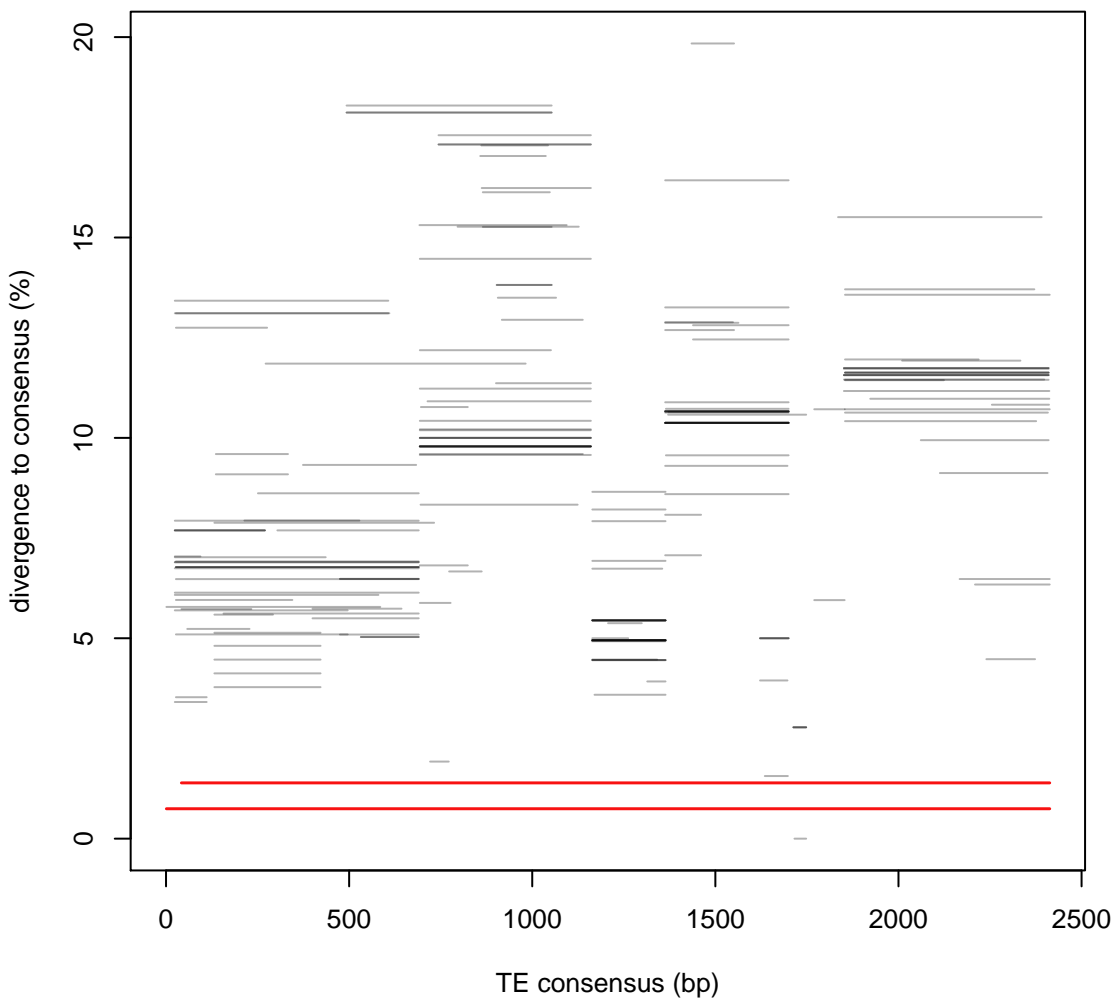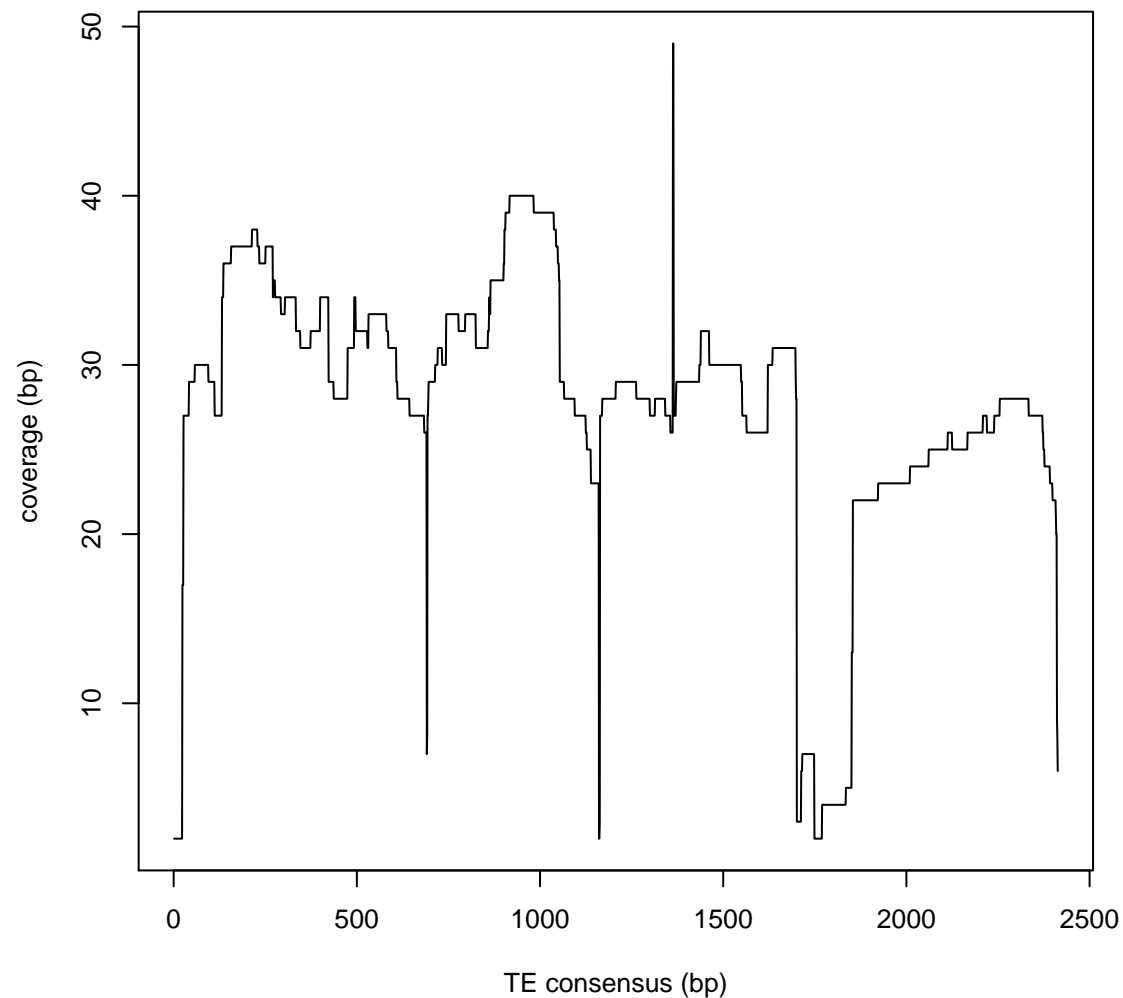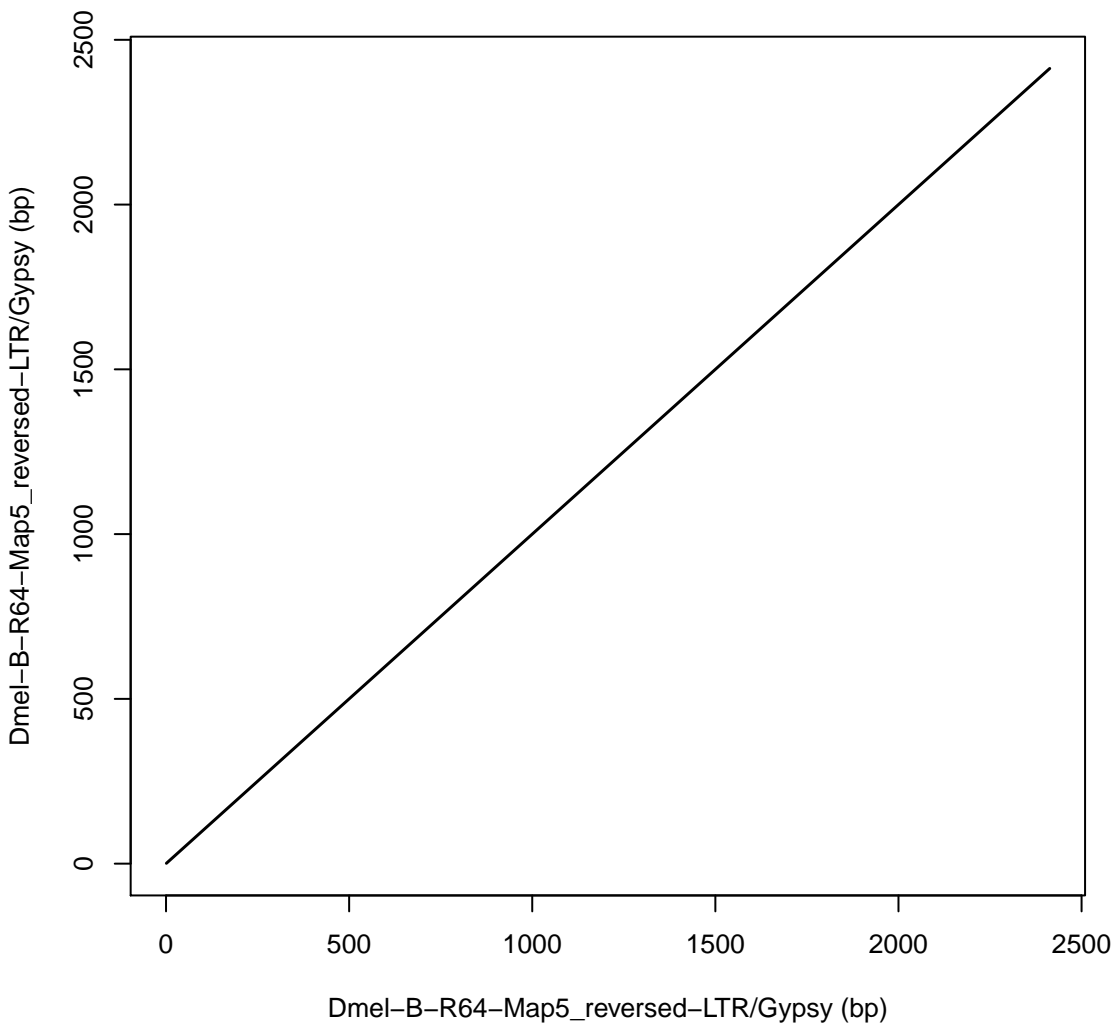**TE: Dmel−B−R64−Map5_reversed−LTR/Gypsy**
consensus size: 2413bp; fragments: 191; full length: 2 (>=2171.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected