on 50<sup>th</sup> day, number of grains per panicle, flag leaf breadth

and days to fifty per cent flowering (Table 1). Akinwale *et al.* (2011) <sup>[7]</sup> and Hossain *et al.* (2018) <sup>[5]</sup> also reported days to fifty per cent flowering and number of grains per panicle respectively having high heritability. High heritability combined with high genetic advance was found in total tillers followed by productive tillers, filled grains and number of grains per panicle. So it's simply conveys that these traits are controlled by additive gene action and selection remains effective. Hence during selection more importance has to be given to these traits for selecting better rice genotypes.

#### **Correlation analysis**

#### Association between yield and component traits

Number of filled grains had highly positive association with single plant yield both genotypic and phenotypic level which also implied the importance of characters for grain yield improvement. The higher genotypic correlation coefficient ( $r_g$ ) than the phenotypic correlation coefficient ( $r_p$ ) reveals the association between these two traits is due to genetic control and the phenotypic value is narrowed by environmental influence. These findings were earlier reported by Kalyan *et al.* (2017) <sup>[8]</sup>.

Number of grains per panicle had positive correlation with single plant yield at genotypic and phenotypic level. Lalitha *et al.* (2019) <sup>[9]</sup>, also observed same results of positive correlation between single plant yield and number of grains per panicle. Number of filled grains had highly positive association with number of grains per panicle, because number of filled grains is determined by product of seed setting rate and number of grains per panicle. Hence the number of grains per panicle number, compactness and branching types. So it is directly contributed to higher grain yield compare with other yield contributing traits (Li *et al.*, 2019) <sup>[10]</sup>.

The panicle weight exhibit positive genotypic correlation with single plant yield, but there is no significant correlation were observed between single plant yield and panicle weight at phenotypic level. Positive genotypic correlation between these traits was earlier reported by Awasthi and Lal, (2014) <sup>[11]</sup>. Grain length, grain breath and seed weight unveiled high negative association with single plant yield at genotypic and phenotypic level and also exhibit negative and significant correlation at phenotypic level for these three traits. The same negative association for single plant yield and grain weight was computed by Saha *et al.* (2019) <sup>[12]</sup>. Number of grains per panicle increased with decrease in size of the grains but yield per plant decreased with increasing small grains. The smaller grain size of majority of selected mutants may be the reason behind this (Saha *et al.*, 2019) <sup>[12]</sup>.

#### Genotypic correlation between yield component traits

Plant height on 50<sup>th</sup> day had highly positive association with plant height at maturity, panicle length, flag leaf length, grain length and also positive association with grain breadth and seed weight. Days to maturity had significant and positive correlation with days to 50% flowering and flag leaf breadth. Saha *et al.* (2019) <sup>[12]</sup> also reported for days to 50% flowering and Aditya and Bhartiya, (2013) <sup>[6]</sup> for flag leaf breadth. Significant positive correlation of SPAD value with number of grains per panicle and number of filled grains per panicle observed in present study is in accordance with the report of Nithya *et al.* (2020) <sup>[13]</sup>. Because SPAD meter value of particular genotype directionally proportional to total chlorophyll content present in the leaf. Evans and Dunstone,

(1970) <sup>[14]</sup> reported small leaves contained high concentration of chlorophyll content in diploid and tetraploid wheat.

Plant height exhibit highly positive correlation with flag leaf length and positive relationship with panicle length. The positive correlation of plant height with panicle length was earlier supported by Parimala et al., (2020) <sup>[15]</sup> and for plant height with flag leaf length was given by Devi et al. (2017) <sup>[16]</sup> and Kishore *et al.* (2018) <sup>[17]</sup>. Panicle length had a highly positive correlation with flag leaf length, panicle weight, grain length and seed weight. Similar findings were reported by Devi et al. (2019) <sup>[18]</sup> for flag leaf length and Bhutta et al. (2019) <sup>[19]</sup>, Devi et al. (2019) <sup>[18]</sup> for panicle weight, Idris et al. (2012) <sup>[20]</sup> for grain length and seed weight were earlier found Devi et al. (2019) [18] and Bhutta et al. (2019) [19]. Ashrafuzzaman et al. (2009)<sup>[21]</sup> reported yield and yield traits had positive correlation with flag leaf area, because flag leaf is the trait which gives the photosynthetic to the grains. It was previously studied by Asana (1968)<sup>[22]</sup>. Rafiq et al. (2014)<sup>[23]</sup> found that increased flag leaf length would increases panicle length with increased number of branches which directly influences the grain yield.

The trait flag leaf breadth showed high positive association between days to maturity, panicle weight, grain breadth and seed weight. Positive correlation for panicle weight with flag leaf breadth was also reported by Devi et al. (2019) [18]. Productive tillers and total tillers which had highest positive and significant correlation and this were previously mentioned by Kalyan et al. (2017)<sup>[8]</sup> and Saha et al. (2019) <sup>[12]</sup>. Days to maturity had positive correlation with panicle weight; and panicle weight exhibit highly positive relationship with grain length and seed weight. In the current study, number of grains per panicle had significant positive correlation with number of filled grains per panicle. Similar results were previously observed by Saha *et al.* (2019)<sup>[12]</sup> and Parimala et al. (2020) <sup>[15]</sup>. This indicate that spikelet fertility led to higher seed set which in turn resulted in more number of filled grains per panicle.

In this present study seed weight exhibited high positive correlation with both grain length and grain breadth. Sakamoto and Matsuoka, (2008) <sup>[24]</sup> reported higher night temperature at the time of grain filling stage leads to reduction in dry matter production, grain length and breadth, so it may reduce the grain yield of the plant.

#### Path coefficient analysis

The direct and indirect effects of each independent variable in the direction of single plant yield is not clearly given by correlation, and so path coefficient analysis split up the direct and indirect effect of each character on single plant yield. So selection of superior genotypes with true relationship and characters association among diverse genotypes is easiest.

#### **Direct effect**

Among the seventeen characters, number of productive tillers exhibited positive and very high (5.062) directs effect on single plant yield, followed by days to maturity (3.721), seed weight (2.004), flag leaf length (1.241), plant height on 50<sup>th</sup> day (1.180) and SPAD value (1.037) (Table 3). The trait filled grains (0.968) had positive and high direct effect towards yield. Here the association between seed weight and single plant yield is under negative relationship but direct effect of seed weight on grain yield is positive. Hence, to reduce the undesirable indirect effect, direct selection for this trait has to be followed.

The trait total tillers (4.504) had negative but very high indirect effect on single plant yield followed by flag leaf breadth (1.471), plant height (1.462) and days to fifty per cent flowering (1.393).

#### **Indirect effect**

Days to maturity had positive and very high indirect effect on single plant yield through days to maturity (3.658) followed by total tillers (1.944). Plant height showed highly positive and indirect effect through plant height on 50<sup>th</sup> day (0.684). The same rationalization was earlier reported by Srijan et al. (2016). The indirect effect of filled grains (0.545) with SPAD value was observed as high positive effect. Positive and very high indirect effect of total tillers (1.736) followed by days to maturity (1.221) on yield by plant height was observed in this study. Flag leaf breadth exhibited positive and very high indirect effect on single plant yield through days to maturity (2.248) followed by total tillers (1.470). Plant height on 50<sup>th</sup> day (0.698), flag leaf length (0.996), total tillers (0.994) and grain weight (0.993) contributed positively and high indirect effect on grain yield per plant by panicle length whereas it exhibited negative and high indirect effect on plant yield through plant height (0.588), number of productive tillers (0.884), days to maturity (0.566) and grain length (0.643).

By the way of productive tillers (5.045), total tillers gives more positive and very high indirect effect on single plant yield and days to fifty per cent flowering (0.601), plant height (0.563) and flag leaf breadth (0.480) showed its positive and indirect effect on grain yield. Days to maturity showed positive and indirect effect through grain length (0.314), however flag leaf breadth (0.888), plant height (0.480) and seed weight (0.447) had negative with high indirect effect on grain yield. High and positive indirect effect of grain length and breadth on grain yield was observed by the influence of plant height on 50<sup>th</sup> day (0.662, 0.429), productive tillers (0.799, 0.879), and very high positive indirect effect through seed weight (1.941, 1.788).

Characters	Genotypic coeff. of variation (%)	Phenotypic coeff. of variation (%)	Broad sense heritability (h <sup>2</sup> )	Genetic advance as per cent of mean
Plant height on 50 <sup>th</sup> day	7.24	7.53	92.39	14.34
Days to fifty per cent flowering	4.37	4.61	89.74	8.53
SPAD value	7.01	7.91	78.62	12.81
Plant height	3.52	4.13	72.56	6.18
Panicle length	7.55	8.37	81.24	14.02
Flag leaf length	10.62	11.78	81.34	19.74
Flag leaf breadth	13.29	14.02	89.79	25.94
Total tillers	26.39	27.92	89.36	51.39
Productive tillers	26.61	28.63	86.41	50.97
Days to maturity	3.62	4.05	79.97	6.68

**Table 1:** Genetic parameters of thirty six rice genotypes for 17 quantitative traits

Panicle weight	9.05	16.27	30.92	10.36
No. of grains per panicle	17.93	18.74	91.52	35.34
Filled grains	19.11	20.62	85.83	36.47
Grain length	4.88	5.33	83.64	9.19
Grain breadth	7.97	8.69	84.01	15.04
100 grain weight	16.17	16.46	96.45	32.72
Single plant yield	16.69	20.08	69.08	28.57

Table 2: Genotypic correlation coefficient among 36 rice genotypes for 17 quantitative traits

Chanastana	PH(50)	DFF	SPAD	PH	PL	FL	FB	TT	РТ	DM	PW	NGPP	FG	GL	GB	100GW	SPY
Characters	(cm)	(days)	value	(cm)	(cm)	(cm)	(cm)	(nos.)	(nos.)	(days)	(g)	(nos.)	(nos.)	(mm)	(mm)	(g)	(g)
PH(50)	1	-0.126	-0.401*	0.580**	0.592**	0.680**	0.184	-0.073	-0.058	-0.109	0.027	-0.328	-0.404*	0.561**	0.364	0.416*	-0.220
DFF		1	-0.154	0.326	-0.224	-0.079	0.595**	-0.432**	-0.420*	0.983**	0.193	-0.054	-0.196	-0.289	-0.172	-0.208	0.126
SPAD value			1	0.003	-0.182	-0.456**	-0.188	0.026	0.055	-0.204	0.024	0.504	0.563**	-0.064	-0.080	-0.177	0.279
PH				1	0.402*	0.454**	0.011	-0.385*	-0.372*	0.328	0.089	0.216	0.072	0.123	-0.201	-0.100	0.130
PL					1	0.803**	0.258	-0.221	-0.175	-0.152	0.478	-0.008	-0.204	0.572**	0.270	0.495**	0.124
FL						1	0.126	-0.214	-0.208	-0.107	-0.025	-0.133	-0.246	0.229	0.037	0.234	0.116
FB							1	-0.326	-0.253	0.604**	0.733**	-0.139	-0.450**	0.329*	0.495**	0.430**	-0.063
TT								1	0.997**	-0.497**	-0.423*	-0.391	-0.300	0.113	0.106	0.114	-0.144
PT									1	-0.477**	-0.333*	-0.414*	-0.354*	0.158	0.174	0.172	-0.170
DM										1	0.393*	0.093	-0.073	-0.279	-0.218	-0.223	0.198
PW											1	0.370*	0.087	0.424**	0.314	0.460**	0.381*
NGPP												1	0.915**	-0.448**	-0.484**	-0.545**	0.500**
FG													1	-0.602**	-0.672**	-0.707**	0.569**
GL														1	0.748**	0.969**	-0.472**
GB															1	0.893**	-0.514**
100GW																1	-0.479**
SPY																	1

**Note:** \* indicates significance at 5%,\*\* indicates significance at 1%, PH (50)= Plant height on 50<sup>th</sup> day, DFF= Days to fifty per cent flowering, SPAD value= SPAD meter value, PH= Plant height, PL= Panicle length, FL= Flag leaf length, FB= Flag leaf breadth, TT= Total tillers, PT= Productive tillers, DM= Days to maturity, PW= Panicle weight, NGPP= Number of grains per panicle, FG= Number of filled grains, GL= Grain length, GB= Grain breadth, HGW=Hundred grain weight, SPY= Single plant yield

Table 3: Direct and indirect effects of the trait on yield in 36 rice genotypes for 17 quantitative traits

Characters	PH(50)	DFF	SPAD	PH	PL	FL	FB	ТТ	РТ	DM	PW	NGPP	FG	GL	GB	100GW	SPY
	(cm)	(days)	value	(cm)	(cm)	(cm)	(cm)	(nos.)	(nos.)	(days)	(g)	(nos.)	(nos.)	(mm)	(mm)	(g)	(g)
PH(50)	1.180	0.175	-0.416	-0.848	-0.131	0.844	-0.271	0.330	-0.292	-0.407	-0.005	-0.037	-0.391	-0.630	-0.156	0.833	-0.220
DFF	-0.148	-1.393	-0.160	-0.476	0.049	-0.098	-0.876	1.944	-2.125	3.658	-0.035	-0.006	-0.190	0.324	0.074	-0.417	0.126
SPAD value	-0.473	0.215	1.037	-0.004	0.040	-0.566	0.277	-0.115	0.276	-0.758	-0.004	0.057	0.545	0.072	0.034	-0.355	0.279
PH	0.684	-0.453	0.003	-1.462	-0.089	0.564	-0.016	1.736	-1.883	1.221	-0.016	0.024	0.069	-0.138	0.086	-0.201	0.130
PL	0.698	0.312	-0.189	-0.588	-0.221	0.996	-0.380	0.994	-0.884	-0.566	-0.086	-0.001	-0.197	-0.643	-0.116	0.993	0.124
FL	0.803	0.110	-0.473	-0.664	-0.177	1.241	-0.185	0.964	-1.053	-0.396	0.005	-0.015	-0.238	-0.258	-0.016	0.468	0.116
FB	0.217	-0.829	-0.195	-0.016	-0.057	0.156	-1.471	1.470	-1.282	2.248	-0.132	-0.016	-0.435	-0.370	-0.211	0.862	-0.063
TT	-0.087	0.601	0.026	0.563	0.049	-0.266	0.480	-4.504	5.045	-1.851	0.076	-0.044	-0.291	-0.127	-0.045	0.229	-0.144
PT	-0.068	0.585	0.057	0.544	0.039	-0.258	0.373	-4.488	5.062	-1.777	0.060	-0.047	-0.343	-0.177	-0.074	0.344	-0.170
DM	-0.129	-1.369	-0.211	-0.480	0.034	-0.132	-0.888	2.240	-2.417	3.721	-0.070	0.011	-0.070	0.314	0.093	-0.447	0.198
PW	0.032	-0.269	0.025	-0.130	-0.106	-0.032	-1.078	1.905	-1.686	1.461	-0.180	0.042	0.085	-0.477	-0.134	0.922	0.381
NGPP	-0.387	0.075	0.523	-0.316	0.002	-0.165	0.205	1.759	-2.093	0.348	-0.066	0.113	0.885	0.503	0.207	-1.092	0.500
FG	-0.476	0.273	0.584	-0.105	0.045	-0.306	0.661	1.352	-1.793	-0.270	-0.016	0.103	0.968	0.677	0.287	-1.417	0.569
GL	0.662	0.402	-0.067	-0.179	-0.126	0.285	-0.484	-0.510	0.799	-1.040	-0.076	-0.051	-0.583	-1.124	-0.32	1.941	-0.472
GB	0.429	0.240	-0.083	0.294	-0.060	0.045	-0.727	-0.479	0.879	-0.811	-0.056	-0.055	-0.650	-0.840	-0.428	1.788	-0.514
100GW	0.490	0.289	-0.184	0.147	-0.109	0.290	-0.633	-0.516	0.870	-0.830	-0.083	-0.062	-0.684	-1.089	-0.382	2.004	-0.479

**Note:** \* indicates significance at 5%, \*\* indicates significance at 1%, PH (50)= Plant height at  $50^{\text{th}}$  day, DFF= Days to fifty per cent flowering, SPAD value= SPAD meter value, PH= Plant height, PL= Panicle length, FL= Flag leaf length, FB= Flag leaf breadth, TT= Total tillers, PT= Productive tillers, DM= Days to maturity, PW= Panicle weight, NGPP= Number of grains per panicle, FG= Number of filled grains, GL= Grain length, GB= Grain breadth, 100GW=Hundred grain weight, SPY= Single plant yield



**Note:** PH (50)= Plant height at  $50^{\text{th}}$  day, DFF= Days to fifty per cent flowering, SPAD value= SPAD meter value, PH= Plant height, PL= Panicle length, FL= Flag leaf length, FB= Flag leaf breadth, TT= Total tillers, PT= Productive tillers, DM= Days to maturity, PW= Panicle weight, NGPP= Number of grains per panicle, FG= Number of filled grains, GL= Grain length , GB= Grain breadth, 100GW=Hundred grain weight, SPY= Single plant yield.



#### Conclusion

In this study, wide range of variation has been observed among the 36 rice genotypes. Among the quantitative characters studied, number of grains per panicle had highly significant and positive correlation with grain yield per plant but direct effect of this trait on single plant yield is comparatively low (0.113). In the case of correlation, number of filled grains and single plant yield had highly significant and its direct effect towards grain yield is high. The heritability (85.53%) and genetic advance (36.47%) of this trait is also very high. So selection of traits such as total tillers, productive tillers, number of grains per panicle and number of filled grains per panicle for further improvement in grain yield of rice.

#### References

- 1. Duwayri M, Tran DV, Nguyen VN. Reflections on yield gaps in rice production: how to narrow the gaps, In; Papademetriou *et al.* (Eds.), Bridging the rice yield gap in the asia-pacific region, FAO, 2000; 26-46.
- Tiwari DK, Pandey P, Tripathi S, Giri PS, Dwivedi JL. Studies on genetic variability for yield components in rice (*Oryza sativa* L.). Advances in Agriculture & Botanics. 2011; 3(1):76-81.
- Kishore NS, Srinivas T, Nagabhushanam U, Pallavi M, Sameera SK. Genetic variability, correlation and path analysis for yield and yield components in promising rice (*Oryza sativa* L.) genotypes. SAARC Journal of Agriculture. 2015; 13(1):9-108
- 4. Mishra SS, Behera PK, Panda D. Genotypic variability for drought tolerance-related morpho-physiological traits among indigenous rice landraces of Jeypore tract of Odisha, India. Journal of Crop Improvement. 2019; 33(2):254-278.
- 5. Hossain S, Salim Md, Azam G Md, Noman S. Variability, correlation and path analysis in drought

tolerant rice (*Oryza sativa* L.). Journal of Bioscience and Agriculture Research. 2018; 18(02):1521-1530.

- 6. Aditya JP, Bhartiya A. Genetic variability, correlation and path analysis for quantitative characters in rainfed upland rice of Uttarakhand hills. Journal of Rice Research. 2013; 6(2):24-34.
- Akinwale MG, Gregorio G, Nwilene F, Akinyele BO, Ogunbayo SA, Odiyi AC. Heritability and correlation coefficient analysis for yield and its components in rice (*Oryza sativa* L.). African Journal of plant science. 2011; 5(3):207-212.
- Kalyan B, Radha KKV, Subba RLV. Correlation coefficient analysis for yield and its components in rice (*Oryza sativa* L.) genotypes. Int. J Curr. Microbiol. App. Sci. 2017; 6(7):2425-2430
- Lalitha R, Mothilal A, Arunachalam P, Senthil N, Hemalatha G. Genetic variability, correlation and path analysis of grain yield, grain quality and its associated traits in EMS derived M4 generation mutants of rice (*Oryza sativa* L.). Electronic Journal of Plant Breeding. 2019; 10(3):1140-1147.
- Li R, Li M, Ashraf U, Liu S, Zhang J. Exploring the relationships between yield and yield-related traits for rice varieties released in China from 1978 to 2017. Frontiers in plant science. 2019; 10:543.
- 11. Awasthi S, Lal JP. Genotypic correlation and path analysis study in rice (*Oryza sativa* L.) Under irrigated and rainfed conditions. Annals of Plant Sciences. 2014; 3(12):916-920.
- Saha SR, Hassan L, Haque Md A, Islam MM, Rasel Md. Genetic variability, heritability, correlation and path analyses of yield components in traditional rice (*Oryza* sativa L.) landraces. Journal of the Bangladesh Agricultural University. 2019; 17(1):26-32.
- 13. Nithya N, Beena R, Stephen R, Abida PS, Jayalekshmi VG, Viji MM *et al.* Genetic Variability, Heritability, Correlation Coefficient and Path Analysis of

Morphophysiological and Yield Related Traits of Rice under Drought Stress. Chemical Science Review and Letters.2020; 9 (33): 48-54.

- 14. Evans, LT., and Dunstone, R.Ls. Some physiological aspects of evolution in wheat. Australian Journal of Biological Sciences. 1970; 23(4): 725-742.
- Parimala, K., Raju, S. CH., Prasad, H. AS., Kumar, S. S. and Reddy, N. S. Studies on genetic parameters, correlation and path analysis in rice (*Oryza sativa* L.). Journal of Pharmacognosy and Phytochemistry. 2020; 9(1): 414-417.
- Devi KR, Chandra BS, Lingaiah N, Hari Y, Venkanna V. Analysis of variability, correlation and path coefficient studies for yield and quality traits in rice (*Oryza Sativa* L.). Agricultural Science Digest-A Research Journal. 2017; 37(1):1-9.
- 17. Kishore C, Pal A, Kumar AV, Prasad V, Kumar ABD. Character Association and Path Analysis for Yield Components in Traditional Rice (*Oryza sativa* L.) Genotypes. International journal of current microbiology and applied sciences. 2018; 7(3):456-460.
- Devi KR, Chandra BS, Venkanna V, Hari Y. Variability, correlation and path studies for yield and quality traits in irrigated upland rice (*Oryza sativa* L.). Journal of Pharmacognosy and Phytochemistry. 2019; 8(6):676-684.
- Bhutta MA, Munir S, Qureshi MK, Shahzad AN, Kashif A, Hamid M *et al.* Correlation and path analysis of morphological parameters contributing to yield in rice (*Oryza sativa* L.) under drought stress. Pak. J Bot. 2019; 51(1):73-80.
- 20. Idris AE, Justin FJ, Dagash YMI, Abuali AI. Genetic variability and inter relationship between yield and yield components in some rice genotypes. Journal of Experimental Agriculture International. 2012; 2(2):233-239.
- Ashrafuzzaman M, Islam MdR, Ismail MR, Shahidullah SM, Hanafi MM. Evaluation of six aromatic rice varieties for yield and yield contributing characters. Int. J Agric. Biol. 2009; 11(5):616-620.
- 22. Asana RD. In quest of yield. Indian J. Plant Physiol. 1968; 11:1-10.
- 23. Rafiq MT, Aziz R, Yang X, Xiao W, Rafiq MK, Ali B. *et al.* Cadmium phytoavailability to rice (*Oryza sativa* L.) grown in representative Chinese soils. A model to improve soil environmental quality guidelines for food safety. Ecotoxicology and environmental safety. 2014; 103:101-107.
- 24. Sakamoto T, Matsuoka M. Identifying and exploiting grain yield genes in rice. Current opinion in plant biology. 2008; 11(2):209-214.
- 25. Srijan A, Kumar SS, Raju ChD, Jagadeeshwar R. Character association and path coefficient analysis for grain yield of parents and hybrids in rice (*Oryza sativa* L.). Journal of Applied and Natural Science. 2016; 8(1):167-172.