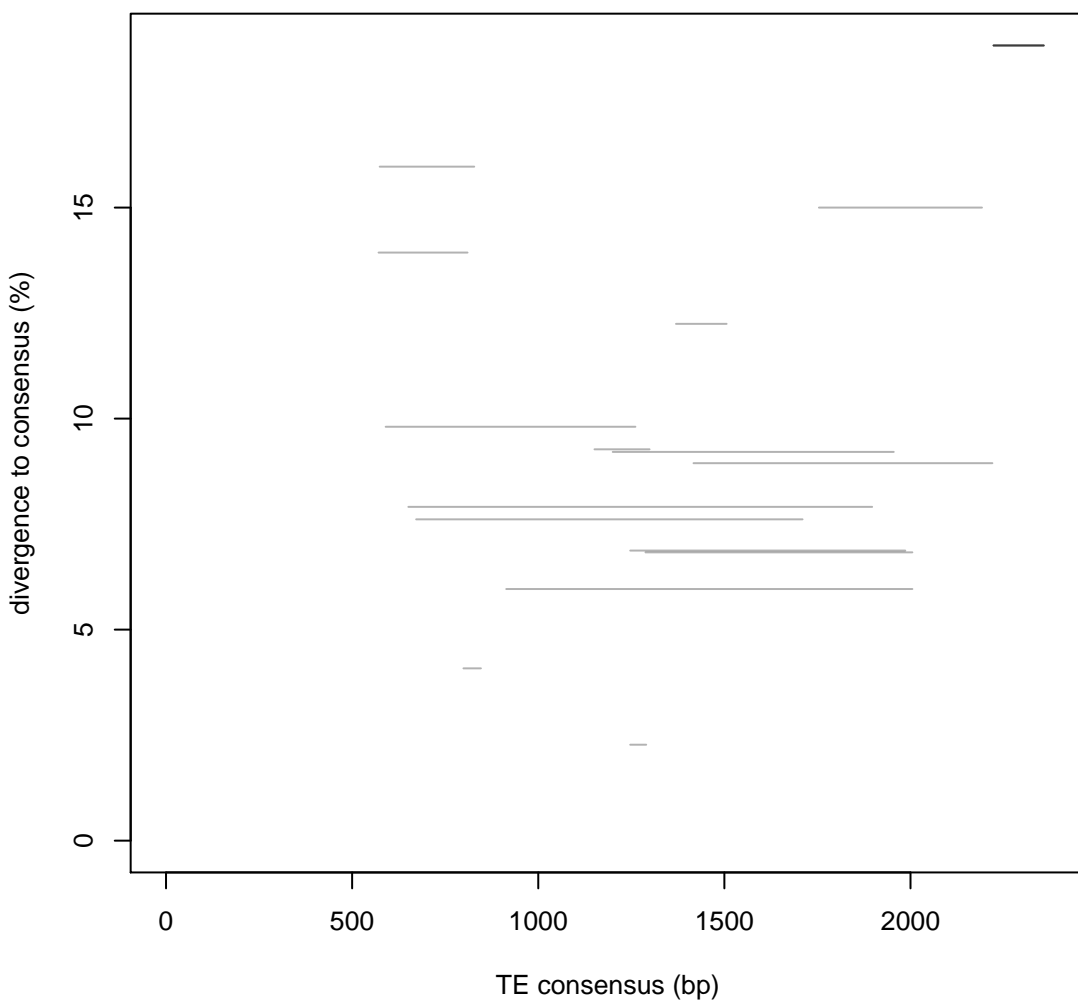
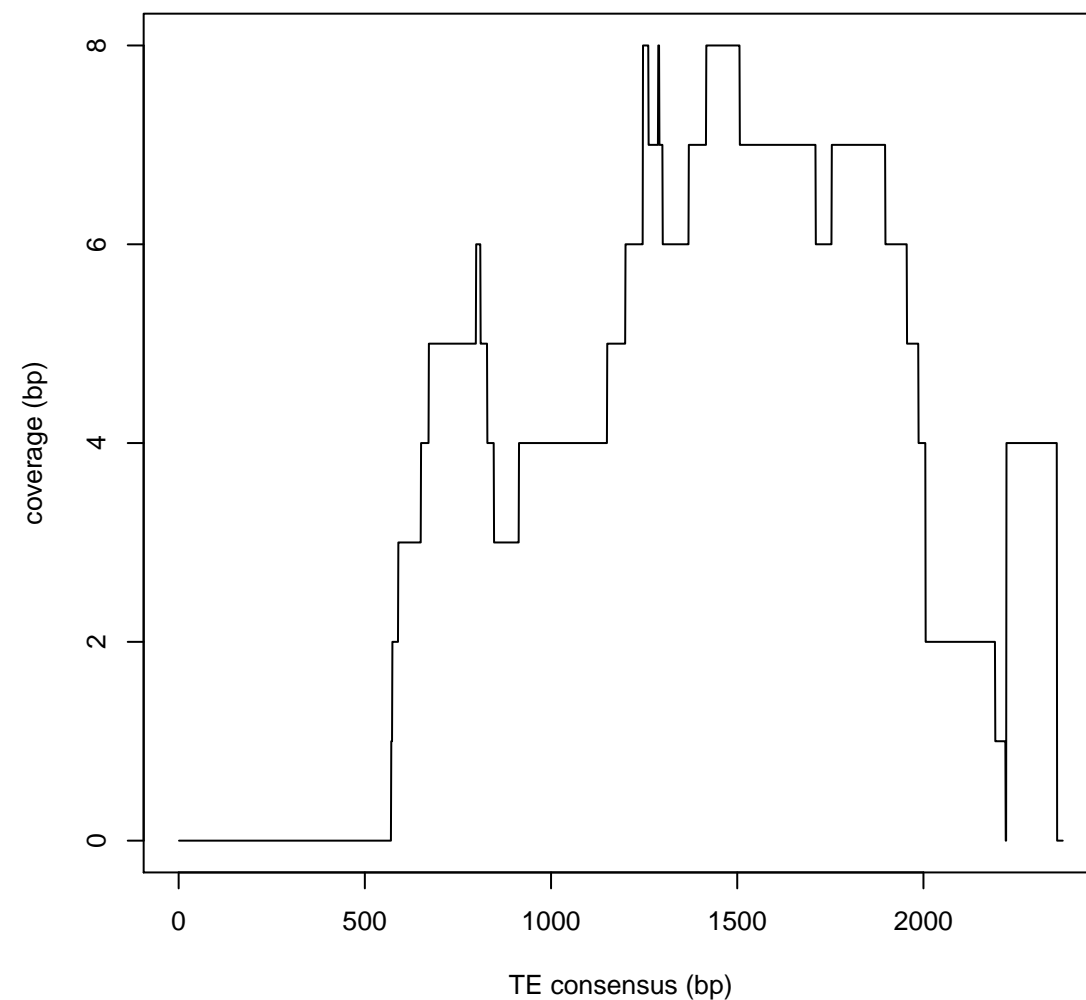


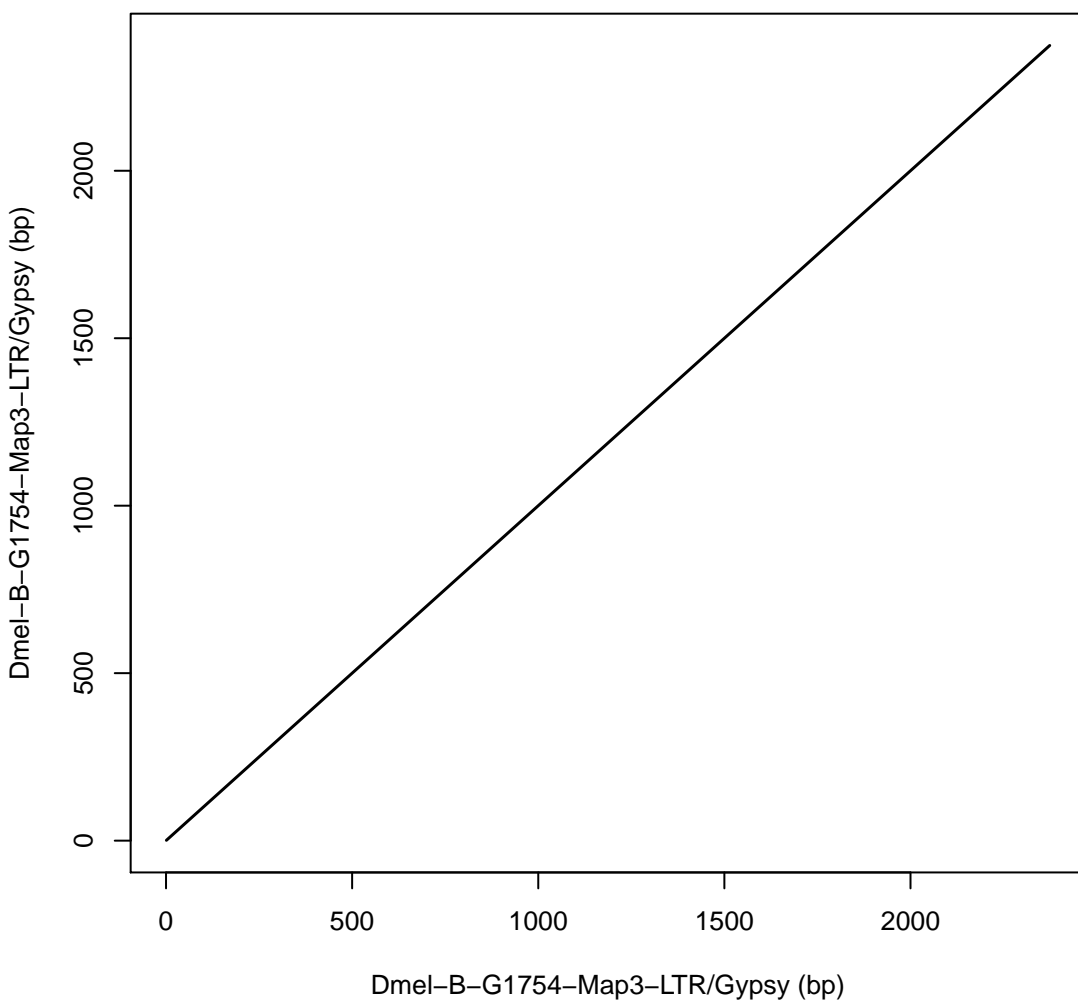
TE: Dmel-B-G1754-Map3-LTR/Gypsy
consensus size: 2374bp; fragments: 19; full length: 0 (≥ 2136.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

