

Table S2. Population genetic summary data for the loci used in this study. The top nine are candidate loci and the rest of the list are control loci named according to [1]

Loci	Length	n	S	θ_π	θ_w	Tajima's D
PoGI	2668	92	32	0.00216	0.00220	-0.015
PoCCA1	3534	83	81	0.00383	0.00392	-0.31
PoPhyN	3167	72	20	0.00089	0.00102	-0.11
PoPhyO	2187	76	6	0.00026	0.00051	-0.61
PoPhyP	1459	85	9	0.00113	0.00108	0.18
PoPRR3	1448	93	28	0.00079	0.00272	-1.79
PoPRR7	2634	86	56	0.00146	0.00209	-0.98
PoFTL2	1373	83	35	0.00156	0.00285	-0.91
PoMFTL1	2081	91	99	0.00736	0.0066	0.21
Sequence 8	652	77	32	0.00984	0.00999	-0.05
Sequence 12	539	95	10	0.00132	0.00362	-1.63
Sequence 14	429	96	3	0.00073	0.00136	-0.84
Sequence 18	821	90	2	0.00026	0.00048	-0.73
Sequence 22	683	95	11	0.00157	0.00314	-1.32
Sequence 24	723	94	11	0.00245	0.00297	-0.46
Sequence 28	1141	82	19	0.00111	0.00335	-1.97
Sequence 31	649	95	10	0.00334	0.00301	0.28
Sequence 32	726	95	9	0.00092	0.00242	-1.56
Sequence 33	803	92	20	0.00402	0.00489	-0.52
Sequence 37	629	96	16	0.00272	0.00495	-1.27
Sequence 49	535	93	9	0.00193	0.00330	-1.05
Sequence 56	572	93	21	0.00832	0.00719	0.46
Sequence 58	600	95	23	0.00964	0.00748	0.86
Sequence 59	740	95	9	0.00316	0.00237	0.84
Sequence 60	434	96	2	0.00032	0.00090	-1.00
Sequence 62	744	96	23	0.00689	0.00602	0.43
Sequence 68	601	92	16	0.00574	0.00523	0.28
Sequence 69	708	95	6	0.00098	0.00165	-0.92
Sequence 70	929	95	28	0.00446	0.00588	-0.73
Sequence 77	507	95	17	0.00478	0.00654	-0.77
Sequence 87	633	90	6	0.00127	0.00187	-0.73
Sequence 88	599	95	26	0.00743	0.00847	-0.37
Sequence 89	642	95	8	0.00225	0.00243	-0.18
Sequence 92	740	91	16	0.00317	0.00425	-0.72
Sequence 99	557	82	5	0.00179	0.00180	-0.02
Sequence 100	459	90	0	0	0	<i>n.a.</i>