

Research Article GROWTH VARIATION AMONG PROGENIES OF Acacia nilotica CPTS OF DIFFERENT PROVENANCES

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Abstract: Acacia nilotica, belonging to family Fabaceae, is one of the most important, versatile and multipurpose tree species. Sufficient quantity of quality pods was collected from ten phenotypically superior candidate plus trees (CPTs) from six provenances/geographical locations to test the performance of progeny of individual tree selected. Significant variation was observed in progenies for field emergence, shoot length, root length, fresh seedling weight, numbers of branches per seedling, root-shoot ratio and straightness, open a way to go ahead for further improvement in tree species through selection.

Keywords: shoot length, root length, fresh seedling weight

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Introduction

Acacia nilotica, belonging to family Fabaceae, is one of the most important, versatile and multipurpose tree species found in dry areas of the Indian subcontinent and Africa. The species seldom occurs above 500 m or in areas with more than annual 1500 mm of rainfall, apart from on harsh porous soils on the river bed sand can grow on a various soil like compact sandy loam, shallow stony, riverine alluvial, black cotton, alluvial loam, saline, mild alkaline, ravines and soils comprising of calcareous concretions. Genetic variation in Acacia nilotica has mainly been investigated using morphological and physiological traits. This differentiation is usually pertinent to habitat conditions. Morphological variation is strongly influenced by various environmental factors and morphological variability signifies the adaptation of the species to its environment and it may be genetically determined or environmentally introduced. Thus, morphological differentiation seems to have an evolutionary significance that result in the gradual adaptation of the species to its environment [1][2]. Acacia nilotica has wide ecological amplitude and is found growing in a variety of habitats. The species is remarkable variable for various morphological expressions such as leaf, stipular spines morphology, pod and seed characteristics. The population of the species can be differentiated even at a very short distance. The available literature reveals that a little work has been done on growth variation in Acacia nilotica. Keeping in view the above facts, the present study was initiated to find extant growth variation in progenies of CPT's of Acacia nilotica among different geographical.

Materials and Methods

A survey was conducted in six geographical locations during June 2014 for collection of *Acacia nilotica* quality pod [Table-1]. Randomly ten phenotypically superior candidate plus trees (CPTs) were selected from each geographical area on the basis traits of economic importance *viz*. clear bole height, straightness, self-pruning ability, low branching habit, disease resistance *etc.*, keeping an isolation distance of 200 m having true representation of each geographical area. Adequate amount of quality pods was collected from each selected tree and were kept distinctly in cloth bags for further study purpose. To test the progeny of individual tree selected from six provenances, 300 seeds in three replications were sown in polythene bags filled with equal proportions of sandy soil and FYM.

The experiment was conducted in nursery of Department of Forestry, CCS Haryana agricultural University, Hisar during second week of July 2015. The poly bags soon after seed sowing were irrigated with the help of garden sprinkler and thereafter, proper moisture was maintained as and when considered necessary. The field emergence percentage was observed by counting the number of seeds successfully germinated under each replication. Observations on field emergence were recorded regularly upto 21 days of sowing. Fifteen plants per progeny per replication were selected at random for growth observations. The plant height and collar diameter were recorded at an interval of three months and final observations were recorded at the age of 12 months. Phenotypic, genotypic and environmental coefficients of variation were calculated [3]:

Phenotypic coefficient of variation % (PCV) = $\sqrt{PV/x \times 100}$ Genotypic coefficient of variation % (GCV) = $\sqrt{GV/x \times 100}$

x = population mean for each trait

Heritability values and genetic advance were worked out following the methodology of Johnson, *et al.* [4]. Genetic advance (GA) is the estimated increase in the extent of a particular character when a selection pressure of chosen intensity (i) is applied. This was calculated as per Johanson, *et al.* [4].

 $GA = GV/PV \times i \times \sqrt{PV}$ Where, i = selection intensity

Results and Discussion

Selection exploits the natural variability existing within a population of the chosen tree species. The extent and nature of heritable variation in any tree species could be assessed through progeny testing. The germination and the plant growth characters differed significantly among progeny of CPT's [Table-2]. The germination percentage varied from 25 to78 with the general mean of 46.51. The seed lot from RH7 tree from Hanumangarh provenance showed highest (78%) germination followed by, PF1 (75%) from Firozpur provenance. However, minimum (25%) percent germination was recorded in PR6 from Roopnagar provenance. The collar diameter ranged from 6.61 in PR5 progeny to 10.03 in HS1 (Roopnagar) progeny (Sonipat) at the age of 9 months. Similar variation was observed in collar diameter of different progenies.

Growth Variation Among Progenies of Acacia nilotica Cpts of Different Provenances

Table-1 Geographical information of Acacia nilotica provenance	es
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Dictrict	State				Altitude	(M) Dei	ofall (mm)			
	Dunich			Altitude	(W) Ral	11dll (11111)				
Perozpur	Punjab Punjab		njab 31 /5 11			9/ m /31 mm				
Rupnagar	Punjab	ab 31 77 262			n 776 mm					
Hanumangarh	Rajasthan 29			75	181 m	1	241 m			
Sonipat Haryana		29		77 224 m			653 mm			
Nagaur	Rajasthan	27		75 295 m			310 mm			
Dausa	Rajasthan	27		77	342 m	า 5	98 mm			
Table-2 Germination and performance of different progenies of A. nilotica at different stages of growth in nurserv										
Seed source Germination (%)Collar diameter (mm)Shoot length (cm)										
		3 month	6 month	9 month	3 month	6 month	9 month			
PF1	75.00	2 11	3.35	7 28	23.36	39.72	103 38			
PF2	40.67	2.68	4 37	7.66	20.00	38.41	106.50			
DE3	56.00	1 80	4.07	6 07	10 0/	2/1 2/1	85.40			
DE/	25.67	2.24	4.07	8 NQ	21 17	28.04	100 71			
DE5	32.07	2.24	3.02	7 69	17.05	30.34	103.71			
PEG	52.00	2.00	3.90	7.00	10.05	25.40	100.30			
	32.07	2.90	4.00	0.02	19.00	20.25	109.79			
	30.00	2.14	4.00	0.03	20.07	30.30	120.51			
PFO	43.07	2.12	5.02	0.00	20.12	43.11	121.90			
PF9	31.07	2.45	4.13	7.41	22.02	34.07	105.44			
PF10	48.00	2.90	4.82	8.90	24.26	44.//	122.52			
PR1	43.67	2.29	4.00	7.45	22.80	40.75	120.05			
PR2	39.67	2.37	3.70	7.80	22.99	33.96	90.92			
PR3	64.00	2.39	4.49	8.35	21.76	36.55	108.45			
PR4	39.67	2.35	4.01	8.19	21.91	37.83	125.59			
PR5	50.00	2.42	3.70	6.61	21.12	30.12	86.88			
PR6	25.00	2.25	3.59	7.44	24.54	36.64	115.84			
PR7	60.67	2.35	4.29	7.70	21.29	38.00	124.40			
PR8	50.00	2.56	4.66	8.61	20.99	35.05	109.28			
PR9	67.67	2.12	3.86	7.21	22.53	33.84	113.44			
PR10	51.33	2.31	4.03	8.13	21.37	36.86	122.96			
HS1	57.00	3.43	5.79	10.03	26.18	46.69	140.48			
HS2	42.00	3.19	5.21	8.20	22.57	35.91	107.81			
HS3	57.00	3.31	5.14	8.26	26.80	40.98	115.58			
HS4	48.67	3.09	4.75	8.01	23.06	34.86	112.46			
HS5	28.33	2.79	4.57	7.41	24.94	40.47	121.47			
HS6	53.67	3.39	5.40	9.10	27.23	43.51	116.01			
HS7	55.00	2.32	4.20	7.30	18.90	38.28	100.68			
HS8	28.67	2 68	4 65	8 24	20.08	37 21	107.01			
HSQ	41 00	2.00	4 18	7 46	25.68	35.14	106.44			
HS10	43.67	3 50	5 58	8 70	25.00	45.08	122.08			
RD1	35.67	2 71	4 55	8 35	24 95	47.21	122.00			
RD2	28.67	2.11	3 80	7 9/	10.61	30.76	96.76			
RD2	20.07	2.21	1.09	8.24	25.90	10.20	122.10			
	21.00	2.00	4.09	0.01	23.09	40.30	122.10			
	50.00	2.99	4.07	0.00	24.22	44.42	120.19			
	09.33 40.00	J.20 0.00	4.04	1.01	29.00	44.00	137.00			
	42.00	2.02	4.42	0.40	25.04	43.33	100.00			
	30.07	2.04	4.03	0.5Z	20.90	44.73	127.23			
RD8	31.33	2.89	4.84	0.11	21.8/	48.41	123.67			
KD9	29.33	2.85	4.38	8.07	29.54	43.85	123.44			
RD10	55.33	2.49	4.22	7.95	29.44	42.21	123.23			
KN1	29.67	2.20	4.00	/.61	17.90	35.56	109.96			
RN2	59.33	2.62	4.44	8.42	22.81	38.13	109.73			
RN3	44.33	1.97	3.76	7.57	19.21	34.66	97.16			
RN4	56.00	2.66	4.45	8.55	23.99	41.29	125.29			
RN5	48.00	2.22	4.09	7.49	20.29	37.68	118.10			
RN6	58.00	2.53	4.14	7.96	22.45	40.00	116.80			
RN7	63.00	2.91	4.66	7.82	25.13	42.77	129.18			
RN8	27.00	2.81	4.58	8.94	25.54	42.29	128.70			
RN9	40.00	3.13	4.89	8.19	29.85	45.48	118.68			
RN10	45.00	3.10	5.00	8.50	26.53	43.47	119.52			
RH1	60.00	2.40	4.23	8.06	20.86	37.99	110.59			
RH2	48.33	2.90	4.65	8.49	21.85	34.96	106.18			
RH3	61.33	2.51	4.31	8.37	22.20	34.96	93.55			
RH4	46.67	2.42	4.17	9.15	22.17	38.74	102.14			
RH5	57.00	2.35	4.13	8.39	22.07	40.46	136.86			
RH6	64.00	1.79	3.50	7.44	22.02	38.44	116.54			
RH7	78.00	2.19	3.77	7.93	22.36	39.13	113.13			
RH8	44.00	2.53	4.38	8.95	16.94	29.90	107.30			
RH9	37,33	2.45	4.05	7,67	22 34	36 64	114 04			
RH10	56.33	3.33	5.06	9.41	28.17	42 34	113.94			
Range	25 00-78 00	1 79-3 50	3 35-5 70	6 61-10 03	16 94-29 88	29 9-48 41	85 4-140 8			
MEAN	<u>46</u> 51	2.63	4 40	8 08	23.25	39.00	114.87			
CD at 5%	8 80	0.40	0.78	0.84	4 78	6 64	7 51			

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Onest	Observations	Destination	No. of		Dest shart			The state of the second st	Descentilities
Seed	Shoot length	Root length	NO. Of	Straight-ness	Root-shoot	Shoot weight	Root weight (g)	Fresh seedling	Dry seedling
source	(cm)	(cm)	branches		ratio	(g)		weight (g)	weight (g)
PF1	150.88	70.00	8.00	4.00	0.47	373.45	92.12	465.57	209.49
DE2	156.81	82.60	10.00	3 33	0.53	325.56	100.58	126.14	217 52
FTZ	100.01	02.00	10.00	0.00	0.00	323.30	100.30	420.14	217.JZ
PF3	129.40	54.90	11.67	2.83	0.42	340.56	102.79	443.35	236.00
PF4	170.31	92.40	10.33	3.67	0.54	342.75	98.48	441.23	216.85
PF5	150.88	66 90	8.33	4 33	0 44	318.96	92 52	411 48	202 81
	100.00	75.00	10.00	4.00	0.47	010.00	00.42	411.40	202.01
PF0	101.07	75.90	10.00	3.50	0.47	301.52	98.43	459.95	207.94
PF7	177.21	95.40	9.67	3.33	0.54	375.24	111.44	486.68	234.16
PF8	179 79	102 90	8 00	4 33	0.57	425 15	90.63	515 78	242 59
	164.64	106.50	0.00	1.00	0.60	240 56	106.66	455.00	212.00
PF9	104.04	100.00	0.33	4.00	0.69	340.30	100.00	400.22	213.41
PF10	180.68	98.40	13.00	2.50	0.54	415.46	115.47	530.93	236.96
PR1	182.15	107.60	9.00	4.00	0.59	360.49	114.39	474.88	240.07
DD2	138 72	78.20	10.00	3 33	0.56	362.40	103.06	165.46	233.46
DD2	150.72	70.20	10.00	0.50	0.00	102.40	100.00	403.40	200.40
PR3	158.66	72.80	12.00	2.50	0.46	409.47	100.69	510.16	246.50
PR4	182.99	87.40	10.33	3.33	0.48	359.35	95.74	455.09	229.96
PR5	133 18	83.00	8.00	4 17	0.62	348 59	100 71	449 30	237.60
	160.10	70.00	0.00	2.50	0.47	260.05	115.00	470.00	201.00
PRO	100.00	70.00	9.55	3.50	0.47	302.45	115.00	470.20	200.20
PR7	182.91	84.50	10.67	3.00	0.46	405.46	100.26	505.72	243.35
PR8	166 96	74 70	12 67	2 33	0 45	410 45	107 20	517 65	242 13
DD0	161.44	52.30	10.00	3.00	0.32	340.54	05.80	115 31	220.67
FR9	101.44	52.50	10.00	3.00	0.32	349.34	95.00	440.04	230.07
PR10	170.83	59.00	10.00	2.83	0.35	413.47	100.50	513.97	248.20
HS1	201.28	102.40	14.00	2.17	0.51	416.53	107.62	524.15	249.16
HS2	152 71	82.50	12.00	3.00	0.54	386.42	00.13	185 55	2/8 50
1102	132.71	02.00	12.00	0.00	0.04	070.50	33.13	403.33	240.00
HS3	173.48	66.90	11.00	3.00	0.39	378.56	102.56	481.12	222.29
HS4	170.36	92.50	10.00	3.17	0.54	355.66	100.19	455.85	233.94
HS5	183 37	104.30	8.00	4 00	0.57	358 79	100.80	459 59	222.81
1100	170.07	70 00	12.00	9.47	0.44	404.00	110.00	-100.00 F24.00	000.07
H20	173.11	76.90	13.00	Z.17	0.44	424.98	110.00	534.98	200.37
HS7	154.58	105.50	9.00	3.67	0.68	366.64	120.12	486.76	235.70
HS8	166.33	110.00	11.00	2.67	0.66	411.35	98.47	509.82	238.13
021	165.77	02.70	1/ 00	1.93	0.56	385.46	103 58	180 04	22/ 12
1109	103.77	92.10	14.00	1.05	0.00	303.40	103.30	409.04	204.12
HS10	184.38	110.50	13.00	2.50	0.60	3/2.4/	122.41	494.88	251.80
RD1	192.70	90.60	15.00	2.00	0.47	420.41	98.40	518.81	254.15
RD2	146.06	74 90	13.00	2.50	0.51	422 51	101 46	523 97	247 55
	170 15	08.80	10.33	3.67	0.55	374.28	120 /8	101 76	226.05
RD3	179.15	90.00	10.55	3.07	0.00	374.20	120.40	494.70	220.00
RD4	184.19	115.50	11.33	3.17	0.63	379.62	105.63	485.25	266.34
RD5	202.70	129.00	10.67	3.67	0.64	378.67	106.78	548.30	275.95
RD6	197.63	101 90	13.00	2.83	0.52	429.67	118.63	533 71	251 38
	101.00	00.00	10.00	2.00	0.52	440.00	110.00	405.45	201.00
RD7	184.03	96.80	14.33	2.33	0.53	418.69	112.10	485.45	260.59
RD8	180.90	114.60	9.33	3.83	0.63	365.58	114.40	479.98	233.66
RD9	184 24	104 00	14 33	1 83	0.56	424 68	126 55	551 23	256 77
	185.43	112.40	12.33	2.83	0.61	400.88	100.63	523.51	242.65
RDTU	103.43	112.40	12.00	2.00	0.01	400.00	122.00	JZJ.J1	242.0J
RN1	158.31	70.90	9.00	4.00	0.45	3/4.56	103.28	477.84	238.69
RN2	152.53	86.60	10.33	3.67	0.57	361.46	113.56	475.02	240.28
RN3	148 62	97 80	12 67	2 50	0.66	423 46	110 25	530 79	266 41
DNA	170 46	00 70	0.00	1 17	0.50	250.05	00.15	457.40	200.11
r.iN4	170.40	90.70	9.00	4.17	0.00	309.20	90.10	457.40	220.01
RN5	171.26	104.60	11.00	3.00	0.61	405.69	108.60	514.29	234.48
RN6	176.30	100.00	12.67	2.50	0.57	375.48	124.86	500.34	246.26
RN7	185.01	106.20	14.67	2 00	0.57	408.46	106.49	51/ 95	258.03
	100.01	100.20	14.00	2.00	0.07	400.40	100.45	502.00	200.00
KN8	184.38	120.00	11.33	3.00	0.65	402.94	120.15	523.09	260.71
RN9	173.58	106.40	9.33	3.83	0.61	368.49	106.14	474.63	233.01
RN10	177 72	87.00	15.67	1.67	0.49	422 15	101 78	523 93	261 18
	160.76	78 20	11.00	3 00	0.46	375 56	100.10	175 74	220.61
KI I	109.70	10.30	11.55	3.00	0.40	3/5.50	100.18	4/5./4	230.01
RH2	161.28	96.60	9.67	3.83	0.60	388.45	103.13	491.58	247.03
RH3	148.65	98.70	11.33	3.00	0.66	365.50	115.19	480.69	230.87
RHA	160 57	101.00	7 67	4 50	0.63	356 20	102 3/	458.63	232 12
DUE	100.07	101.00	1.01		0.00	000.23	102.04	407.00	202.10
RH5	198.06	115.00	9.33	3.83	0.58	363.77	103.23	467.00	230.60
RH6	177.74	89.80	9.00	4.00	0.51	419.31	116.78	536.09	263.43
RH7	173 80	95 70	7 67	4 67	0.55	360 40	118.30	478 70	227 60
	150.60	05 50	0 00	1 50	0.60	270.00	04.00	166 67	224.06
	109.00	90.00	0.00	4.00	0.00	512.29	94.20	400.37	204.90
RH9	165.04	78.90	10.67	3.50	0.48	369.22	99.62	468.84	234.12
RH10	173.74	100.80	8.00	4.50	0.58	366.74	122.14	488.88	246.40
Range	129 40-202 70	52 30-129 00	7 67-15 67	1 67-4 67	0.32-0.60	318 96-429 67	90 63-126 55	411 48-551 23	202 81-275 05
Nange	470.07	00.00	1.01-10.01	1.07-4.07	0.02-0.03	010.00-420.07	400 70	400 70	202.01-213.33
Mean	170.27	92.26	10.78	3.24	0.54	382.00	106.73	488.73	239.76
CD at 5%	7.48	1.04	3.62	1.47	0.03	8.79	6.94	12.18	24.35

Table-3 Growth performance of different progenies of A. nilotica at the age of 12 months in nursery

Growth Variation Among Progenies of Acacia nilotica Cpts of Different Provenances

Table-4 Magnitude of variation for	percent germination an	d various growth	characters of differen	nt progenies of A	A nilotica
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U		U	1 0	
Growth characters	GCV	PCV	Heritability	GA (% mean)
Germination (%)	26.96	29.39	84.19	50.96
Shoot length (cm)	7.62	9.70	74.52	14.86
Root length (cm)	12.82	17.84	70.44	25.54
No. of branches per plant	15.05	25.62	34.51	18.22
Straightness	17.50	33.01	28.11	19.12
Root-shoot ratio	9.21	15.17	58.71	17.94
Shoot weight(m)	4.84	7.46	61.80	9.46
Root weight(m)	8.18	9.11	80.63	15.13
Fresh seedling weight (m)	4.35	6.52	63.41	8.50
Dry seedling weight (m)	4.73	9.14	26.78	5.04

Table-5 Phenotypic correlations among percent germination and various growth characters of different progenies of A. nilotica

	Germination	Shoot length	Root length	No. of branches	Straightness	Root-shoot ratio	Shoot weight	Root weight	Fresh seedling weight
Shoot length	-0.04						Ĭ		Ĩ
Root length	-0.17*	0.56**							
No. of branches	-0.05	0.22**	0.03						
Straightness	0.06	0.24	0.06	-0.94**					
Root-shoot ratio	-0.18*	0.09	0.82**	-0.10	0.16*				
Shoot weight	0.03	0.38**	0.15*	0.46**	-0.43**	-0.05			
Root weight	-0.02	0.27**	0.43**	0.20**	-0.17*	0.34**	0.20**		
Fresh seedling weight	0.02	0.42**	0.27**	0.47**	-0.44**	0.06	0.95**	0.48**	
Dry seedling weight	0.006	0.21**	0.19	0.29**	-0.27**	0.03	0.49**	0.24**	0.51**

*,** Significant at 5% and 1%, respectively

Table-6 Genotypic correlations among percent germination and various growth characters of different progenies of A. nilotica

	Germination	Shoot	Root	No. of	Straightness	Root-shoot	Shoot	Root	Fresh seedling
		length	length	branches		ratio	weight	weight	weight
Shoot length	-0.04								
Root length	-0.19*	0.56**							
No. of branches	-0.06	0.37**	0.05						
Straightness	0.03	-0.27**	0.12	-0.76**					
Root-shoot ratio	-0.22**	0.07	0.84**	-0.18*	0.31**				
Shoot weight	0.09	0.40**	0.15*	0.81**	-0.84**	-0.05			
Root weight	-0.06	0.31**	0.48**	0.26**	-0.18*	0.38**	0.20**		
Fresh seedling weight	0.05	0.45**	0.27**	0.81**	-0.81**	0.06	0.94**	0.47**	
Dry seedling weight	-0.05	0.45**	0.27**	0.91**	-0.86**	0.05	0.88**	0.53**	0.86**

*,** Significant at 5% and 1%, respectively

Shoot length was largest (140.48 cm) in HS1 from Sonipat provenance, followed by RD5 (137.60 cm) at the age of nine months [Table-2]. In the present study, the progenies of six different provenances of *A. nilotica* at the age of 12 months showed substantial variations in mean performance of growth performance in nursery [Table-3]. The progenies of RD5 from Dausa showed fast growth with higher values for shoot length, root length, root-shoot ratio and straightness. It was closely followed by the progenies of HS1 from Sonipat. Progenies of other selected trees RD1, RD4, RD5, RD6 and RD7 from Dausa provenance, HS1 and HS10 from Sonipat showed desirable growth performance in nursery [Table-3]. Among different provenances under study, the progenies of the selected trees from Dausa provenances showed highest vigorous growth for desirable characters in nursery. It seems that the progenies from Dausa provenance adapted better to the climatic conditions under which they were raised. These findings are in line with the observations of Kedharnath, *et al.* [5], Bagchi, *et al.* [6], Bagchi [7], Bangarwa [8], Sidhu [9], Mishra [10] and Solanki, *et al.* [11].

Variability, Heritability and Genetic advance

The values of phenotypic coefficient of variation [Table-4] ranged from 6.25 (Frersh seedling weight) to 33.01 percent (straightness). Overall, the estimate of genotypic coefficient of variation was less in magnitude than the phenotypic coefficient of variation indicating environmental effect on these characters. Heritability estimates in broad sense were observed higher than 50 percent under study (except dry seedling weight, straightness, no. of branches per seedling). The high estimates of heritability help the breeder in the selection programme. Johanson, *et al.* [4] suggested that heritability estimates in conjunction with genetic advance are usually more helpful in predicting its resultant effect for selecting the best individuals. High heritability (broad sense) may be due to non-

additive gene action (dominance and epistasis), so it will be reliable only if accompanied high genetic gain. The genetic advance (in percent of mean) ranged from 5.04 (dry seedling weight) to 50.96 (germination) [Table-4]. In present study high heritability coupled with moderate to high genetic advance (% mean) for straightness, root length and germination which indicated that high heritability obtained for these characters was probably due to additive gene effects. High heritability accompanied by moderate to high genetic advance for growth parameters have earlier been reported by Solanki, *et al.* [12] in prosopis cineraria; Gera, *et al.* [13] in Dalbergia sissoo and Dhillon, *et al.* [14] in *Azadirachta indica.*

Correlation coefficients

Phenotypic and genotypic correlation coefficients between various seedling characters showed that the magnitude of correlation coefficients at genotypic level was higher than their corresponding correlations at phenotypic level [Table-5] and [Table-6]. This clearly indicated the genotypic association among the characters. Shoot length showed positive correlations with root length, number of branches, shoot weight, root-shoot ratio, fresh seedling weight and dry seedling weight. Such results also have been reported by Thakur and Thakur [15] in *Melia azedarach*. The present observations indicate highly significant variation due to progenies for field emergence, shoot length, root length, fresh seedling weight, numbers of branches per seedling, root-shoot ratio and straightness, open a way to go ahead for further improvement in tree species through selection. Best way for this would be to select enormous amount of seeds from superior trees (through progeny testing) and raise their progenies for second cycle of selection.

Research Category: Forestry, Tree Improvement

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