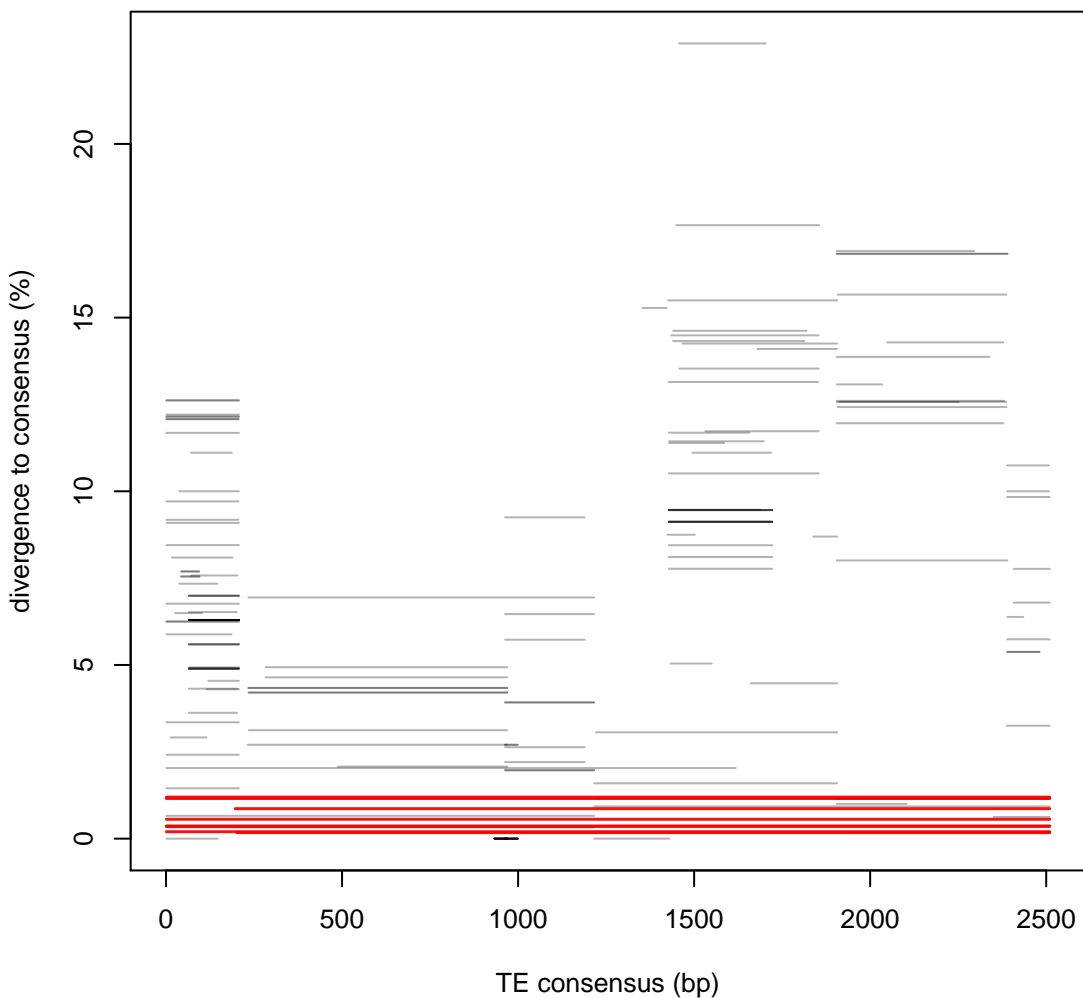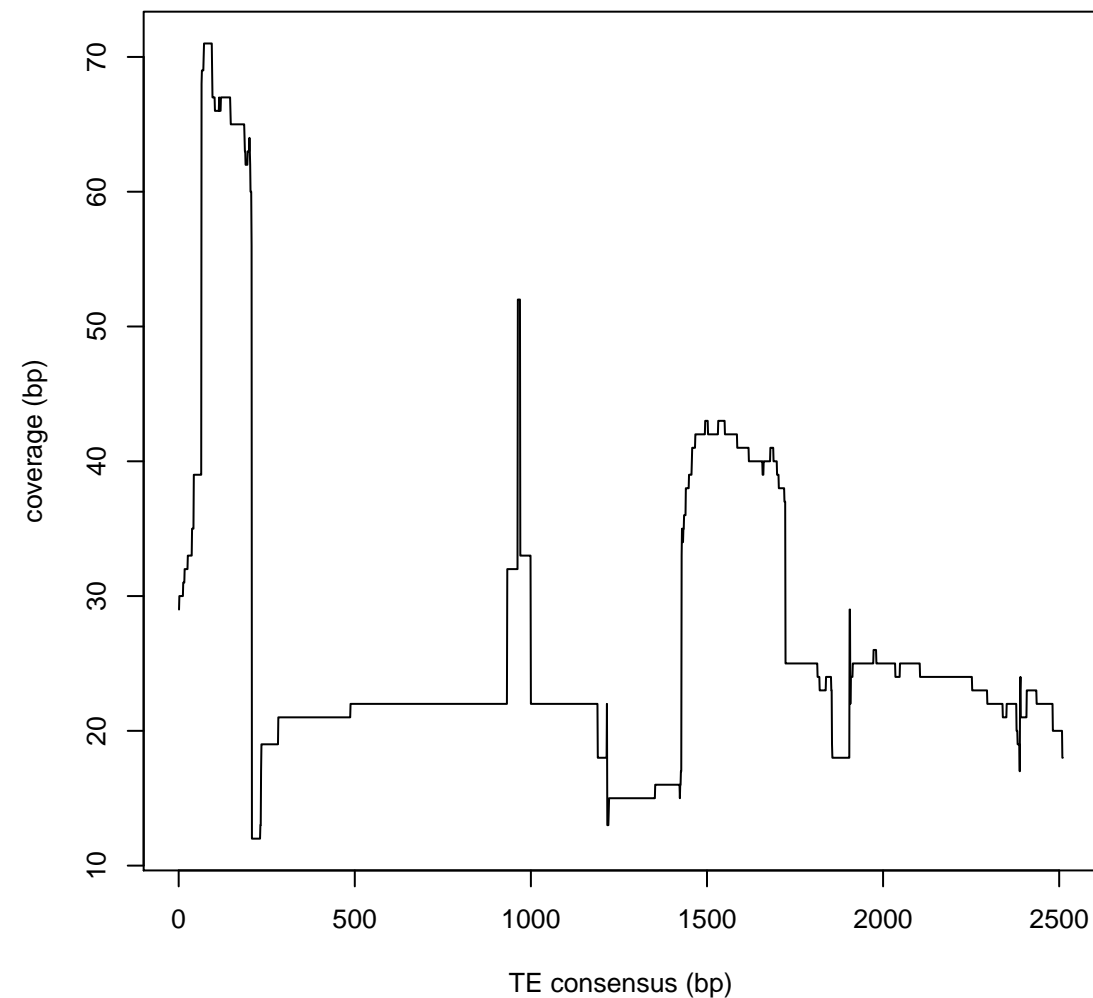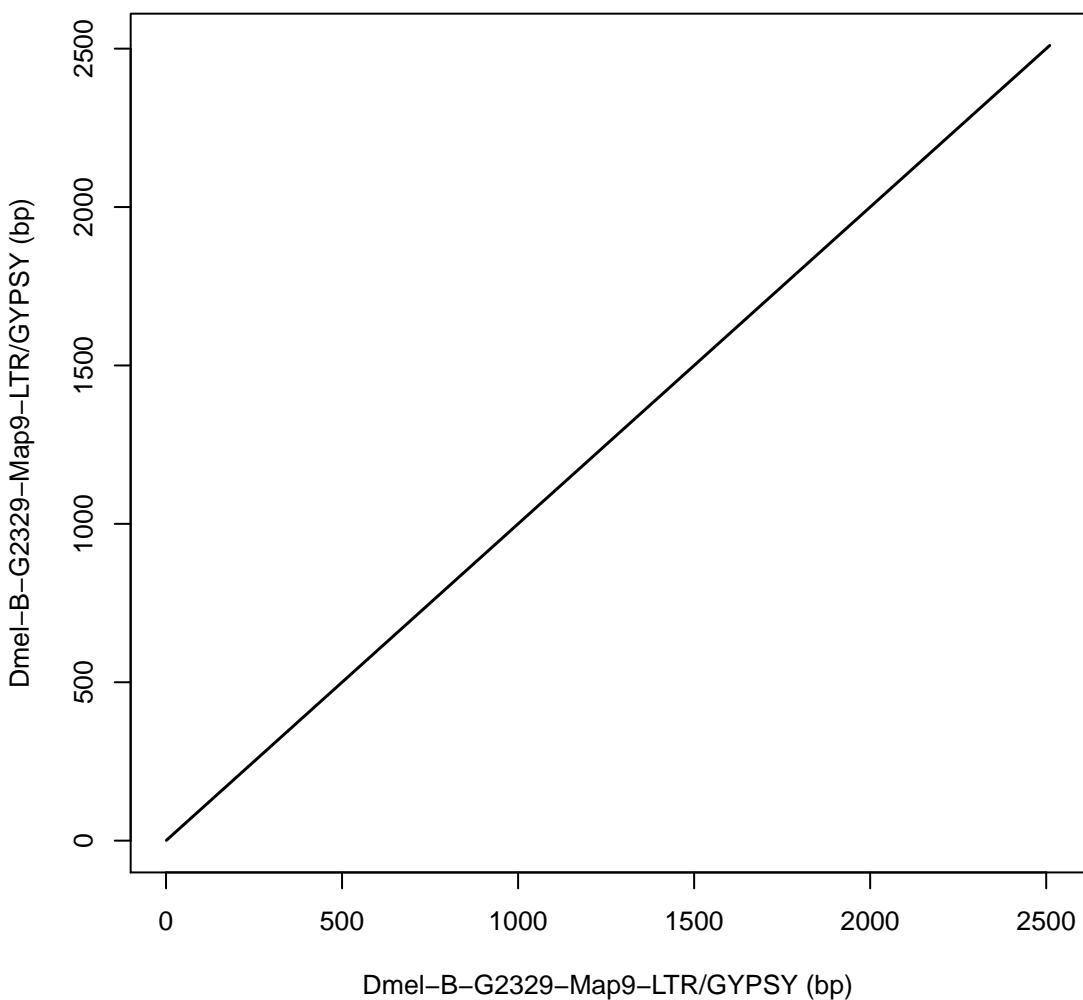**TE: Dmel−B−G2329−Map9−LTR/GYPSY**
consensus size: 2510bp; fragments: 182; full length: 8 (>=2259bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Copia−101_BRa LTR/Copia