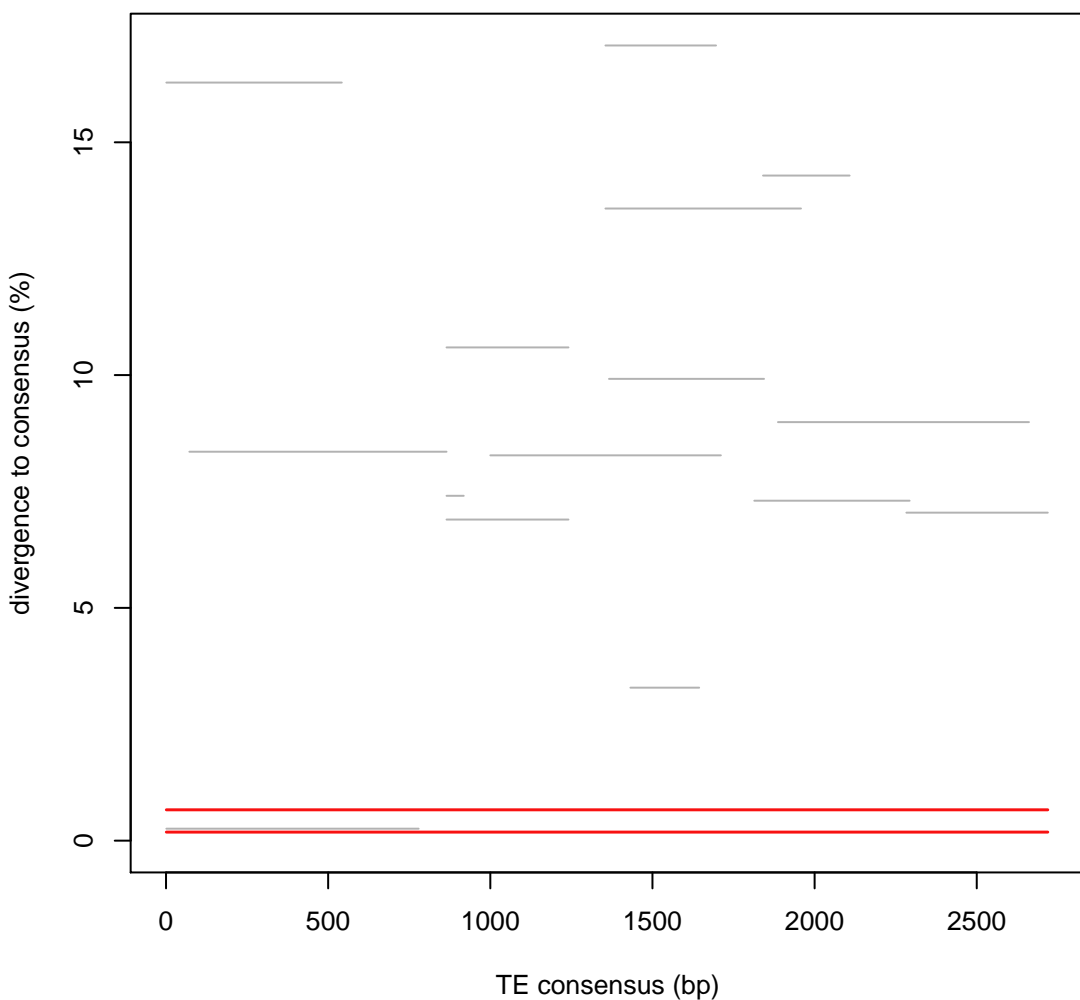
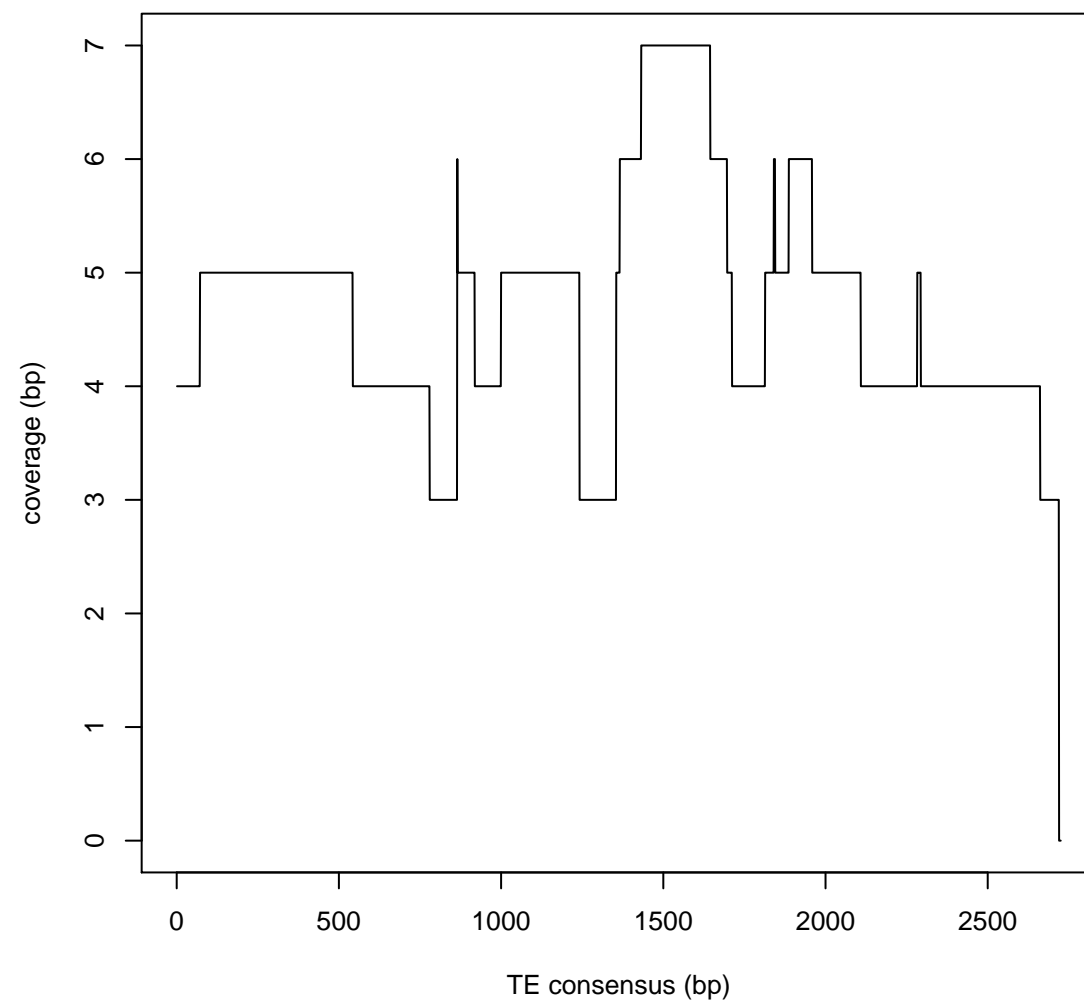


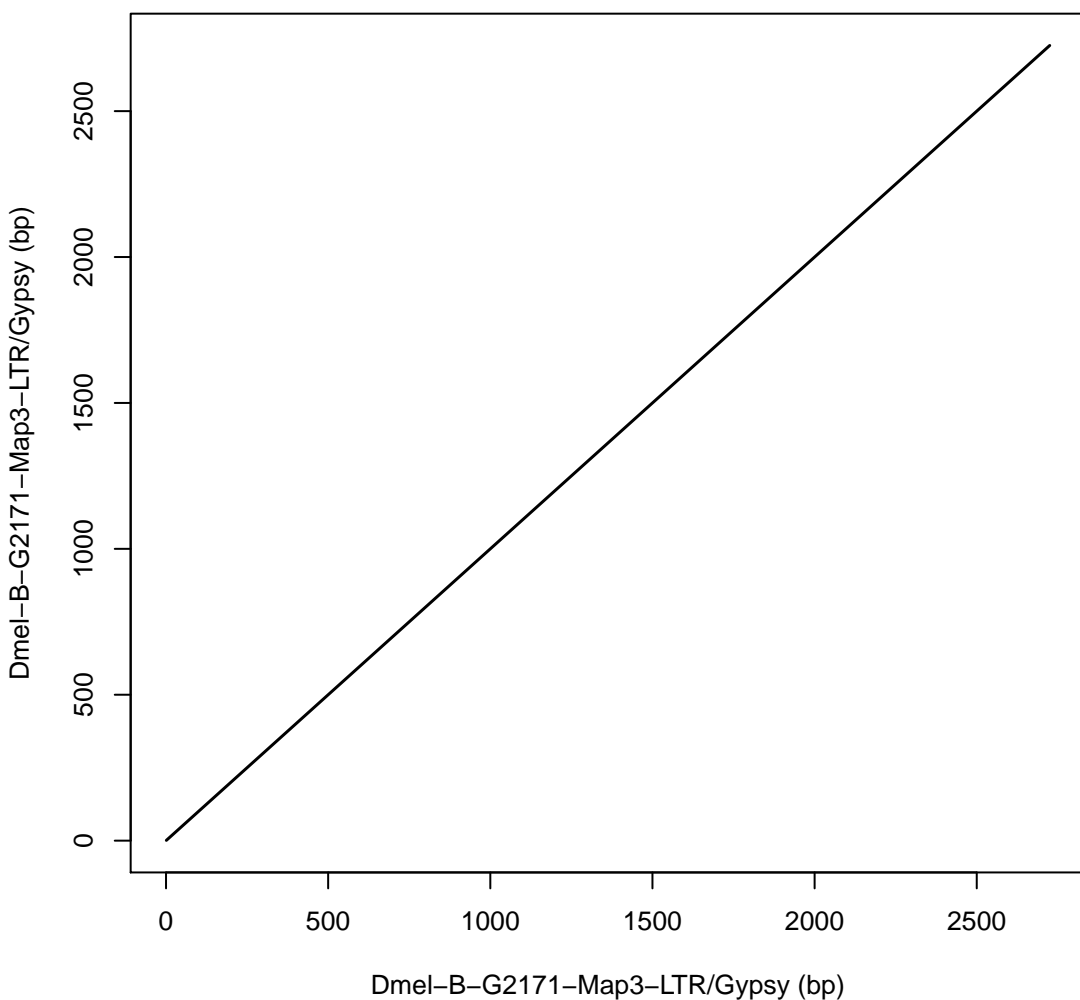
TE: Dmel-B-G2171-Map3-LTR/Gypsy
consensus size: 2725bp; fragments: 17; full length: 2 (≥ 2452.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

