**TE: Dmel−B−G1975−Map15_s_2−UNCLASSIFIED**
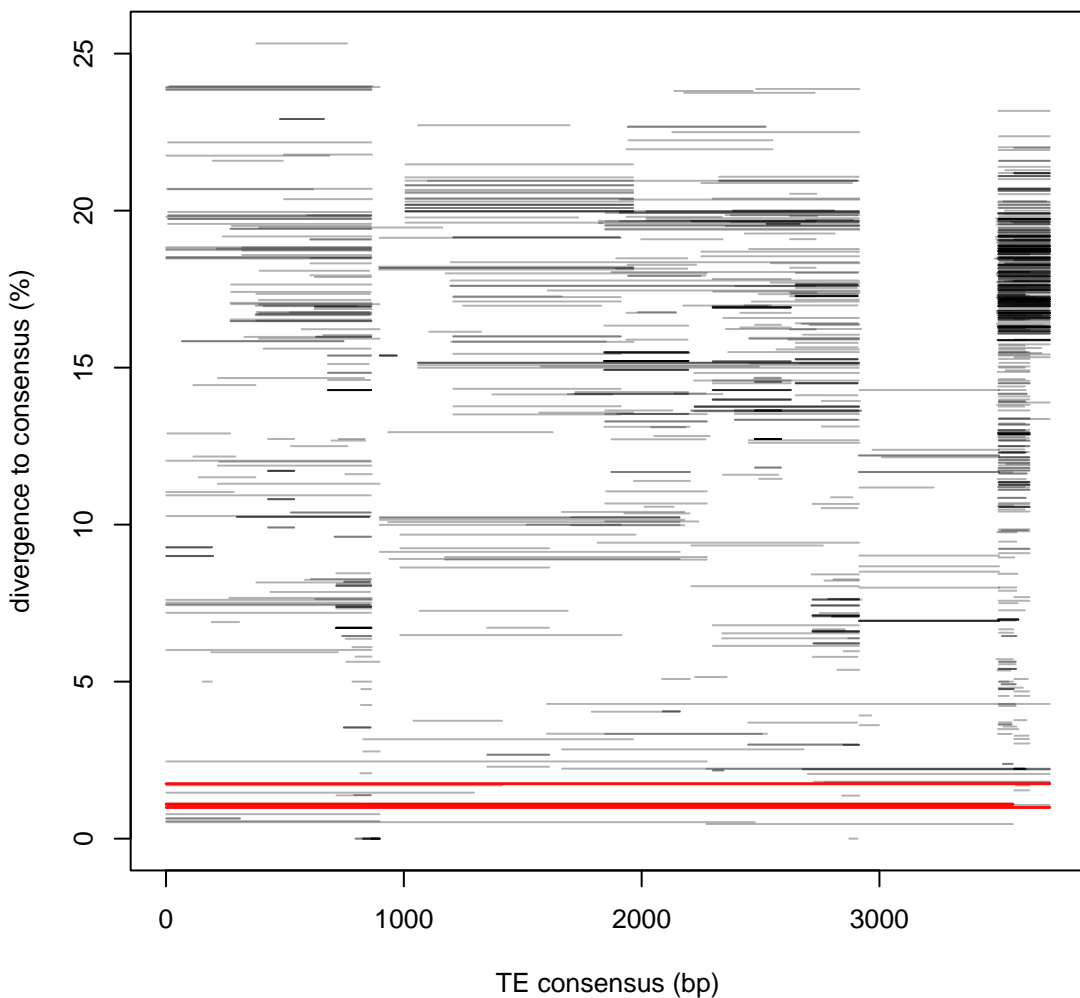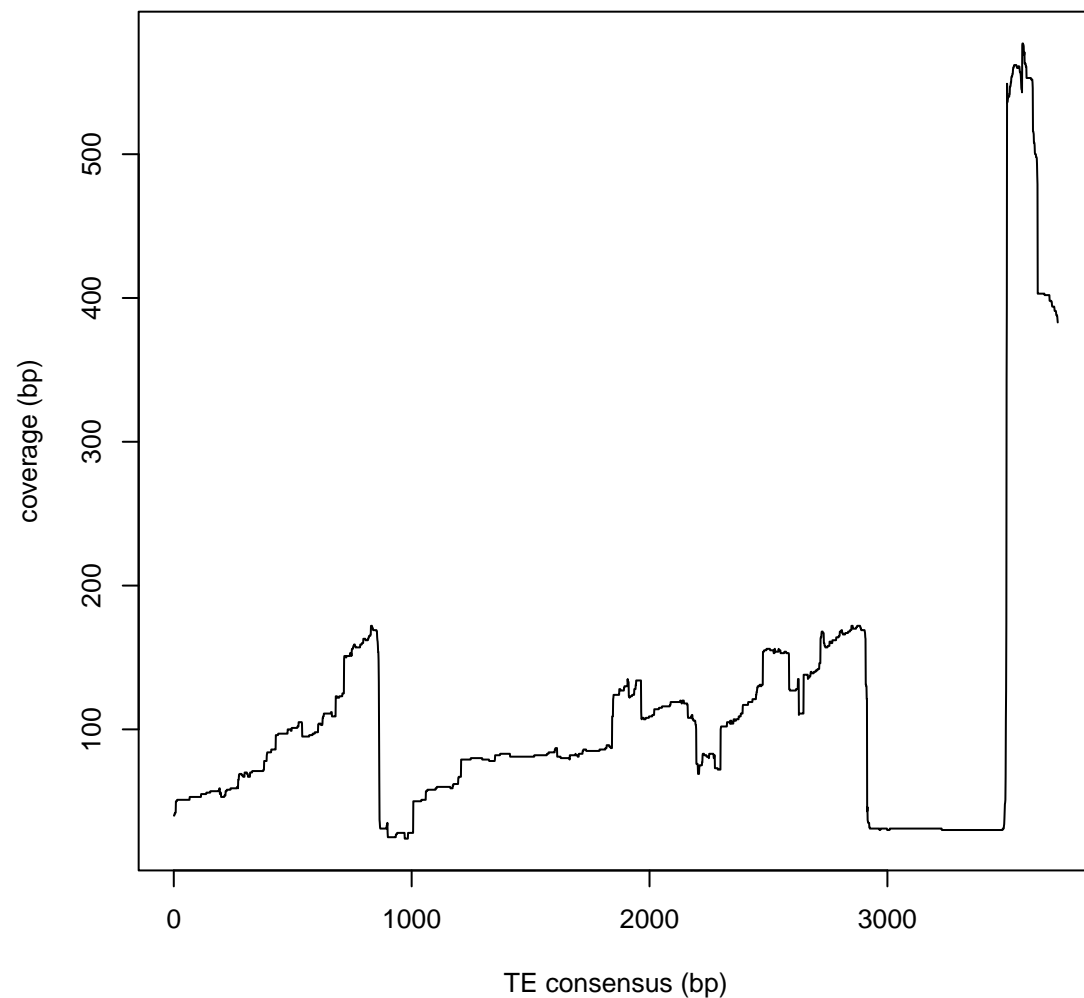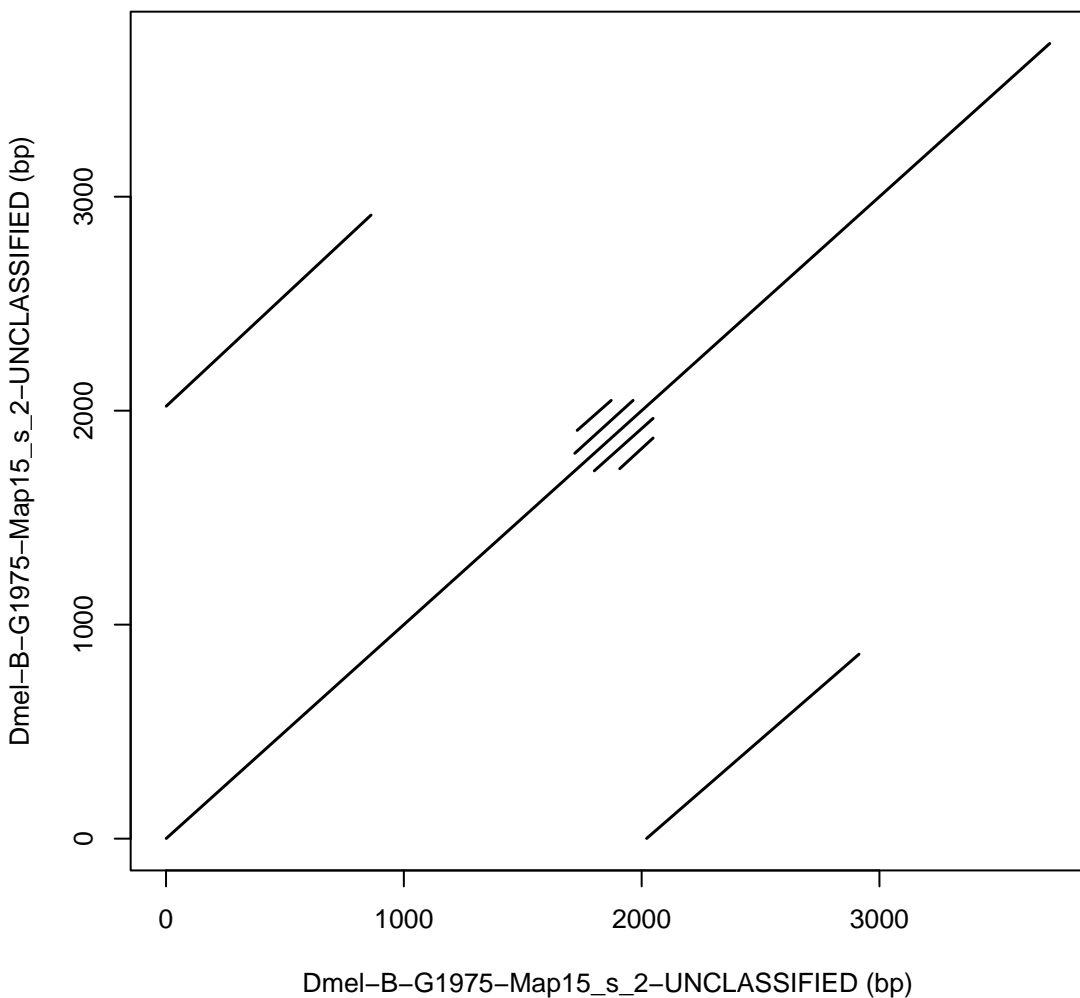**consensus size: 3716bp; fragments: 1215; full length: 4 (>=3344.4bp)**

**TE consensus genomic coverage**

**TE consensus self dotplot (blastn)**

**TE consensus structure and protein hits**

Jockey−14_DAn_pol LINE/I−J