**TE: Dmel−B−R1144−Map6−LTR/GYPSY**
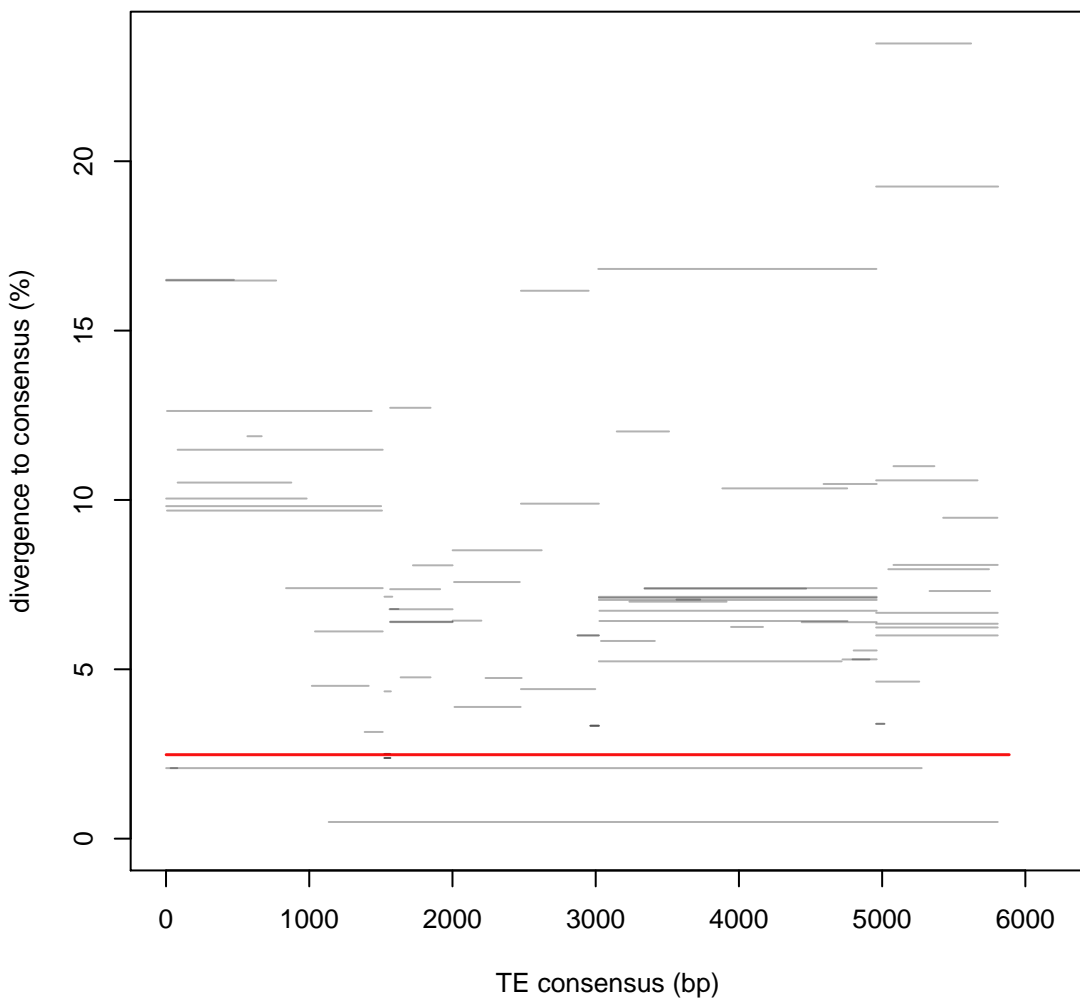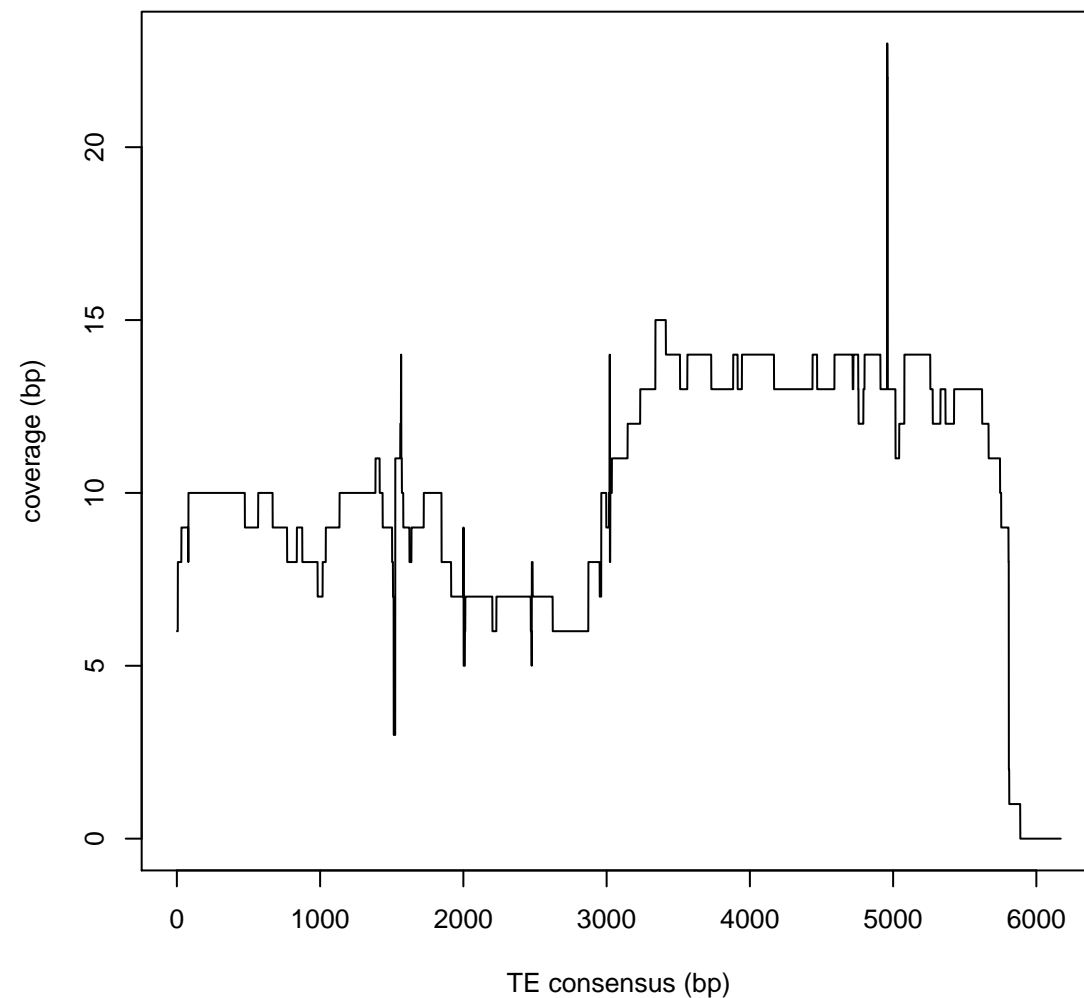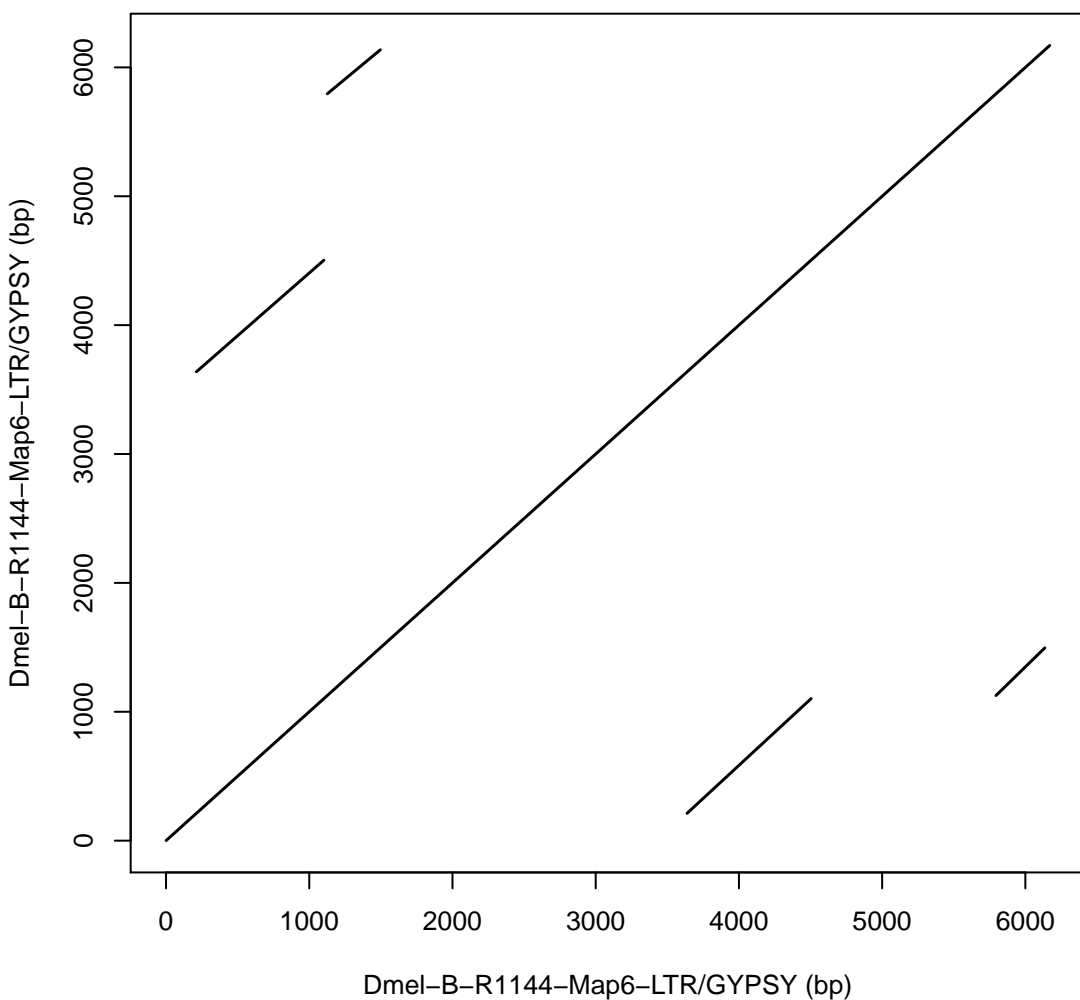consensus size: 6170bp; fragments: 81; full length: 1 (>=5553bp)

**TE consensus genomic coverage**

**TE consensus self dotplot (blastn)**

**TE consensus structure and protein hits**

divergence to consensus (%)

TE consensus (bp)

coverage (bp)

TE consensus (bp)

Dmel−B−R1144−Map6−LTR/GYPSY (bp)

Dmel−B−R1144−Map6−LTR/GYPSY (bp)

Dmel−B−R1144−Map6−LTR/GYPSY (bp)

AM_pol LTR/Gypsy
GYPSM1_pol LTR/Gypsy
UN−TIRANT_pol LTR/Gypsy
Gypsy5_pol LTR/Gypsy
ACCORD_pol LTR/Gypsy
YOYO_pol LTR/Gypsy