

**Table S2** Genetic diversity indices for the different SSRs over all populations

SSR	Seedlings				Adults			
	$N_a$	$A_R$	$H_o$	$H_e$	$N_a$	$A_R$	$H_o$	$H_e$
sfc0018	10.2	9.5	0.689	0.700	8.7	7.6	0.681	0.698
sfc0161	13.2	12	0.785	0.794	12.5	9.6	0.783	0.786
sfc1063	9.5	9	0.778	0.792	8.7	7.9	0.779	0.781
sfc1143	10.1	9.5	0.754	0.764	9.5	8.1	0.764	0.763
GOT066	3.2	2.9	0.131	0.125	3.1	2.5	0.135	0.128
FIR065	4.1	4.1	0.685	0.671	4	4	0.708	0.673
FIR004	7.1	6.7	0.480	0.534	6.2	5.1	0.446	0.502
FS3-04	3.7	3.5	0.422	0.421	3.3	2.8	0.402	0.408
mfs11	8.3	7.7	0.663	0.668	7.8	6.5	0.681	0.663

$N_a$  - number of alleles,  $A_R$  - allelic richness,  $H_o$  - observed heterozygosity,  $H_e$  - expected heterozygosity