

SELECTION OF PROMISING LINES FROM F₄ GENERATION OF CHICKPEA GENOTYPES THROUGH CORRELATION AND PATH COEFFICIENT ANALYSIS

KEYWORDS

Chickpea
Correlation
Path coefficient analysis

ABSTRACT

The investigation was conducted with 45 genotypes of Chickpea comprising of 30 elite lines of F₄ generation, 12 parental lines and 3 checks in randomized block design. These genotypes which were evaluated for correlation among agronomical yield components and their direct and indirect effects on seed yield. The correlation coefficient analysis revealed that seed yield per plant exhibited a positive and significant correlation with plant height (0.2736, 0.2957), number of branches per plant (0.4912, 0.5492), pods per plant (0.4726, 0.4837), number of seeds per plant (0.4736, 0.4899), 100 seed weight (0.5892, 0.6193) and harvest index (0.4453, 0.4806) at both genotypic and phenotypic level. The character traits biological yield per plant and days to maturity depicted showed a positive but non-significant association and days to 50% flowering exhibited showed a negative and non-significant association with seed yield per plant at both genotypic and phenotypic level. The perusal of Path analysis indicated revealed that the high positive direct effect exhibited by branches per plant (0.2782) followed by 100-seed weight (0.2415), harvest index (0.2107) and number of pods per plant (0.1569) on grain yield at both genotypic and phenotypic level. Thus the study suggests that plant height, number of branches per plant, pods per plant, number of seeds per plant, 100 seed weight and harvest index are important traits for the selection criteria in crop improvement program of chickpea.

INTRODUCTION

The Chickpea (*Cicer arietinum* L.) is third most vital important pulse in the world (Padmavathi et al., 2013) which grown in over forty country which occupies upper position in the territory prominent position in cultivated areas (Dhingan et al., 2013). It is an important crop of rabi season in India. It has manifold multiple benefit for human and livestock as pod and seed coat can be utilized used as fodder. It provides good source of protein, energy, fiber, vitamin and minerals and considered a healthy food in many developed countries (Maiti, 2001; Nair and Mehta, 2014). It is also good source of amino acids like tryptophan and lysine (Awasthi et al., 1991). Apart from providing its dietary benefit to human beings, chickpea is very useful in management of soil fertility due to its nitrogen fixation ability (Gul et al., 2011).

Seed yield is the most important economic character trait and very complex character in nature because it is governed by polygenes and greatly influenced by environmental factors (Singh et al., 2014). Therefore, direct selection for yield per seed. Hence, it is necessary to consider the yield contributing traits during selection (Singh, 2006). The contribution of different character towards yield varies significantly different from each other. Thus, it is imperative to identify those characters,

which would play an important role during selection. The correlation and path co-efficient analysis provide valuable information about the importance of various yield components in formulation of appropriate selection strategy. Studies on above these

aspects of the available germplasm under the specific environment where it is to be exploited are essential for successful utilization of germplasm resources for the development in developing of superior chickpea varieties.

The present study was undertaken to find out determine the association among different character traits, their direct and indirect effect among yield and yield contributing characters and their contribution to define seed yield.

MATERIALS AND METHODS

The experiment was conducted at the Field Experimental Center of the Department of Genetics and Plant Breeding, Sam Higginbottom Institute of Agriculture Technology and Sciences, Allahabad in rabi from 2007-08 to 2011-12.

The experiment was carried out involved 45 genotypes of Chickpea (30 elite lines + 12 parents + 3 checks) which were grown in randomized block design with two replications. Six crosses (ICC-12234 x IPC-9494, BG-396 x ICC-11316, ICC-5742 x ICC-12234, IPC-94-94 x ICC-12234, ICC-596 x ICC-12234 and IC-264091 x ICC-596) were carried out made in 2007-08. Elite lines were selected from F_3 generation in 2009-10.

The observations like such as days to 50% flowering, plant height, number of branches per plant, number of pods per plant, number of seeds per plant, days to maturity, 100 seed weight, harvest index, seed yield per plant and seed yield per hectare were recorded from the F_4 generation in 2011-12 which were utilized to estimate correlation and path analysis. The statistical

Table1: Phenotypic correlation (above the diagonal) and genotypic correlation (below the diagonal) between different quantitative characters in Chickpea

No	Character	Plant Height	Daysto 50% Flowering	Branches/ Plant	Pods/ Plant	Daysto 50% maturity	Seeds/ Plant	Biological Yield/ Plant	Harvest Index 100	Seed Weight	Seedyield/ Plant
1	PlantHeight	1.0000	-0.2782*	0.3627*	0.2784*	0.1403	0.2457*	0.1075	-0.0127	0.2133*	0.2736*
2	Daysto50%Flowering	-0.3291*	1.0000	0.0067	-0.0813	0.0761	0.0155	0.1785	0.0152	0.0196	-0.0095
3	Branches/Plant	0.4074*	0.0038	1.0000	0.3287*	-0.1635	0.3394*	0.2069	0.2469*	0.5006*	0.4912*
4	Pods/Plant	0.2911*	-0.0889	0.3678*	1.0000	-0.0644	0.9106*	0.1718	0.1787	0.3961*	0.4726*
5	Daysto50%maturity	0.1470	0.0484	-0.1698	-0.0773	1.0000	-0.1208	0.0629	0.0239	-0.0620	0.0478
6	Seeds/Plant	0.2545*	0.0166	0.3725*	0.9159*	-0.1282	1.0000	0.1690	0.2676*	0.4138*	0.4736*
7	BiologicalYield/Plant	0.1014	0.2009	0.2327*	0.1762	0.0571	0.1743	1.0000	-0.2442*	0.0300	0.0855
8	HarvestIndex	-0.0019	0.0072	0.2550*	0.1921	0.0453	0.2778*	-0.2553*	1.0000	0.5523*	0.4453*
9	100SeedWeight	0.2198*	0.0206	0.5462*	0.4071*	-0.0641	0.4250*	0.0240	0.6050*	1.0000	0.5892*
	Seedyield/Plant	0.2957*	-0.0386	0.5492*	0.4837*	0.0590	0.4899*	0.0903	0.4806*	0.6193*	1.0000

Table2: Genotypic direct and indirect effect of component traits attributing to grain yield in Chickpea

No	Character	Daysto 50% Flowering	Plant Height	Branches/ Plant	Pods/ Plant	Daysto 50% maturity	Seeds/ Plant	Biological Yield/Plant	harvest Index	100Seed Weight
1	Daysto50%Flowering	-0.0365	0.0120	-0.0001	0.0032	-0.0018	-0.0006	-0.0073	-0.0003	-0.0008
2	PlantHeight	-0.0091	0.0277	0.0113	0.0081	0.0041	0.0071	0.0028	-0.0001	0.0061
3	Branches/Plant	0.0011	0.1133	0.2782	0.1023	-0.0472	0.1036	0.0647	0.0709	0.1519
4	Pods/Plant	-0.0140	0.0457	0.0577	0.1569	-0.0121	0.1437	0.0276	0.0301	0.0639
5	Daysto50%maturity	0.0064	0.0193	-0.0224	-0.0102	0.1316	-0.0169	0.0075	0.0060	-0.0084
6	Seeds/Plant	0.0014	0.0222	0.0324	0.0797	-0.0112	0.0870	0.0152	0.0242	0.0370
7	BiologicalYield/Plant	0.0056	0.0028	0.0065	0.0049	0.0016	0.0048	0.0278	-0.0071	0.0007
8	harvestIndex	0.0015	-0.0004	0.0537	0.0405	0.0095	0.0585	-0.0538	0.2107	0.1275
9	100SeedWeight	0.0050	0.0531	0.1319	0.0983	-0.0155	0.1026	0.0058	0.1461	0.2415

analysis for genotypic and phenotypic correlation was performed as per the method suggested by Singh and Chaudhari (1985) and path coefficient analysis were worked out according to following the methods described by Dewey and Lu (1959).

to maturity ($rp=0.0478$, $rg=0.0590$) depicted showed a positive but non-significant association with seed yield per plant at both genotypic and phenotypic level. The character traits daysto50%flowering ($rp=-0.0095$, $rg=-0.0386$) exhibited a negative and non-significant association with seed yield per plant at both genotypic and phenotypic

RESULTS AND DISCUSSION

Correlation coefficient

Correlation coefficient is one of the most important statistical analysis tools for amelioration the improvement of any crop. It provides information about the degree and magnitude of association among between the two variables which is may be due to pleiotropic gene action or linkage or both (Kumar, 2013). In the present investigation, genotypic and phenotypic correlation between seed yield and its component character traits were estimated to know understand the nature of their relationship presented in Table 1, which the results indicated that seed yield per plant exhibited significant positive correlation with plant height ($rp=0.2736$, $rg=0.2957$), number of branches per plant ($rp=0.4912$, $rg=0.5492$), pods per plant ($rp=0.4726$, $rg=0.4837$), number of seeds per plant ($rp=0.4736$, $rg=0.4899$), 100-seed weight ($rp=0.5892$, $rg=0.6193$) and harvest index ($rp=0.4453$, $rg=0.4806$) at both genotypic and phenotypic level. The similar finding was reported by Rao and Rao (2005) Yuces et al. (2006), Durga et al. (2007), Yuces and Anlarsal (2010) and Ali et al. (2011) in chickpea. While the characters traits biological yield per plant ($rp=0.0855$, $rg=0.0903$) and days

level.

The results indicate that the genotypic correlation ~~is~~
are higher than phenotypic correlation, which may be due to the
masking effect of environment on the expression of the
genotype. ~~similar~~ Similar findings were reported by Rai and
Dharmatti (2014) in clusterbean.

Correlation coefficient analysis revealed that seed yield exhibited significant and positive correlation at both genotypic and phenotypic level with 100-seed weight, branches per plant, seeds per plant, pods per plant and harvest index. Hence, direct selection for these traits could be helpful in the improvement improving of chickpea breeding.

Path coefficient analysis

The seed yield per plant is a complex character trait with influenced by many component characters, each having positive or negative contribution to the final manifestation expression. Further, the contribution

of any component character to seed yield may be direct and or indirect through other component characters and with its magnitude may also be different than differing from what is revealed — indicated by simple correlations. As a result, the direct and indirect effects of different casual characters on fruit seed yield per plant were estimated and are represented in Table 2. which The analysis indicated that the characters like plant height (0.0277), b Branches per plant (0.2782), number of pods per plant (0.1569), days to maturity (0.1316), seeds per plant (0.0870), biological yield (0.0278), harvest index (0.2107) and 100-seed weight (0.2415) exhibited positive direct effect on seed yield, while in contrast days to 50 per cent flowering exhibited negative direct effect (-0.0365) on seed yield per plant. These findings are in accordance with align with those of Khoragade et al. (1995), Kumare et al. (1999), Bhaduria et al. (2003), Rao and Rao (2005) and Yucesi et al. (2006) in chickpea.

In plant breeding, it is very difficult to have challenging to gain complete knowledge of the effects of all components characters on yield. A residual effect measures the role of other possible potential independent variables. In the present study, the residual effect was observed in present study were to be 0.6884 for genotypic path coefficient analysis.

The perusal of path analysis indicated that branches per plant, 100-seed weight, harvest index and pods per plant had high positive direct effect on grain yield at both genotypic and phenotypic levels. Therefore, these characters traits may serve as effective selection parameters in direct breeding programmes for yield improvement in Chickpea.

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