**TE: Dmel−B−R1687−Map4−LTR**
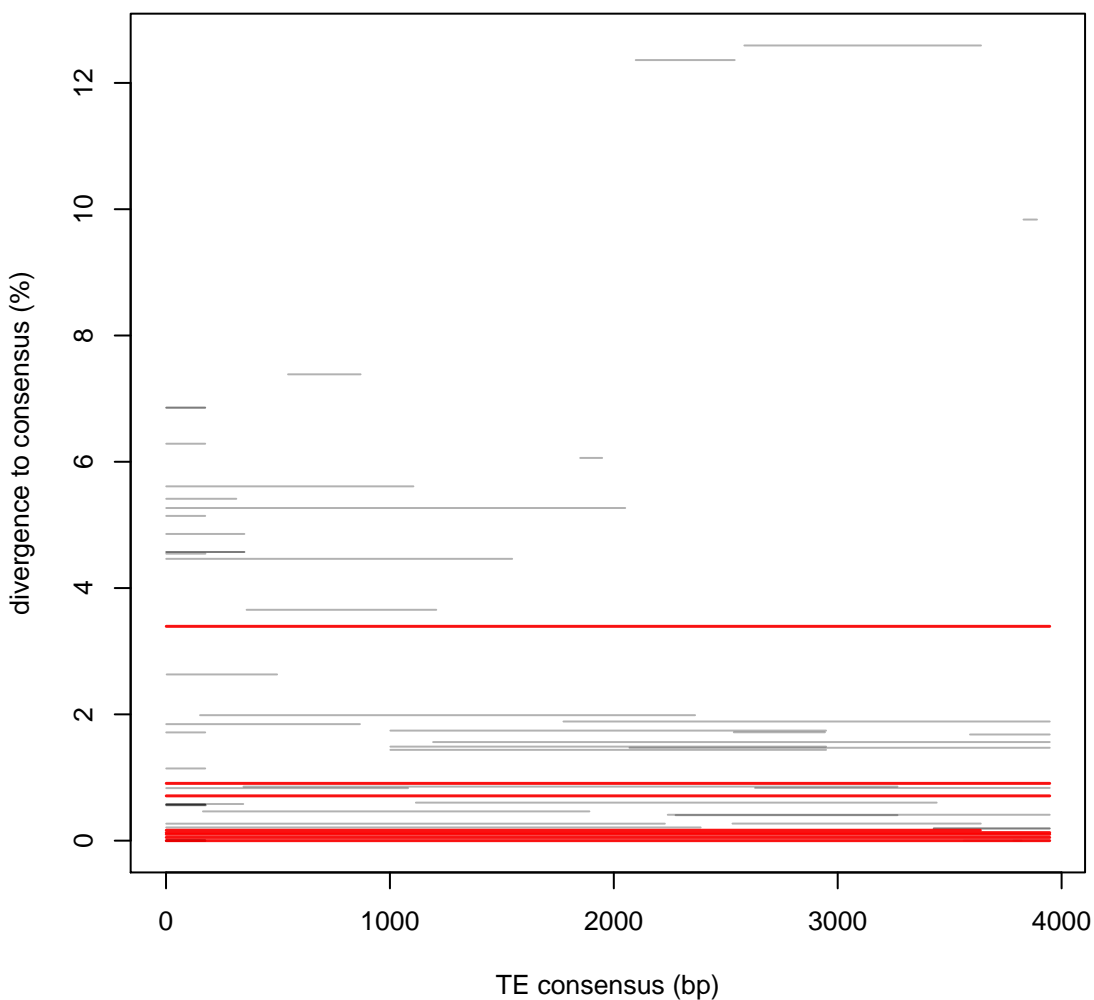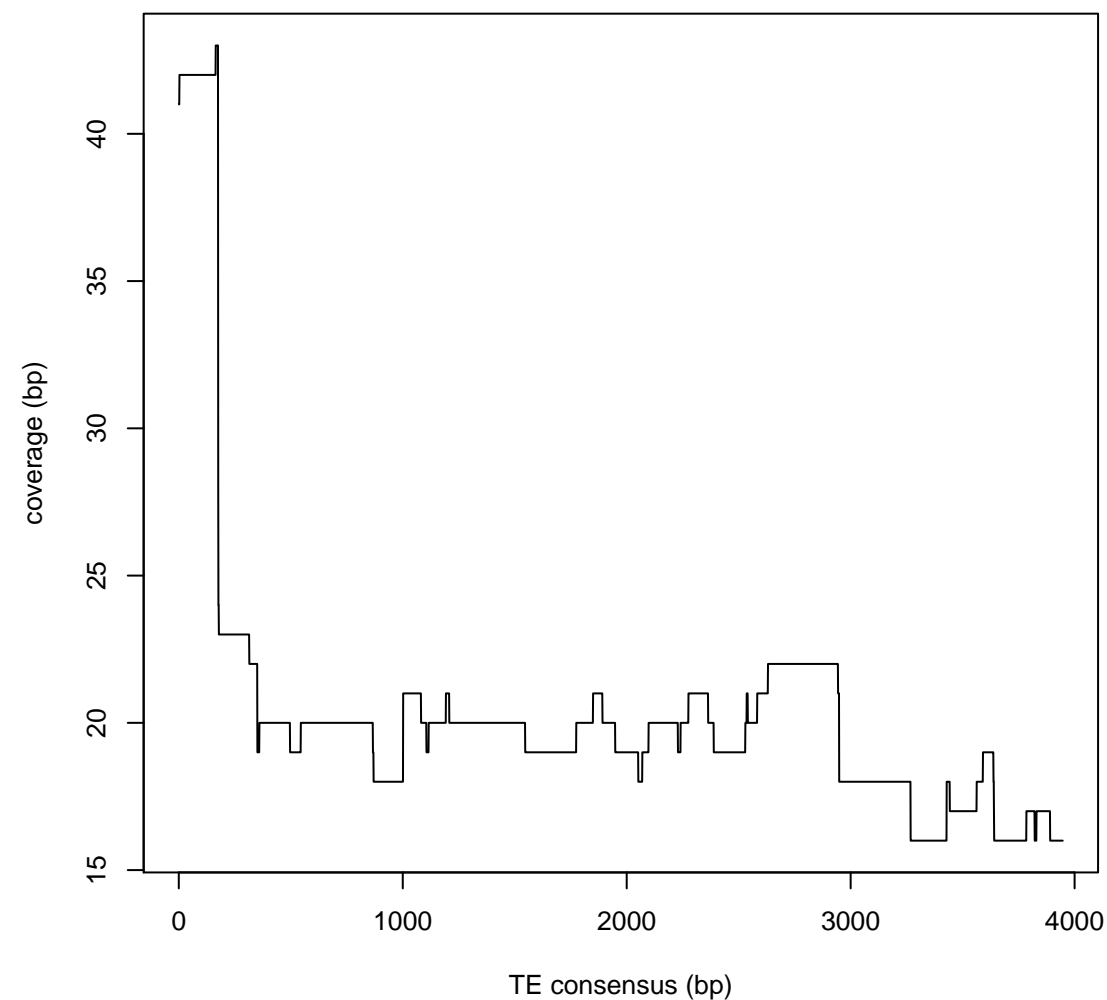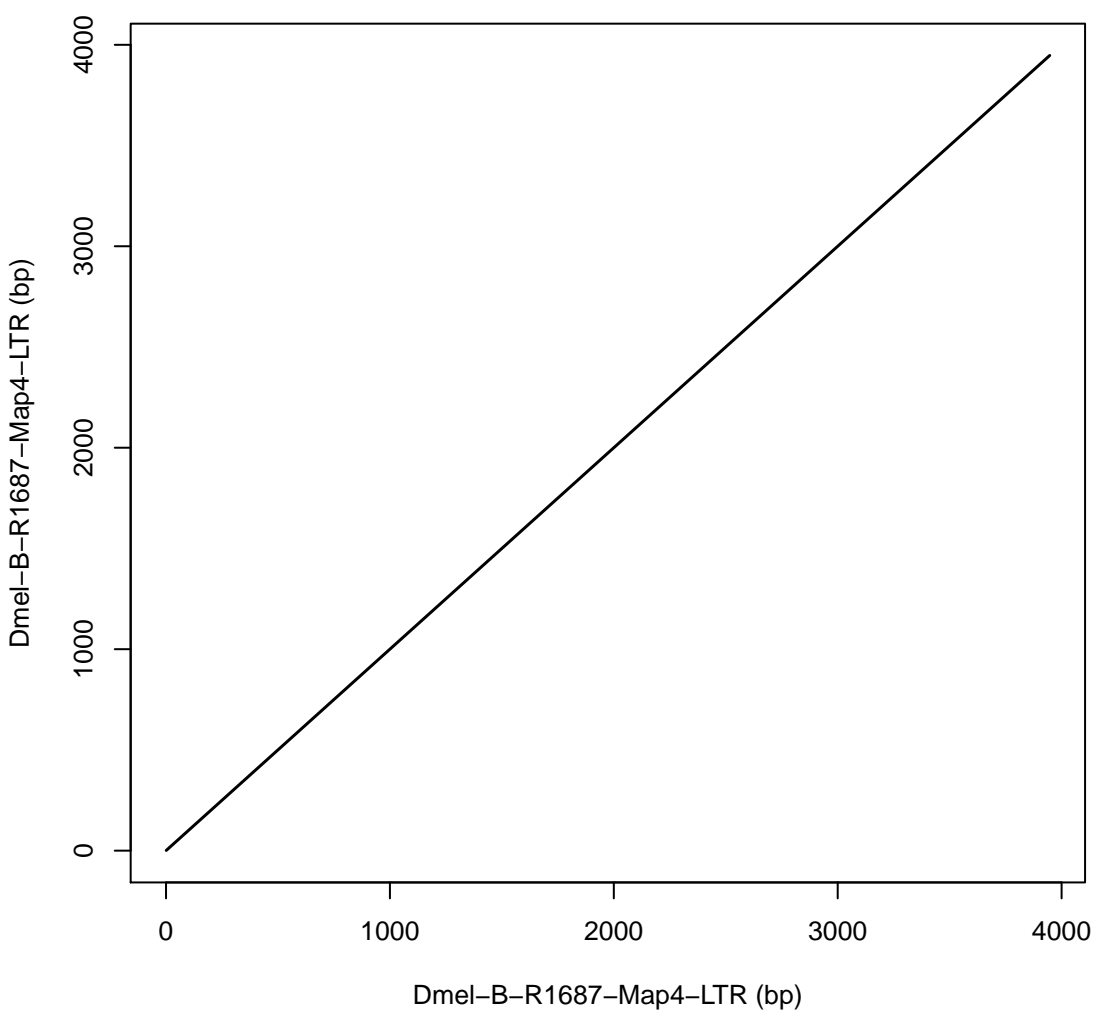consensus size: 3947bp; fragments: 68; full length: 8 (>=3552.3bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

BEL24_DR_pol LTR/Pao
Copia−11_Pru LTR/Copia
Baggins1_pol LINE/R1−LOA
BEL−63_CQ_pol LTR/Pao
DM1731 LTR/Copia