Figure S1  Comparison of BAC sequences from *Boechera drummondii* (*Bdr*, 93 kb), *Capsella rubella* (*Cru*, 75 kb), *Arabidopsis lyrata* (*Aly*, 110 kb), and *A. halleri* (*Aha*, 92 kb) to a syntenic region in *A. thaliana* (*Ath*, 60 kb), from which *A. thaliana* FRIGIDA (*FRI*) originated. Black numbers indicate At5gXXXXX gene identifiers. The *FRI* homolog, which is truncated in *A. thaliana*, is indicated in bold (see also IRWIN, J. A., C. LISTER, E. SOUMPOUROU, Y. ZHANG, E. C. HOWELL et al., 2012 Functional alleles of the flowering time regulator *FRIGIDA* in the *Brassica oleracea* genome. BMC Plant Biol. 12: 21). Grey numbers indicate distances in *A. thaliana* in kb. Dashes indicate non-syntenic sequences.