

Table 2. Mean performance for wheat mutants of some agronomic traits in M₃ generation of pedigree method for Gemmeiza-11 under water stress.

Mutants	Treatments	days to 50% heading	Spike length (cm).	No. of spikelits /spike	No. of spikes/ plant	No. of grains/ pike	Spike grain weight (g).	1000-grain weight (g)	Grain yield/plant (g)	Biological yield / plant (g.)	Harvest index %
	Control	93	14.93	24.4	4.13	72.83	2.65	37.13	7.81	31.72	24.95
Mutant-4	Gem-0.06 SA	94.33	17.53	24.26	8.93	54.13	3.41	63.63	12.55	52.68	23.79
Mutant-5	Gem-0.08 SA	98.33	17.16	24.93	10.7	72.56	3.38	48.76	27.71	93.97	29.52
Mutant-6	Gem-350Gy1	97	16.8	24	8.13	41.73	1.40	33.66	8.7	49.31	17.63
Mutant-9	Gem-350Gy2	100	8.7	20.66	8.4	58.2	3.29	56.83	17.41	49.7	35.06
	F. test	**	**	**	**	**	**	**	**	**	**
	L.S.D_{0.05}	1.97	1.52	1.42	1.38	12.24	0.55	3.54	1.91	6.44	4.41

*,**Significant at 0.05 and 0.01, respectively

Table 3. Mean performance for wheat mutants of some agronomic traits in M₃ generation of pedigree method for shandawel-1, under water stress.

Mutants	Treatments	days to 50% heading	Spike length (cm).	No. of spikelits/ Spike	No. of spikes/ plant	No. of grains/ pike	Spike grain weight (g).	1000-grain weight (g)	Grain yield/plant (g)	Straw yield/plant (g)	Biological yield / plant (g)	Harvest index %
	Control	102.33	15.8	24.4	6.13	79.46	2.84	36.36	13.51	37.2	50.71	26.66
Mutant-2	Sh-0.04SA	94.33	13.2	23.73	6.33	88.8	3.62	40.98	14.72	28.9	43.62	33.71
Mutant-5	Sh-0.06SA	90.66	15.5	24.93	8.86	72.52	2.50	34.51	14.63	46.25	60.88	23.02
Mutant-7	Sh-250 Gy	98.33	16.6	23.86	8.93	65.26	3.003	45.94	18.04	51.01	69.05	26.11
Mutant-8	Sh-350Gy	104.33	16.46	26.53	6.4	60.4	2.56	42.28	11.75	35.57	47.32	24.79
	F. test	**	*	Ns	Ns	*	Ns	**	*	**	**	*
	L.S.D_{0.05}	2.98	1.67	2.17	3.91	11.65	0.66	3.85	2.81	4.81	5.78	4.76

*, **Significant at 0.05 and 0.01 of levels probability, respectively

Table 4. Mean performance for wheat mutants of some agronomic traits in M₃ generation of pedigree method for Sids-12 under water stress.

Mutants	Treatments	days to 50% heading	Spike length (cm).	No. of spikelits/spike	No. of spikes/plant	No. of grains/pike	Spike grain weight (g).	1000-grain weight (g)	Grain yield/plant (g)	Biological yield / plant (g)	Harvest index %
	Control	93.33	12.4	21.6	5	75.66	2.98	39.54	11.69	30.02	38.99
Mutant – 2	Sid-0.04SA	95.66	14.1	23.06	5.66	107.3	4.26	39.55	20.62	48.32	42.68
Mutant-3	Sid-0.06 SA	101	13.73	24.4	7.8	82	4.43	54.37	28.15	79.5	35.44
Mutant-4	Sid-0.08 SA	92.33	15.16	22.93	6.6	120.46	6.20	51.59	34.08	75.04	45.39
Mutant-5	Sid-250 Gy	91.66	16.2	23.46	9.6	91.53	3.83	41.90	21.21	75.94	27.49
Mutant-6	Sid-350 Gy1	93.33	14.73	22.13	5.26	128.2	5.93	45.95	25.64	64.32	39.87
Mutant-9	Sid-350Gy2	94.33	8.66	21.2	4.73	75.06	3.6	48.24	12.86	35.34	36.54
Mutant-10	Sid-350Gy3	92.66	10.13	21.6	4.86	102.73	4.91	47.75	18.17	44.55	40.74
	F. test	**	**	*	*	**	**	**	**	**	**
	L.S.D_{0.05}	1.4862	1.3254	1.4938	2.394	15.781	1.234	3.7818	2.3896	4.6433	3.529

*,**Significant at 0.05 and 0.01of levels probability, respectively

Table 5. Mean performance for wheat mutants of some agronomic traits in M₃ generation of pedigree method for Sahel -1, under water stress.

Mutants	Treatments	days to 50% heading	Spike length (cm).	No. of spikelits/spike	No. of spikes/plant	No. of grains/pike	Spike grain weight (g).	1000-grain weight (g)	Grain yield/plant (g)	Biological yield / plant (g)	Harvest index %
	Control	96	11	22.66	4.46	54.46	2.10	38.56	7.76	23.06	33.65
Mutant-1	Sah-0.04SA	96.66	14.4	24	6	80.2	3.81	47.50	20.52	53.43	38.40
Mutant-2	Sah-0.06SA	95.33	15	24.66	5.46	117.91	5.15	43.67	16.41	54.36	30.18
Mutant-3	Sah-0.08 SA1	92.66	13.33	24.26	4.66	83.41	4.75	56.94	17.44	43.52	40.07
Mutant-4	Sah-0.08 SA2	94.33	12.86	22.26	4.76	89.83	4.18	46.53	13.58	33.78	40.20
Mutant-5	Sah-250 Gy1	90.33	14.13	23.2	3.53	96.55	5.40	55.92	16.44	42.94	38.28
Mutant-6	Sah-250 Gy2	92.33	9.33	22	9.7	61.6	2.90	47.07	19.33	53.16	36.36
Mutant-7	Sah-350 Gy1	94.33	11.53	24.4	5.8	54	2.32	42.96	8.17	30.67	26.63
Mutant-9	Sah-350 Gy2	89.33	14.53	23.6	4.8	104.2	5.45	52.30	21.43	55.72	38.46
Mutant-12	Sah-350Gy3	92.33	12.93	22.13	7.8	61.83	2.98	48.19	18.52	57.99	31.93
Mutant-14	Sah-350Gy4	93.33	11	22.93	4.26	57.6	3	52.08	10.28	28.76	35.74
	F. test	**	**	**	**	**	**	**	**	**	**
	L.S.D _{0.05}	1.72	1.03	1.15	1.50	8.00	0.55	6.39	3.46	8.40	5.79

*,** Significant at 0.05 and 0.01 of levels probability, respectively

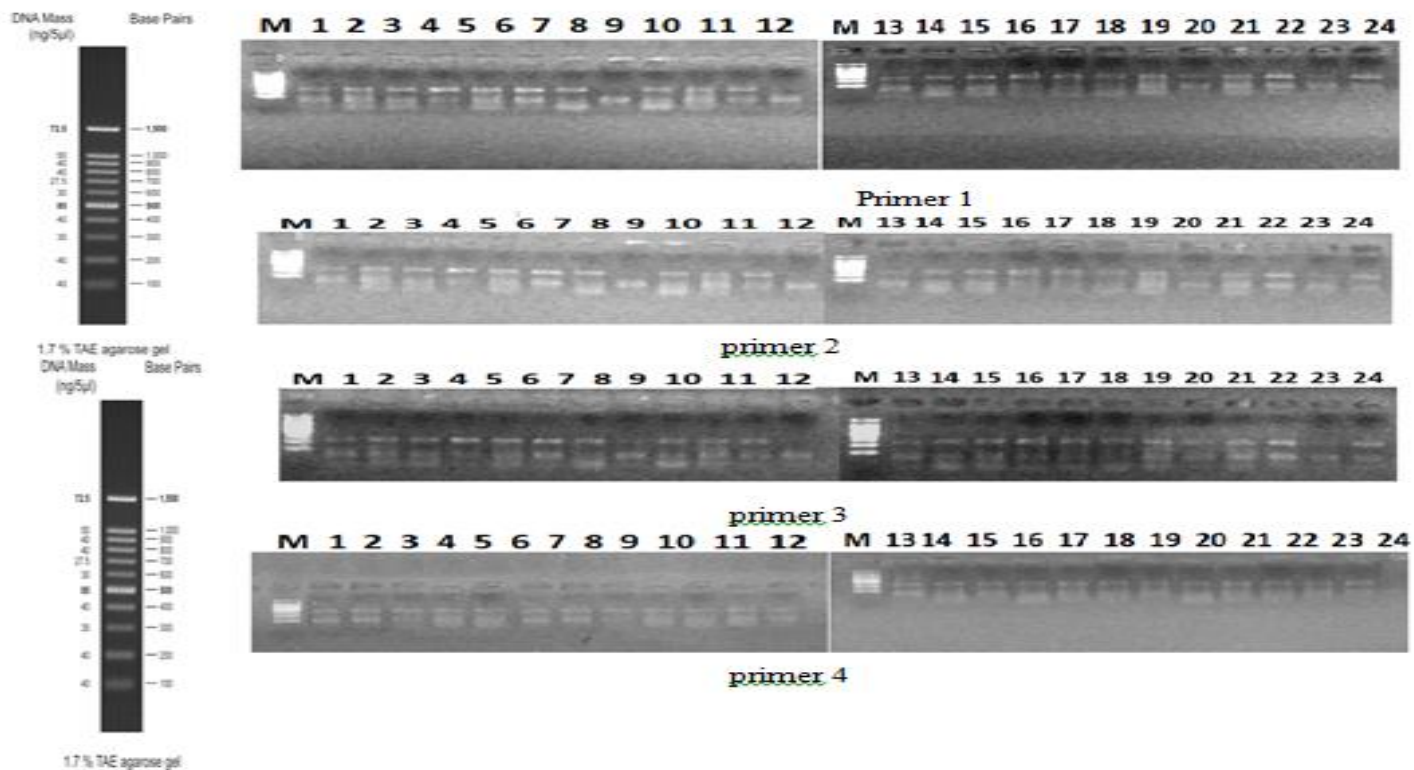


Figure 1. RAPD pattern obtained by five primers for DNA marker

1:Sahel-1 control, 2:Sahel-1 250Gy, 3:Sahel-1 350Gy1, 4: Sahel-1 350Gy2 5: Sahel-1 0.04 % SA, 6:Sahel-1 0.06 % SA, 7:Sahel-1 0.08 % SA, 8: Sids-12 control, 9:Sids-12 0.04 %SA, 10:Sids-12 0.06 %SA, 11:Sids-12 0.08 % SA, 12:Sids-12 250Gy, 13: Sids-12 350Gy1, 14:Sids-12 350 Gy2, 15:Shandaweel-1 control, 16:Shandaweel-1 0.06 % SA, 17:Shandaweel-1 0.04 %SA, 18:Shandaweel-1 250Gy, 19:Shandaweel-1 350Gy, 20: Gemmieza-11control, 21: Gemmieza-11 0.06% SA, 22: Gemmieza-110.08 %SA, 23:Gemmieza-11 350 Gy1, 24:Gemmiza-11 350Gy2.

Table (8): The similarity values between the wheat genotypes computed according to RAPD data.

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23
1	1																						
2	0.3	1																					
3	0.3	0.3	1																				
4	0.4	0.7	0.5	1																			
5	0.3	0.2	0.3	0.6	1																		
6	0.3	0.3	0.5	0.4	0.1	1																	
7	0.4	0.5	0.3	0.4	0.3	0.4	1																
8	0.3	0.6	0.6	0.7	0.5	0.3	0.9	1															
9	0.3	0.3	0.4	0.5	0.1	0.2	0.2	0.6	1														
10	0.3	0.1	0.4	0.7	0.1	0.2	0.3	0.6	0.1	1													
11	0.2	0.4	0.6	0.3	0.2	0.1	0.5	0.3	0.3	0.3	1												
12	0.5	0.8	0.8	0.7	0.5	0.3	0.9	0.1	0.6	0.6	0.3	1											
13	0.2	0.3	0.4	0.5	0.2	0.1	0.5	0.3	0.3	0.3	0.1	0.4	1										
14	0.3	0.3	0.2	0.4	0.1	0.3	0.1	0.7	0.2	0.2	0.3	0.7	0.3	1									
15	0.3	0.3	0.2	0.6	0.1	0.3	0.3	0.5	0.2	0.2	0.3	0.5	0.3	0.1	1								
16	0.3	0.5	0.5	0.3	0.3	0.3	0.3	0.7	0.2	0.3	0.3	0.7	0.3	0.3	0.3	1							
17	0.3	0.6	0.4	0.3	0.3	0.3	0.3	0.6	0.3	0.4	0.4	0.6	0.4	0.3	0.2	0.1	1						
18	0.6	0.7	0.5	0.4	0.6	0.6	0.4	0.9	0.5	0.7	0.7	0.9	0.7	0.6	0.4	0.3	0.2	1					
19	0.3	0.3	0.3	0.7	0.2	0.3	0.3	0.6	0.3	0.1	0.4	0.6	0.4	0.2	0.1	0.3	0.3	0.5	1				
20	0.3	0.5	0.7	0.4	0.3	0.1	0.6	0.2	0.3	0.3	0.1	0.2	0.2	0.4	0.4	0.4	0.5	0.8	0.5	1			
21	0.4	0.2	0.3	0.8	0.1	0.3	0.4	0.5	0.2	0.1	0.3	0.5	0.3	0.3	0.1	0.4	0.3	0.6	0.1	0.4	1		
22	0.3	0.4	0.4	0.5	0.2	0.1	0.5	0.3	0.3	0.3	0.1	0.3	0.1	0.3	0.2	0.3	0.3	0.5	0.3	0.2	0.2	1	
23	0.4	0.7	0.7	0.6	0.4	0.3	0.8	0.2	0.5	0.5	0.2	0.1	0.3	0.6	0.4	0.6	0.5	0.8	0.5	0.1	0.4	0.2	1
24	0.3	0.5	0.5	0.4	0.3	0.1	0.6	0.2	0.3	0.3	0.1	0.2	0.2	0.4	0.3	0.4	0.3	0.6	0.3	0.1	0.3	0.1	0.1

1:sahel-1control, 2:Sahel-1250Gy, 3:Sahel-1350Gy1, 4: Sahel-1 350Gy2 5: Sahel-1 0.04 % SA, 6:Sahel-1 0.06 % SA, 7:Sahel-10.08 % SA, 8: Sids-12 control, 9:Sids- 12 0.04 %SA, 10:Sids-12 0.06 %SA, 11:Sids- 12 0.08 % SA, 12:Sids-12 250Gy, 13: Sids-12 350Gy1, 14:Sids-12 350 Gy2, 15:Shandaweel-1 control, 16:Shandaweel-1 0.06 % SA, 17:Shandaweel-1 0.04 %SA, 18:Shandaweel-1 250Gy, 19:Shandaweel-1 350Gy, 20: Gemmieza-11control, 21: Gemmieza-11 0.06% SA 22: Gemmieza-110.08 %SA, 23:Gemmieza-11 350Gy1, 24:Gemmiza-11 350Gy2.

