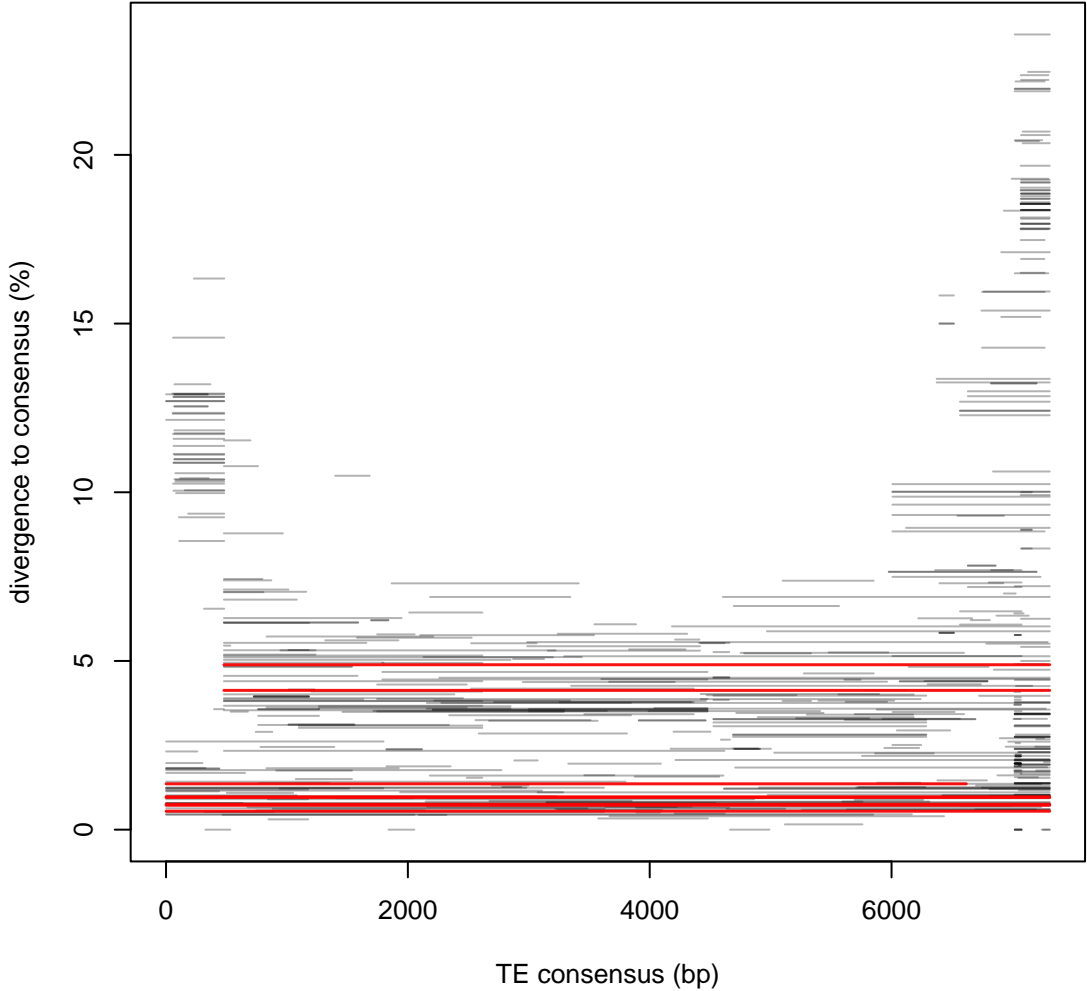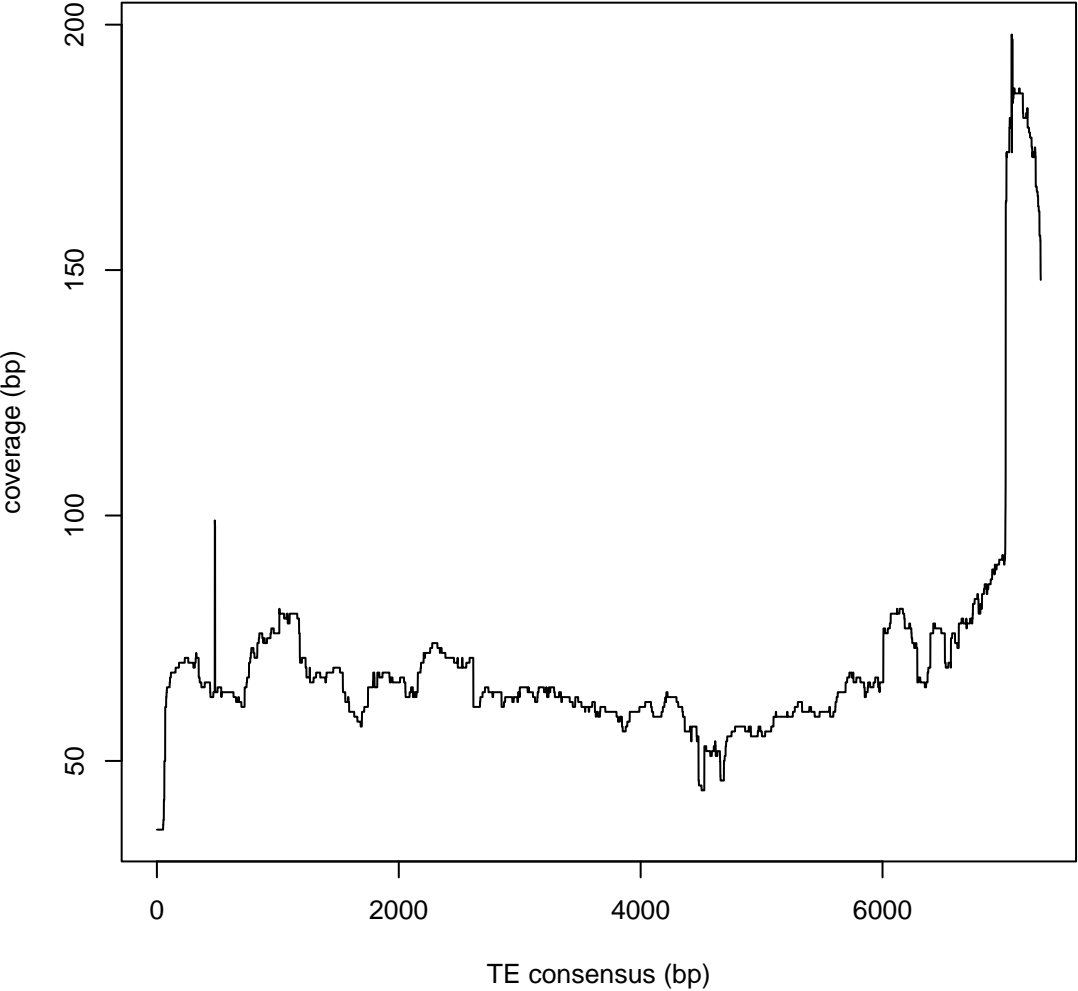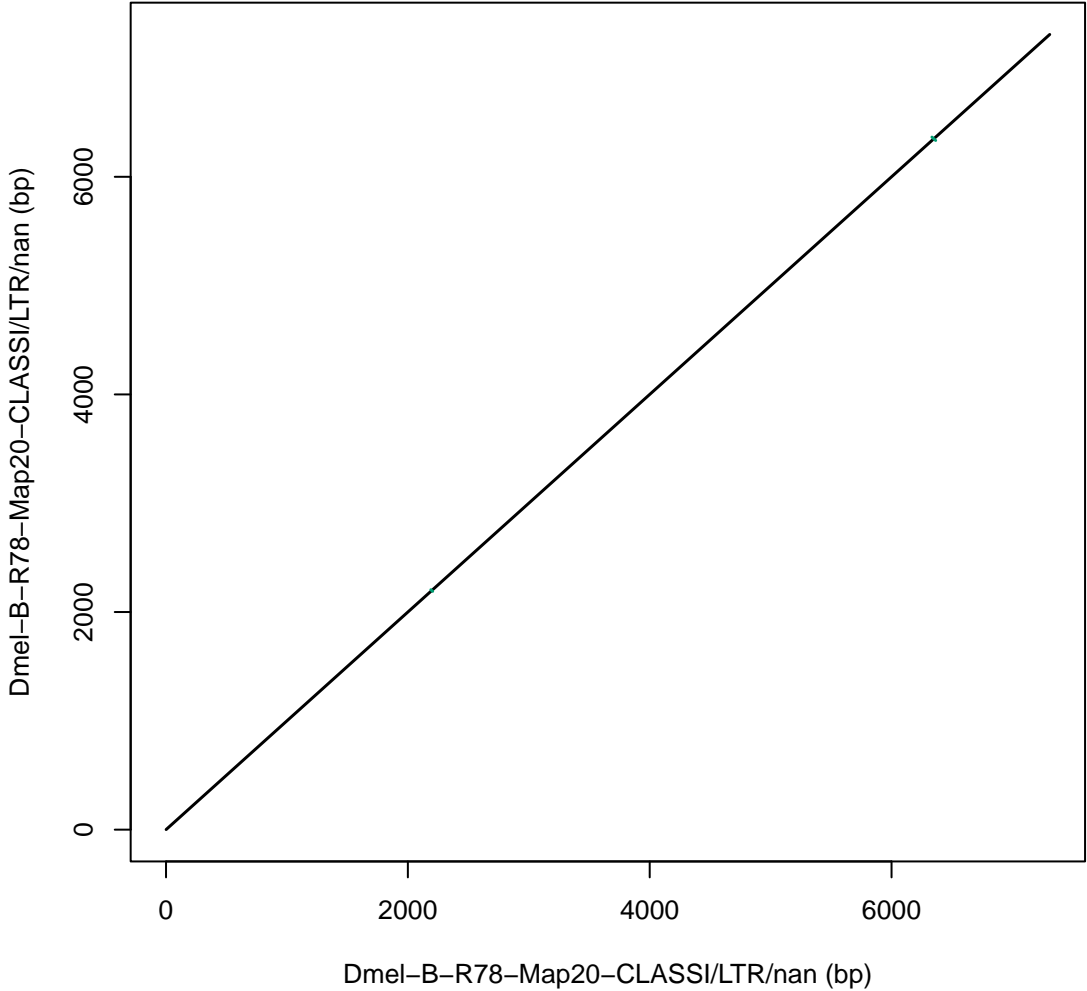**TE: Dmel−B−R78−Map20−CLASSI/LTR/nan**
consensus size: 7308bp; fragments: 524; full length: 8 (>=6577.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Gypsy−40_PT LTR/Gypsy

BEL−5_DBp_pol LTR/Pao

I_Ele17_gag LINE/I

BEL−1_PBa_pol LTR/Pa