

## Accessory Table

### Means and standard variation for genotypic combinations at minor QTL locations on chromosomes 1H and 3H for germination percentage in the three environments tested

$R^2$  values are shown for  $***P < 0.0001$ . Allele 'A' is the Stirling allele and allele 'B' is the Harrington allele. The first allele of the genotypes is representative of the allele at the 1H QTL and the second allele is representative of the allele at the 3H QTL, e.g. AB = Stirling allele at the 1H QTL and Harrington allele at the 3H QTL

Genotype	2007MB		2007WH		2006WH	
	Mean	s.d.	Mean	s.d.	Mean	s.d.
AA	72.5	4.2	82.7	3.2	55.7	5.5
AB	86.6	5.4	92.9	4.2	72.6	7.1
BA	90.6	5.0	96.1	3.9	73.9	6.7
BB	26.6	4.3	40.6	3.3	11.5	5.7
$R^2$	0.41***		0.47***		0.28***	