TE: Dmel−B−R1438−Map6_reversed−CLASSI/LTR/Gypsy
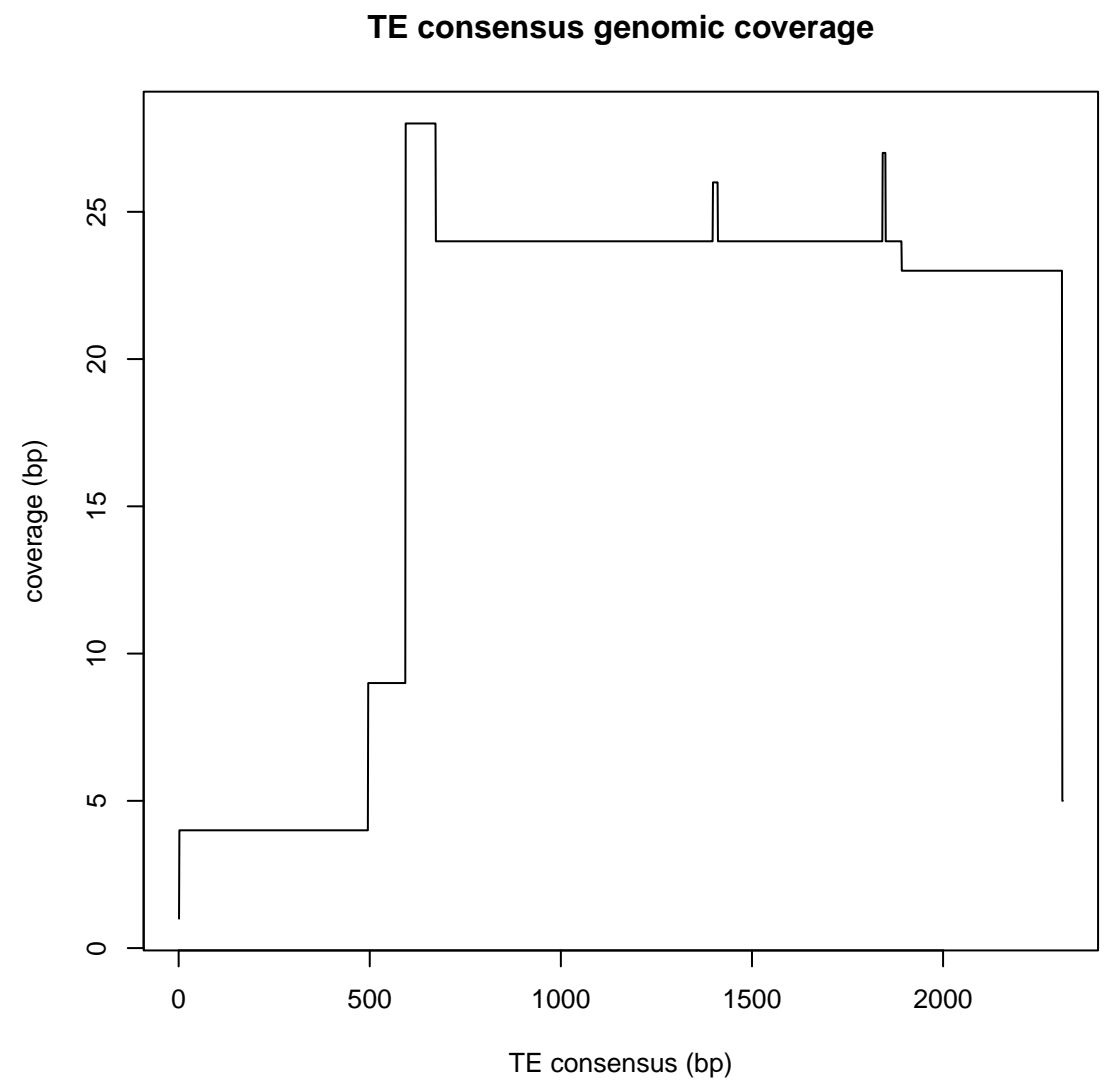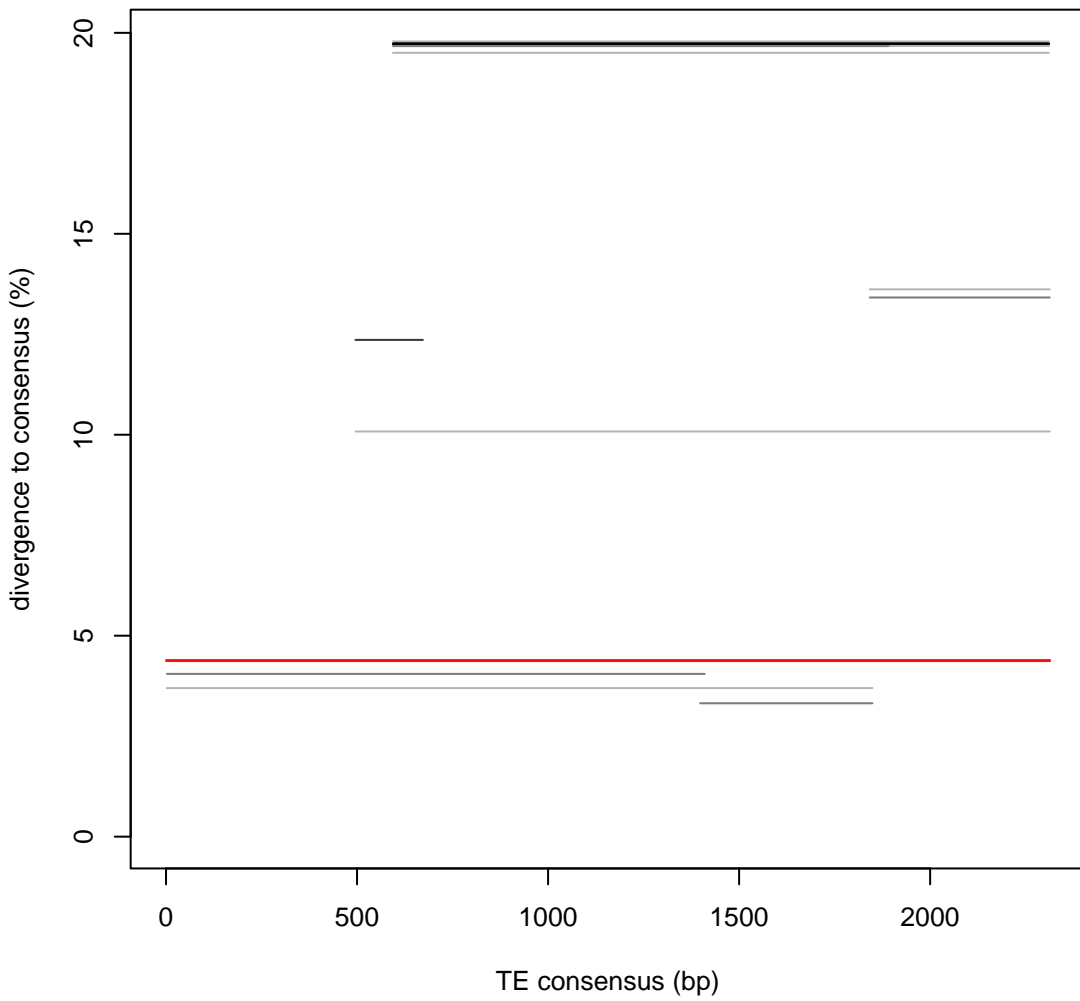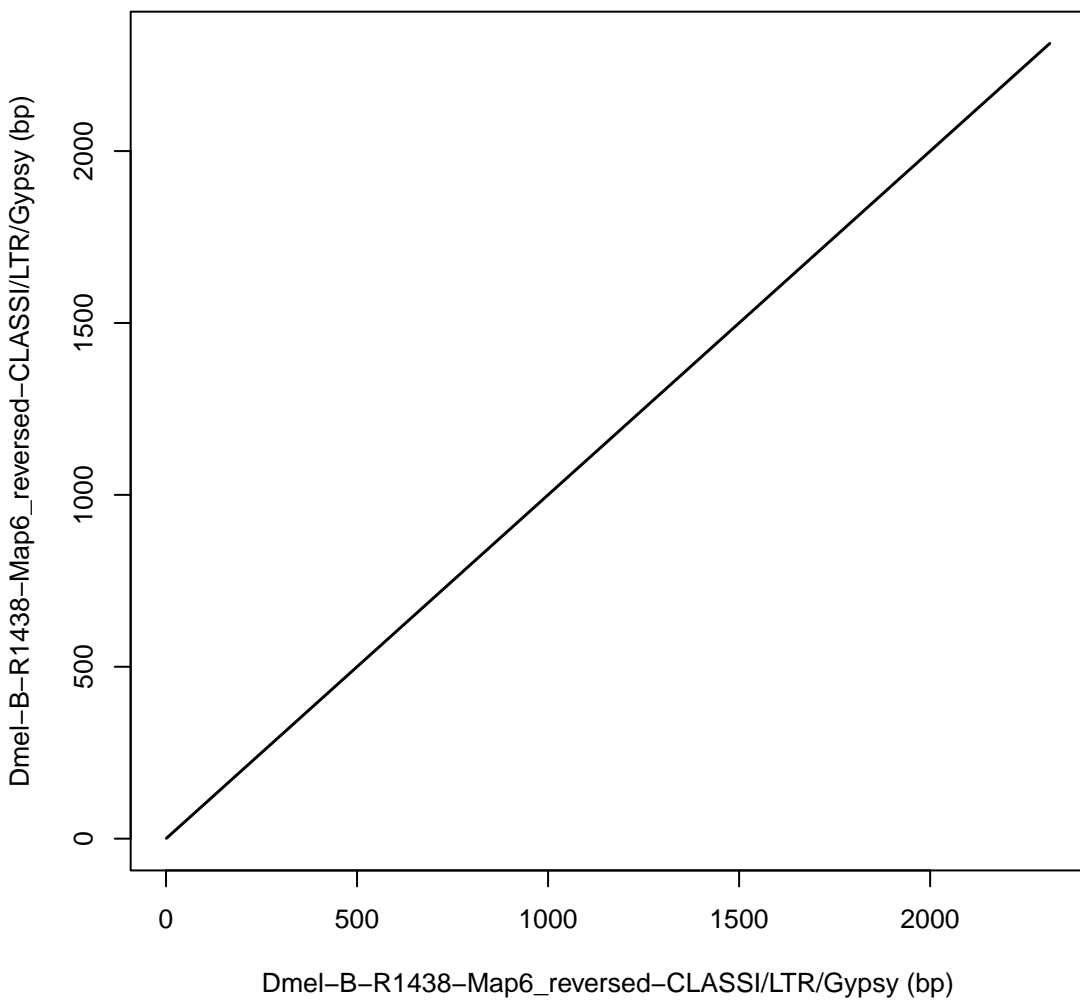**consensus size: 2313bp; fragments: 33; full length: 1 (>=2081.7bp)**

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected