

- 1 **Supplemental Table S2.** Significant QTLs from interval mapping of binary classification of
- 2 flowering time phenotypes in the ABR6 x Bd21 F<sub>4:5</sub> families.

ENV <sup>a</sup>	Chr <sup>b</sup>	cM	EWT <sup>c</sup>	LOD
1	Bd1	298.0	3.84	13.1
5	Bd3	302.0	3.64	5.08

- 3 <sup>a</sup>Environment (see Supplemental Table S1)
- 4 <sup>b</sup>Chromosome
- 5 <sup>c</sup>Experiment-wide permutation threshold