# Assessment of different drought tolerance indices and identifying drought tolerant lines in backcross population derived from cross between ICC13124 and WR-315 in chickpea

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**Abstract:** Drought is one of the major limiting factor in increasing the productivity of chickpea, as the crop is mainly grown under the residual soil moisture conditions under rainfed ecosystem. Hence, to identify most drought tolerant and productive lines in BC<sub>1</sub>F<sub>3</sub> population, drought tolerance indices were computed by taking yield under irrigated and rainfed conditions. The population was developed by crossing a drought tolerant genotype ICC-13124 as recurrent parent and susceptible parent WR-315 is also donor for wilt resistance. Totally 224 BC<sub>1</sub>F<sub>3</sub> families were subjected for evaluation under irrigated and rainfed situation to identify high yielding and drought tolerant lines based on drought tolerance indices. A drought tolerance index 'Drought and Productivity Index (DPI)' was computed along with other drought tolerance indices *viz*; DSI, TOL or TDS, MP, DTE, YI, RR, GMP and HM. Analysis of variance indicated presence of significance differences between families for seed yield under IR and RF as well as for drought tolerance indices. The highly significant and positive correlation of YIR and YRF with DPI, MP, YI, GMP and HM, indicates that selection could be effective for high DPI, MP, YI, GMP and HM. Superior genotypes can be selected by combining two to three indices and it may leads to bias in the selection if considering single index. Hence, recently reported Drought and Productivity Index (DPI) alone can be effectively used in identifying most productive and drought tolerant genotypes. Based on these indices out of 224 families 10 families were found superior and among these three families *viz*; BC<sub>1</sub>F<sub>3</sub>-4-1, BC<sub>1</sub>F<sub>3</sub>-6-3 and BC<sub>1</sub>F<sub>3</sub>-8-7 were found highly superior with respect to YIR, YRF and DPI compared to both the parents as well as checks.

Key words: Chickpea, Correlation, Drought tolerance indices, Population

#### Introduction

Chickpea is one of the most important crop in the semi arid tropics (SAT) region of the world. India stands first with respect to total world chickpea production followed by Turkey, Pakistan and Iran. In India, it tops the list of pulse crops and is cultivated in an area of 9.6 million ha, producing a total of 8.83 million tons with an average yield of 920.1 kg/ha (Anon., 2014). In SAT regions chickpea is mainly cultivated under rainfed, marginal and resource poor conditions. Under such situations crop usually faces terminal drought as it is cultivated in the residual soil moisture condition. Therefore it is necessary to develop drought tolerant superior genotypes with high yields.

Among the abiotic stresses, drought is most important constraint in improving productivity. Drought is very complex phenomena. It requires a clear understanding of the traits associated with it. Screening for drought tolerance will be very cumbersome because of its quantitative and complex nature. It is very difficult to screen the large number of genotypes for root traits and other physiological traits in field conditions to identify drought tolerant lines. Usually to identify drought tolerance genotypes researchers will go for evaluation of genotypes under irrigated and non-irrigated conditions and select the common lines which are high yielding in both the environments. But measurement of drought tolerance with single parameter will not give clear idea about the potential genotypes because of multiplicity of different factors involved and their interactions contributing to drought tolerance under field conditions. However, it is necessary for a breeder to identify not only drought tolerant line/genotype but also it should give high yield under favourable environments. Therefore, it is necessary to derive selection indices for drought tolerance based on a mathematical relation between yield under stress and non-stress condition along with yield of drought tolerant check to understand response of genotypes to drought. However, it is necessary to identify the most efficient and useful index to identify precisely the highly drought tolerant genotype. Keeping the above in view, an attempt was made to compute and compare different drought tolerance indices to identify efficiently drought tolerant lines in chickpea.

## Material and methods

The material used in this study includes  $BC_1F_2$  population developed by using two parents ICC13124 and WR-315. Parent ICC13124 (Parameswarappa et al., 2012, Shivukumar et al., 2013 and Jha et al., 2014) is drought tolerant and agronomically superior. It was used as recurrent parent and the second one WR-315 is a well known source of resistance was used as a donor for wilt resistance but it is susceptible to drought. Parents were crossed to generate F<sub>1</sub> seeds during Rabi 2010. All the F<sub>1</sub>s were sown in a field and by using molecular markers 18 true F<sub>1</sub>s were identified during off season kharif-2011. These true F<sub>1</sub>s were backcrossed to ICC13124 from which 55 BC, F, seeds were harvested. These seeds were sown in the field during rabi 2011 and 20 true BC<sub>1</sub>F<sub>1</sub>s were identified by using markers linked to fusarium wilt and were selfed to get 300 BC<sub>1</sub>F<sub>2</sub> seeds. These BC<sub>1</sub>F<sub>2</sub> seeds were sown during kharif 2012 and selfed to get  $BC_1F_3$  seeds from 240 healthy plants. Out of 240  $BC_1F_3$  families 224 families having sufficient seeds to replicate in two environments were selected and were evaluated during rabi-2012-13 under both irrigated and rainfed conditions in Botany Garden, Department of Genetics and Plant Breeding, UAS, Dharwad, Karnataka, India. Material was evaluated by following augmented design consisting of 14 blocks and 20 entries in each block including two parents and two local checks. By taking yield of lines under irrigated and rainfed conditions, following drought tolerance indices were computed.

The Drought and Productivity Index (DPI) defined by Ontagodi T. P. (2015), was computed as follows.

$$DPI = \frac{(3YRF + 1YIR - 2DCRF)}{2}$$

Where, YRF - Grain yield of genotype under rainfed condition YIR - Grain yield of genotype under irrigated condition

> DCRF - Grain yield of drought tolerant check under rainfed

Earlier reported drought indices like Drought Susceptibility Index (DSI) of Fischer and Maurer (1978), the Drought Tolerance Efficiency (DTE %) of Fischer and Wood (1981), the Mean Productivity (MP) and TOL or Tolerance to Drought Stress (TDS) of Rosielle and Hamblin (1981), Geometric Mean Productivity (GMP) and Harmonic Mean (HM) of Fernandez (1992), Yield Index (YI) of Gavuzzi et al. (2006) and Relative Reduction (RR) of Sadiki (2006) were computed for selecting drought tolerant lines. Data on yield under irrigated and rainfed along with drought tolerance indices were analysed by using WINDOSAT version 9.1.

## **Results and discussion**

Analysis of variance indicated significant differences among the families and significant differences were also found between families and checks for yield under rainfed. But there was no significant difference among genotypes as well as varieties for yield under irrigated condition. This indicates that genotypes thrown a differential expression under rainfed compared to irrigated condition. There was no significant difference between the blocks indicating that existence of sufficient uniformity across blocks and repetition of checks in each block (Table 1).

Extent of genetic variability expressed by the traits indicates scope for selection of such traits. Existence of variability for yield under rainfed and irrigated in  $BC_1F_3$  indicates scope for selection of favourable genotypes based on the drought tolerance indices (Table 2). This can be evidenced by expression of higher GCV under rainfed (37.79) compared to irrigated (20.89) situation. Only estimation of variability may not give clear idea about the heritable portion of the trait therefore, heritability and genetic advance needs to be estimated. Seed yield per plant under rainfed situation showed highest heritability of 67.84% followed by yield index (66.67%). But yield under irrigated situation recorded least heritability of 23.25%. This indicates that existence of more phenotypic variance because of environmental influence on genotype expression. Heritability indicates only the effectiveness with which selection of genotypes can be based on their phenotypic performance but fail to indicate the amount of progress expected from selection. Therefore, heritability estimates appear to be more meaningful when accompanied by estimates of genetic advance. Highest GA (5.22) and GAM (64.12) were recorded for yield under rainfed (YRF) followed by HM (GA; 4.46 and GAM; 49.40).

Block (eliminating Check+Var.)       13       12.41       8.84       9.45       0.89       4.53       9.49       431.96       0.13*       0.043       9.39       9.3         Sintries (ignoring Blocks)       227       20.17       13.93***       17.50**       0.55       4.64       16.03**       267.54       0.21***       0.027       15.86**       15         Sintries (ignoring Blocks)       3       2.52       0.98       1.48       0.06       1.16       1.46       27.57       0.01       0.003       1.37       1.2         Anietics       233       19.79       13.4***       17.20**       0.56       4.71       15.57*       270.92       0.02***       0.027       15.38**       16.         Anietics       1       15.57*       235.15       2.4***       0.027       15.38**       16.         Checks vs. Varietics       1       15.9**       235.15       2.4***       0.023       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1***       166.1**	ource of variations	$\mathrm{DF}$	YIR	YRF	DPI	DSI	TDS	МΡ	DTE	YI	RR	GMP	ΗМ
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	lock (eliminating Check+Var.)	13	12.41	8.84	9.45	0.89	4.53	9.49	431.96	$0.13^{*}$	0.043	9.39	9.36
Thecks       3       2.52       0.98       1.48       0.06       1.16       1.46       27.57       0.01       0.003       1.37       1.2         /arieties       223       19.79       13.4***       17.20**       0.56       4.71       15.57*       270.92       0.20***       0.027       15.38**       15.         Arieties       1       157.8**       160.0***       171.0**       0.49       0.01       162.9***       235.15       2.4***       0.023       166.1***       16         Anoto       0.01       162.9***       235.15       2.4***       0.023       166.1***       16	ntries (ignoring Blocks)	227	20.17	$13.93^{***}$	$17.50^{**}$	0.55	4.64	$16.03^{**}$	267.54	$0.21^{***}$	0.027	$15.86^{**}$	$15.78^{**}$
Arietics       223       19.79       13.4***       17.20** $0.56$ $4.71$ 15.57* $270.92$ $0.20***$ $0.027$ 15.38** $15$ Checks vs. Varietics       1       157.8**       160.0***       171.0** $0.49$ $0.01$ 162.9***       235.15 $2.4***$ $0.023$ 166.1*** $166$ Concorr       0.02       0.02       0.02       0.02 $0.02$ $0.027$ $156$ $0.02$ $0.02$ $0.023$ $166.1***$ $166$ Concorr       0.02       0.02       0.02 $0.02$	hecks	3	2.52	0.98	1.48	0.06	1.16	1.46	27.57	0.01	0.003	1.37	1.27
Thecks vs. Varieties 1 157.8** 160.0*** 171.0** 0.49 0.01 162.9*** 235.15 2.4*** 0.023 166.1*** 160 models and a second	arieties	223	19.79	$13.4^{***}$	$17.20^{**}$	0.56	4.71	15.57*	270.92	$0.20^{***}$	0.027	$15.38^{**}$	$15.28^{**}$
	hecks vs. Varieties	1	$157.8^{**}$	$160.0^{***}$	$171.0^{**}$	0.49	0.01	$162.9^{***}$	235.15	2.4***	0.023	$166.1^{***}$	$169.1^{***}$
1./ 8/./ 0.0.0 /0.0 %24 8.48 8.00 0.70 59.49 501.79 0.0 %24 1.0 %24 1.0 %24 1.0 %24 1.0 %24 1.0 %24 1.0 %24 1.0	RROR	39	15.48	4.48	8.50	0.75	5.93	8.49	361.79	0.07	0.036	7.78	7.17

Table 2. Estimation of genetic parameters for yield under irrigated and rainfed and drought tolerance indices BC<sub>1</sub>F<sub>3</sub> populations of a cross (ICC-13124xWR-315)xICC-13124 in chickpea 43.95 ΗM GMP 31.06 43.52 50.95 48.77 84.47 33.33 0.11 58.00 RR 37.39 45.80 66.67 Z DTE 12.05 20.31 35.23 43.18 47.04 29.62 МΡ 96.39 27.80 50.82 TDS 50.58 83.88 36.36 0.56 62.83 DSI 34.5 19.5 50.5 0.65 53.5 DPI 45.88 67.84 37.79 YRF 20.8943.31 during rabi 2012-13. YIR **Genetic Parameters** PCV GCV

54.56

4.46

9.40

45.67 4.18

62.90 0.63

11.87

3.88 41.84

55.20

.23

5.22

23.25

h<sup>2</sup> (%)

GA @

2.15 20.75

5% 5%

GAM@

### Assessment of different drought tolerance indices and ......

Determining the associations among various traits is useful to breeders in selecting genotypes in desired direction. A strong positive correlation for the seed yield per plant between irrigated and rainfed situation was recorded (Table 3). This indicates that selecting better yielding genotypes under rainfed can also give higher yields under the irrigated situation. Yield under irrigated (YIR) recorded highly significant positive correlation with drought tolerance indices like DPI (0.91), DSI (0.46), MP (0.96), YI (0.81), RR (0.46), GMP (0.94) and HM (0.92). On the contrary, yield under rainfed (YRF) also showed significant positive correlation with DPI (0.98), MP (0.93), YI (1.0), GMP (0.95) and HM (0.96) but it is negatively correlated with DSI (-0.06) and RR (-0.06). These results are in accordance with Fernandez (1992) in mungbean, Farshadfar and Sutka (2002) in maize and Win (2011), Talebi et al. (2011) and Sabaghnia and Janmohammadi (2014) in chickpea. Similarly, TDS showed negative correlation of -0.74 and -0.21 with YIR and YRF respectively. Computation of TOL or TDS gives negative values. Therefore based on the correlation with yield under IR and RF, selection for lower negative TDS values needs to be done. Another index DTE had negative correlation with YIR (-0.46) but showed positive non-significant correlation with YRF (0.06). Therefore, selecting genotypes having higher mean values for MP, YI, GMP and HM can give high yielding drought tolerant lines. In similar way several studies indicate that the GMP, HM, STI, YI and MP indices are preferred in late drought condition for selecting the most favourable genotypes (Sio-Se Mardeh et al., 2006; Talebi et al., 2011, Pouresmaeil et al., 2012 and Sabaghnia and Janmohammadi, 2014). Selection for lower values of DSI and TDS leads to selection of genotypes with low yield potential under non-stress conditions and high yield under stress conditions (Fernandez, 1992). Selection of genotypes based on single index leads to bias in selection of drought tolerant genotypes hence selection should be made by combination of 2 or 3 indices. Mean productivity is resulted in significantly positive association with all the drought tolerance indices except TDS and DTE. So selection for higher MP along with lower values of TDS and higher values of DTE as it is positively correlated with YRF is going to yield best drought tolerant genotypes. Whereas, the DPI reported by Ontagodi (2015) can alone be used for identifying most

productive and drought tolerant lines in the population or germplasm evaluations.

Superior genotypes under irrigated and rainfed situation were isolated from  $BC_1F_3$  population and their calculated drought tolerance indices is presented in Table 4. Among the selected genotypes BC<sub>1</sub>F<sub>3</sub>-6-3 (24.2), BC<sub>1</sub>F<sub>3</sub>-7-7 (22.10), BC<sub>1</sub>F<sub>3</sub>-8-7 (22.50), BC<sub>1</sub>F<sub>3</sub>-11-3(24.30) and BC<sub>1</sub>F<sub>3</sub>-18-8(22.30) recorded high seed yield per plant(g) under irrigated conditions compared to both parents and checks. But, under rainfed situation BC<sub>1</sub>F<sub>3</sub>-4-1(18.20), BC<sub>1</sub>F<sub>3</sub>-6-3 (21.20), BC<sub>1</sub>F<sub>3</sub>-8-7 and BC<sub>1</sub>F<sub>3</sub>-34-4 (15.50) performed better with respect to seed yield per plant(g). One genotype  $BC_1F_3$ -6-3 (24.2) performed better in both environments. Even though BC<sub>1</sub>F<sub>3</sub>-7-7, BC<sub>1</sub>F<sub>3</sub>-11-3 and BC<sub>1</sub>F<sub>3</sub>-18-8 gave high yields under irrigated and having higher MP values but with respects other drought tolerance indices these families showed lower values and maximum yield reduction under RF. Many reports showed reduction in yield and other related traits may be because of limited moisture available at critical stages in chickpea. So, selection of these families based on mean performance will lead to bias in selecting for drought tolerance. On the contrary, BC<sub>1</sub>F<sub>3</sub>-6-3 family showed maximum yield under both IR and RF along with higher values of drought tolerance indices like MP (22.70), DTE (87.60), YI (2.61), GMP (22.65) and HM (22.60) and lower values for indices like DSI (0.56), TDS (-3.0) and RR (0.12) compared to both the parents and checks. Similarly, selecting superior genotypes based on these indices have been reported by Win (2011), Ganjeali et al. (2011), Talebi et al. (2011), Ulemale et al. (2013) and Sabaghnia and Janmohammadi(2014) in chickpea. Another two families  $BC_1F_3$ -4-1(18.20) and  $BC_1F_3$ -34-4 (15.50) were not so high yielding under irrigated but recorded consistence performance under both environments. Therefore, these two families were superior with respect to DTE (94.30 and 90.38) and YI (2.24 and 1.91) and showed maximum lower values for DSI (0.26 and 0.44), TDS (-1.10 and -1.65) and RR (0.06 and 0.10) compared to drought tolerant parent ICC13124. Parameshwarappa et al., 2012 and Shivukumar et al., 2013 reported that the line ICC-13124 is having more DTE and showed low DSI, similarly in the present study ICC-13124 also recorded superiority for drought tolerance indices. Hence combination of DSI, TDS and RR can be used for selection of superior genotypes under rainfed condition

Table 3. Phenotypic correlation	ion coefficients for yield unde	er irrigated and rainfed and	d drought tolerance indice	es in $BC_1F_3$ population of	a cross
(ICC-13124xWR-31	15)xICC-13124 in chickpea d	luring rabi 2012-13			

Characters	YIR	DPI	DSI	TDS	MP	DTE	YI	RR	GMP	HM
YIR	1.00									
DPI	0.91***	1.00								
DSI	0.46 ***	0.13	1.00							
TDS	-0.74***	-0.44**	-0.84***	1.00						
MP	0.96***	0.98***	0.25***	-0.55***	1.00					
DTE	-0.46***	-0.13	-1.00***	0.84***	-0.25***	1.00				
YI	0.81***	$0.98^{***}$	-0.06	-0.21***	0.92***	0.06	1.00			
RR	0.46***	0.13	0.99***	-0.84***	0.25***	-0.99***	-0.06	1.00		
GMP	0.94***	0.99***	0.21**	-0.49***	0.99***	-0.21**	0.95***	0.21**	1.00	
HM	0.92 ***	0.99***	0.17*	-0.44***	0.99***	-0.16 *	0.96***	0.17*	0.99***	1.00
YRF	0.81	0.98***	-0.06	-0.21	0.93	0.06	1.00	-0.06	0.95	0.96

\*Significance at 5%, \*\* Significance at 1%, \*\*\* Significance at 0.1%

Table 4 Isolation of superior and drought tolerant BC<sub>1</sub>F<sub>3</sub> families based on the yield under irrigated and rainfed situation and drought tolerance indices

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Family Name	YIR (g)	YRF(g)	DPI	DSI	TDS	MP (g)	DTE	YI	RR (g)	GMP (g)	HM (g)
BC <sub>1</sub> F <sub>3</sub> -2-2	17.4	13.4	13.97	1.05	-4.0	15.43	76.79	1.65	0.23	15.29	15.16
BC <sub>1</sub> F <sub>3</sub> -4-1	19.3	18.2	22.10	0.26	-1.1	18.75	94.3	2.24	0.06	18.74	18.73
BC <sub>1</sub> F <sub>3</sub> -6-3	24.2	21.2	29.05	0.56	-3.0	22.70	87.6	2.61	0.12	22.65	22.6
BC <sub>1</sub> F <sub>3</sub> -7-7	22.1	11.1	12.85	2.26	-11.0	16.60	50.23	1.37	0.5	15.66	14.78
BC <sub>1</sub> F <sub>3</sub> -8-7	22.5	14.6	18.30	1.6	-7.9	18.55	64.89	1.80	0.35	18.12	17.71
BC <sub>1</sub> F <sub>3</sub> -11-3	24.3	13.2	17.10	2.08	-11.1	18.75	54.32	1.62	0.46	17.91	17.11
BC <sub>1</sub> F <sub>3</sub> -18-8	22.3	12.4	14.90	2.02	-9.9	17.35	55.61	1.53	0.44	16.63	15.94
BC <sub>1</sub> F <sub>3</sub> -34-4	17.1	15.5	16.97	0.44	-1.6	16.33	90.38	1.91	0.1	16.3	16.28
BC <sub>1</sub> F <sub>3</sub> -37-4	16.5	14.6	15.30	0.52	-1.9	15.55	88.48	1.80	0.12	15.52	15.49
BC <sub>1</sub> F <sub>3</sub> -39-1	16.0	14.0	14.22	0.56	-1.9	15.02	87.78	1.73	0.12	14.99	14.96
JG-11 (LC)	15.4	13.1	12.57	0.67	-2.2	14.27	85.35	1.62	0.15	14.23	14.19
BGD 103 (LC)	14.5	11.9	10.38	0.8	-2.5	13.26	82.43	1.47	0.18	13.19	13.13
ICC13124 (P1)	15.4	14.8	15.15	0.18	-0.6	15.15	96.12	1.83	0.04	15.15	15.14
WR315 (P2)	12.7	7.5	2.82	1.85	-5.1	10.13	59.28	0.93	0.41	9.79	9.47
Mean	18.56	13.98	15.41	1.06	-4.58	16.27	76.68	1.72	0.23	16.01	15.76
S.Em.	1.03	0.85	1.57	0.20	1.01	0.79	4.35	0.10	0.04	0.79	0.79

rather than the irrigated, similar results were reported by Rehman (2009) and Win (2011) in Chickpea. Whereas the families like  $BC_1F_3$ -6-3,  $BC_1F_3$ -4-1 and  $BC_1F_3$ -8-7 were found superior for DPI and most productive under both environments. Even though families  $BC_1F_3$ -11-3 and  $BC_1F_3$ -18-8 showed higher DSI, TDS and RR but superior with respect to DPI, hence these families can also be selected as most productive and drought tolerant ones. Families  $BC_1F_3$ -4-1 and  $BC_1F_3$ -11-3 showed similar MP (18.75) but superiority of these families can be differentiated based on DPI values.  $BC_1F_3$ -4-1 was found superior for DPI(22.10) compared to  $BC_1F_3$ -11-3(17.10).

Screening of large number of genotypes for root related traits to identify drought tolerant lines will be very difficult and cumbersome. Hence, to get idea about the performance of line under irrigated and rainfed and its drought tolerance ability, drought tolerance indices are computed based on yield under

irrigated and rainfed. Strong correlation between YIR and YRF and with drought indices indicates that indirect selection could be effective for drought tolerance. There is a positive and strong correlations exist between DPI, MP, GMP, HM and YI with YIR and YRF. Mere selection based on the mean performance of genotype may not give clear idea about the genotype under drought. Therefore, the parameters like DPI, MP, DTE, GMP, HM and YI can be suggested to select high yielding drought tolerant genotypes along with lower values of DSI and RR. Based on the drought tolerances indices 10 superior families were identified in  $BC_1F_3$  population. Among the selected entries BC<sub>1</sub>F<sub>3</sub>-6-3 was found very superior transgressive segregant followed by  $BC_1F_3$ -4-1 and  $BC_1F_3$ -8-7 with respect all the drought tolerance indices as well as DPI. Hence, these genotypes can be used in the future breeding programmes as tolerant sources. Continuous selfing of these families gives stable genotypes and may be released as drought tolerant varieties.

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