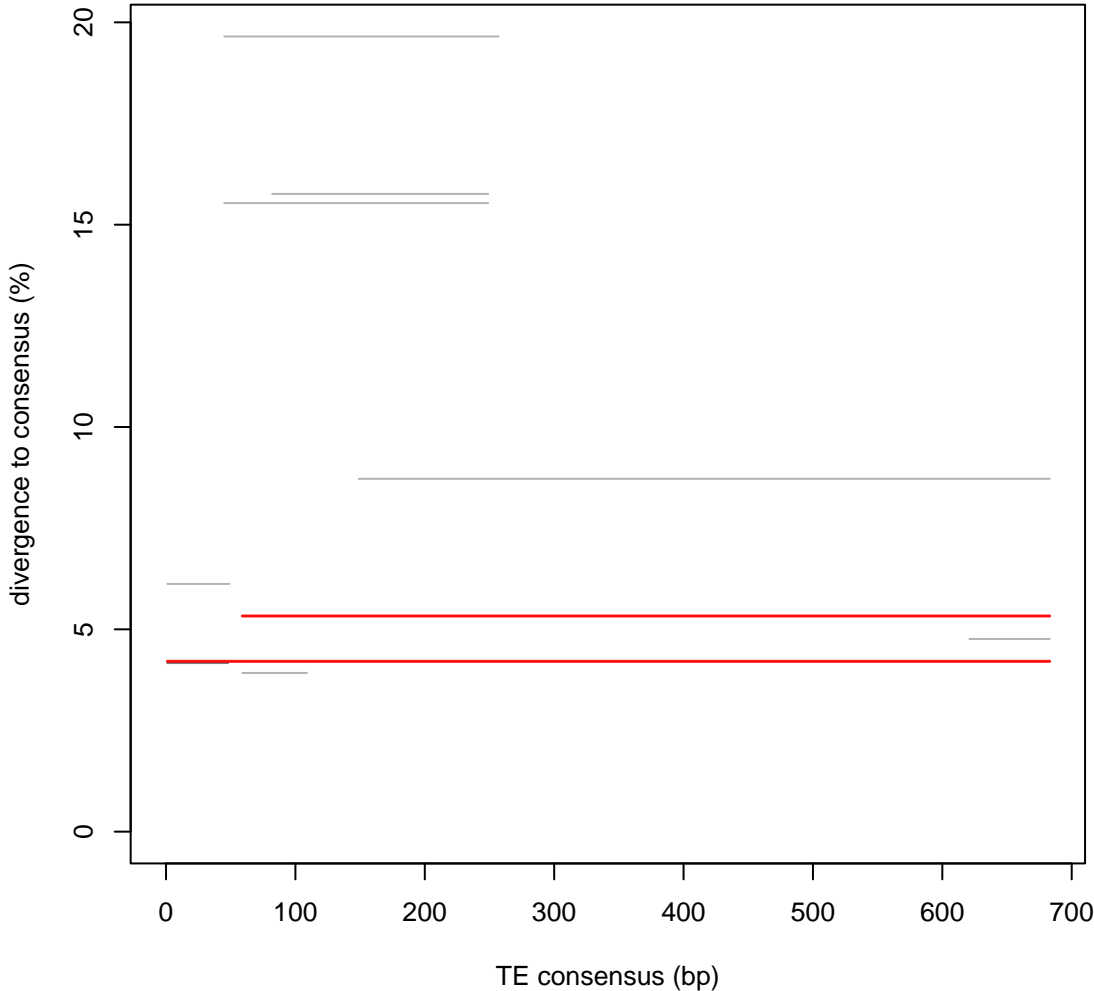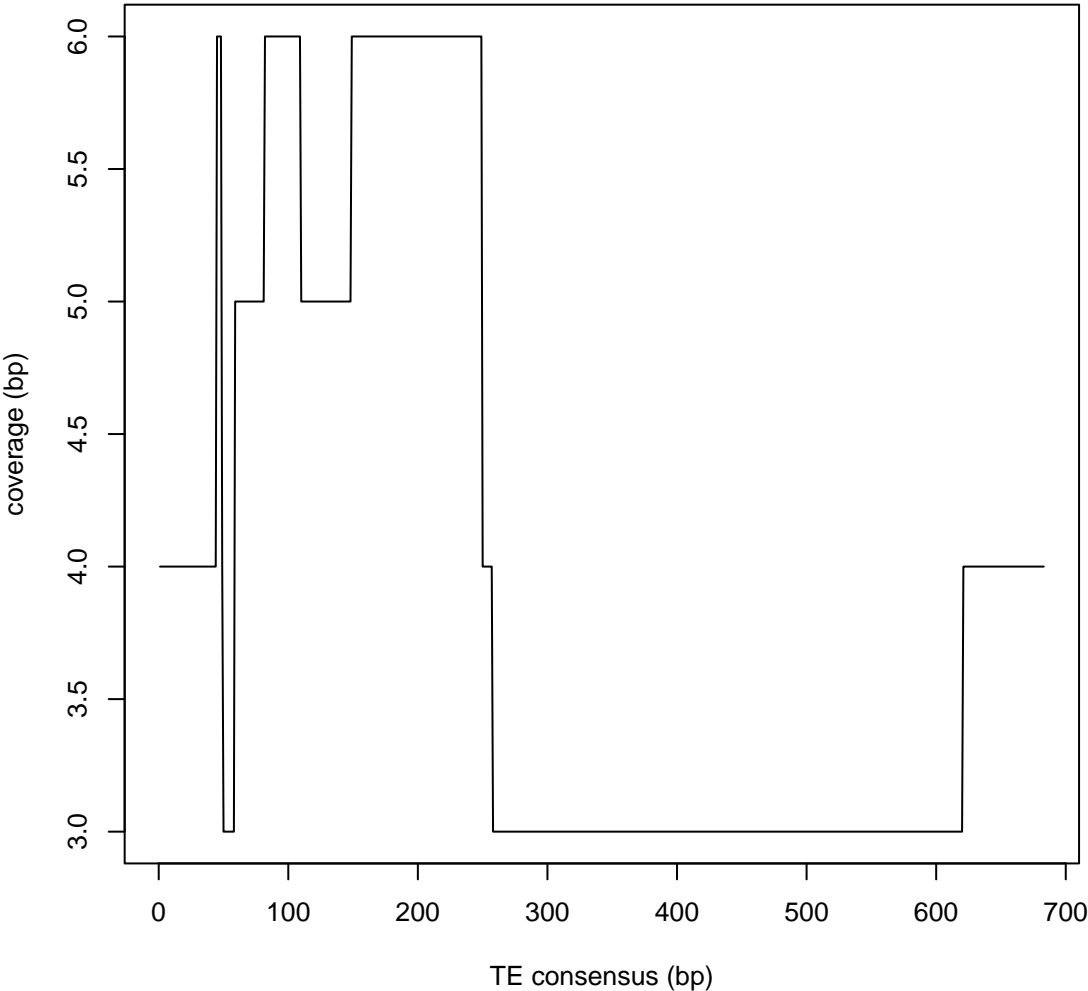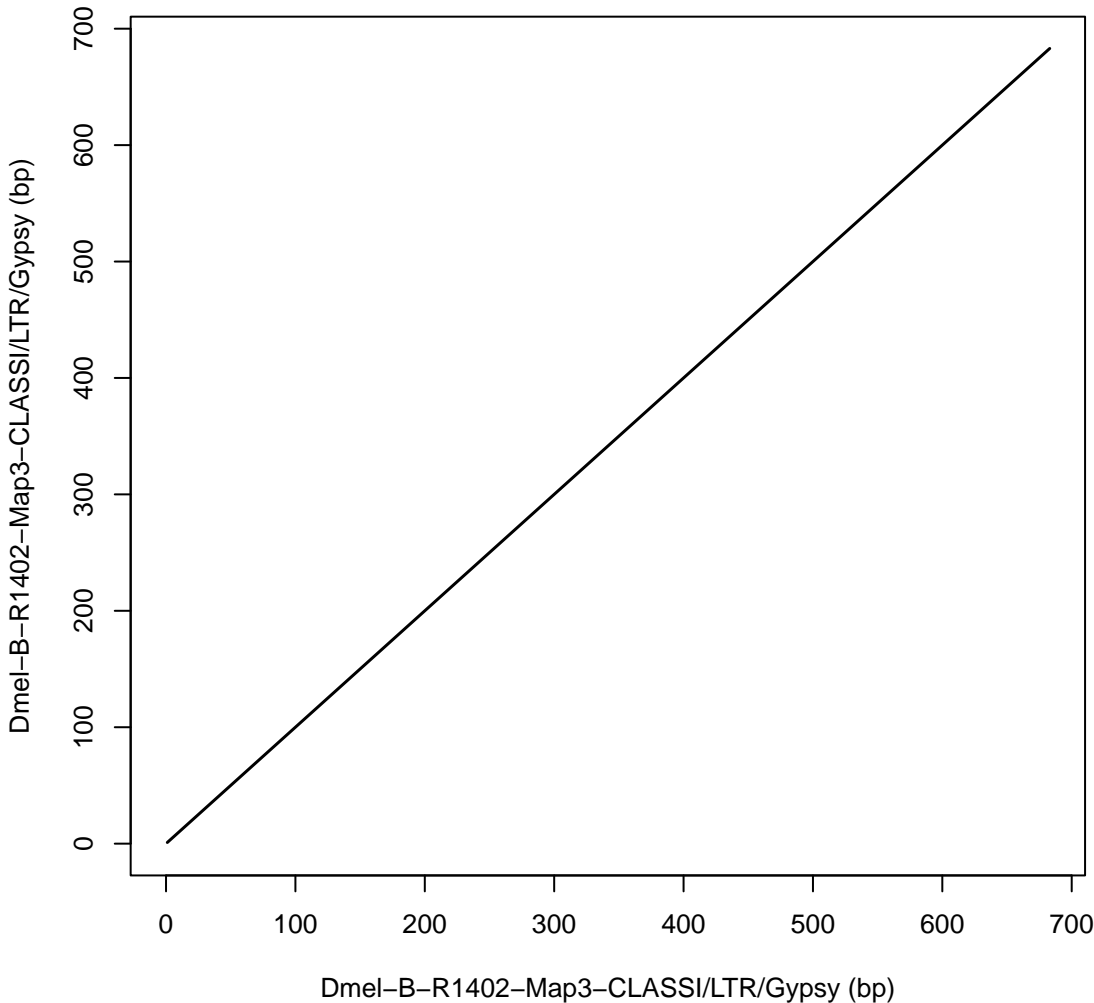**TE: Dmel−B−R1402−Map3−CLASSI/LTR/Gypsy**
consensus size: 683bp; fragments: 11; full length: 2 (>=614.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected