FIGURE S4.—Northern blot analyses of *Wis2-1A* (left) and *Veju* (right) retrotransposon LTR regions in the tetraploid parent (genome BBAA), the diploid parent (genome DD), their hybrid and derived allopolyploid. Methylene blue staining of the membranes shows equal loading of RNA samples. Probes were derived from a consensus sequence from wheat *Wis2-1A* LTRs (a 354bp fragment) and from *Veju1_TM_LTR* entry at TREP (a 339bp fragment).