Table S4. Sequences of two significant SNP markers identified in the HA 300 × RHA 464 mapping population.

SNP marker	Allele	Map position (cM)	SNP sequence
Ha5_11356218	C/T[a]	14.6	AGTGGCTTTCACATATGGTCATAAAGTTCTAAT ACTTCTTGGGAGATCTTTTCTGTTACTAATC[C/T] TTTCATCTGCAACTGACTATTTGGACAGTGAATT CTATTCGGTTACTCAGCTGATAATAATGTTTATA CCATAGTAGTTGAAC
Ha6_8364901	T/C[a]	60.5	TTTAAATGTCCACTTGCTTTGAACAACACTCCAT GGCCTGCGGTACC[T/C]TTTAGATACCTTAGAAT TCTCTGAGCAGCTTCCATGTGGGCAACTTGCGG TTGGTGCATAAACTGACTTACCACTCC

[a] The first allele is from HA 300 and the second allele is from RHA 464.