

Table 1 Disease intensity of different lines, testers and their crosses at two locations. DMRT analysis is needed (treatment means can be compared by using C.D. and S.E. already given in the table hence DMRT is not necessary)

Genotype	Disease intensity (E1)	Disease intensity (E2)
K-08-59 x SKAU-7A	11.778	13.562
K-08-59 x SKAU-11A	9.084	11.064
K-08-59 x IR-68888A	10.778	13.432
Jhelum x SKAU-7A	12.113	15.189
Jhelum x SKAU-11A	11.282	14.122
Jhelum x IR-68888A	11.091	16.765
SKAU-382 x SKAU-7A	3.172	4.760
SKAU-382 x SKAU-11A	2.705	2.123
SKAU-382 x IR-68888A	2.684	2.269
SK-389 x SKAU-7A	8.602	10.876
SK-389 x SKAU-11A	5.931	6.330
SK-389 x IR-68888A	5.491	6.760
Ch-988 x SKAU-7A	13.922	15.187
Ch-988 x SKAU-11A	11.943	16.342
Ch-988 x IR-68888A	5.318	6.975
SR-1 x SKAU-7A	1.535	1.863
SR-1 x SKAU-11A	1.624	1.231
SR-1 x IR-68888A	2.250	2.750
K-08-60 x SKAU-7A	5.350	5.120
K-08-60 x SKAU-11A	4.881	3.768
K-08-60 x IR-68888A	0.713	0.381
K-08-61 x SKAU-7A	1.724	1.532
K-08-61x SKAU-11A	0.238	0.901
K-08-61 x IR-68888A	0.163	0.768
PS-5x SKAU-7A	3.703	3.907
PS-5x SKAU-11A	0.329	0.726
PS-5x IR-68888A	1.564	1.667
K-08-59	15.811	15.811
Jhelum	16.908	16.908
SKAU-382	3.684	4.684
SKAU-389	5.376	7.675
China-988	10.104	13.104
SR-1	1.181	1.235
K-08-60	2.183	2.790
K-08-61	1.251	1.899
PS-5	0.704	0.974
SKAU-7A	3.461	3.764
SKAU-11A	4.653	4.383
IR-68888A	1.377	1.856
Check	14.251	16.908
CD	0.632	0.722
SE (d)	0.216	0.362
SE (m)	0.142	0.256
CV	3.221	3.821