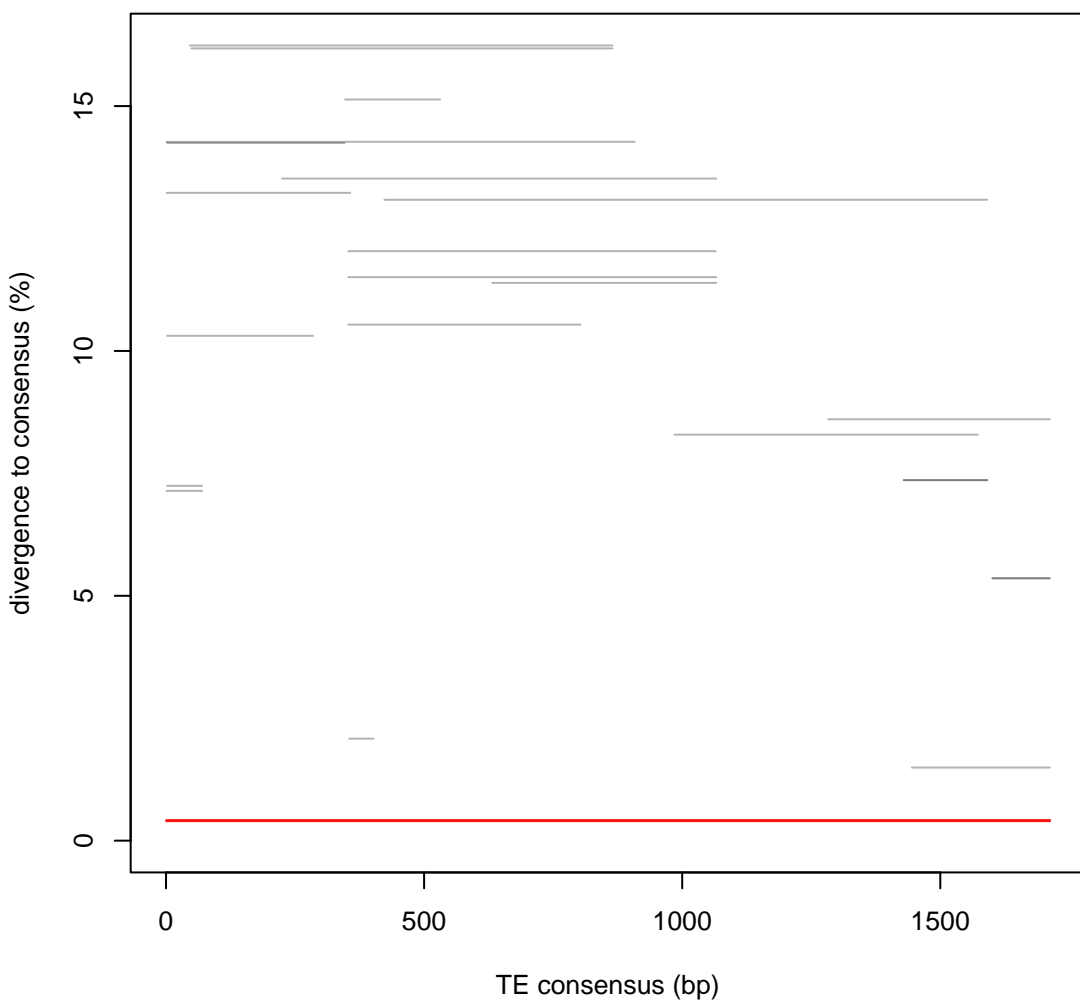
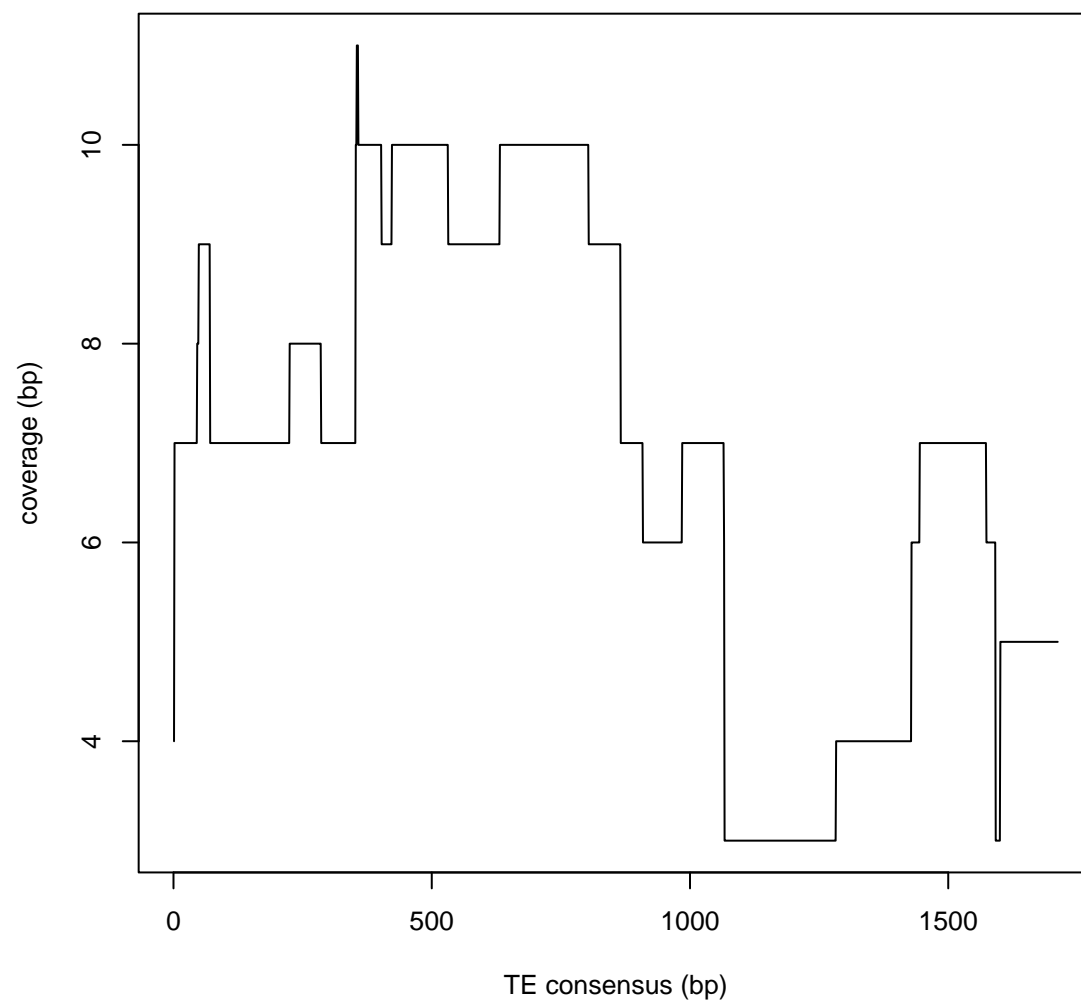


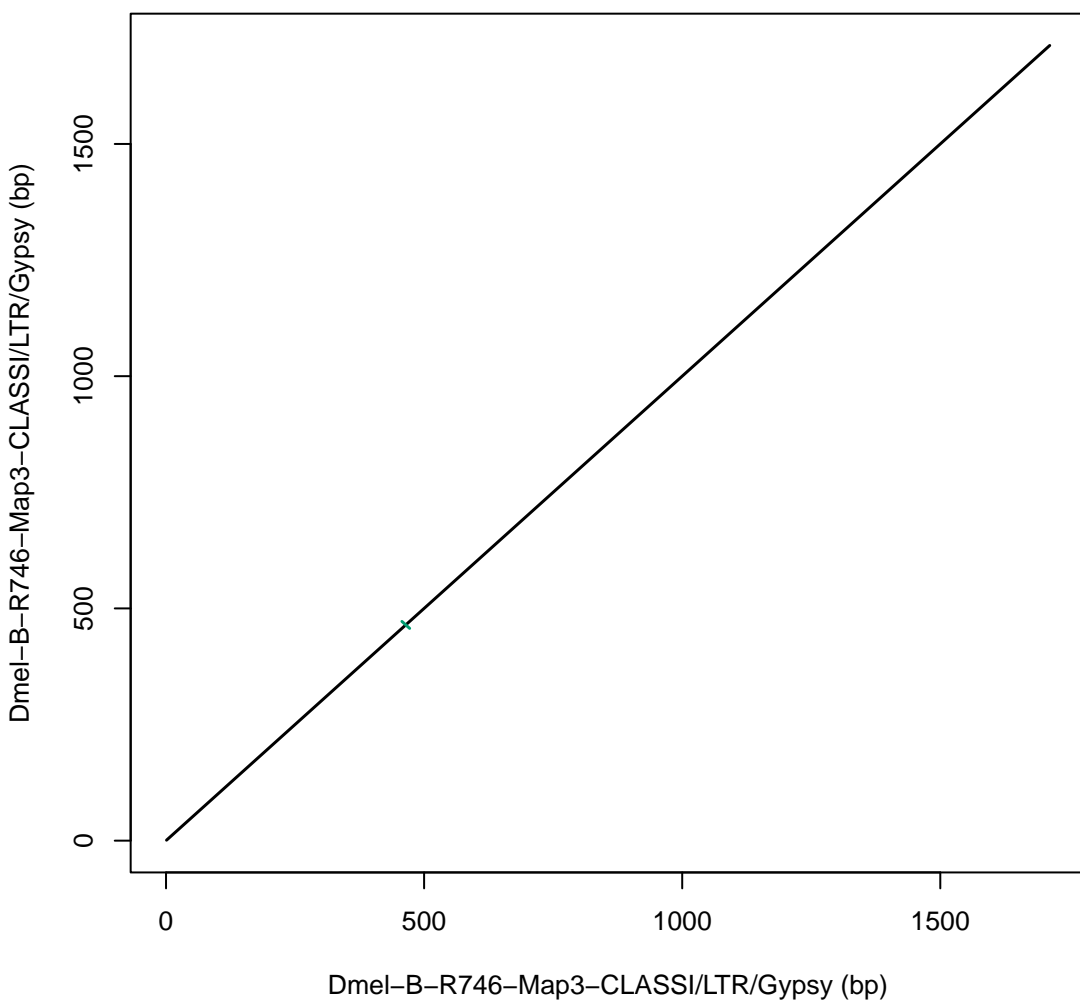
TE: Dmel-B-R746-Map3-CLASSI/LTR/Gypsy
consensus size: 1712bp; fragments: 24; full length: 1 (≥ 1540.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

