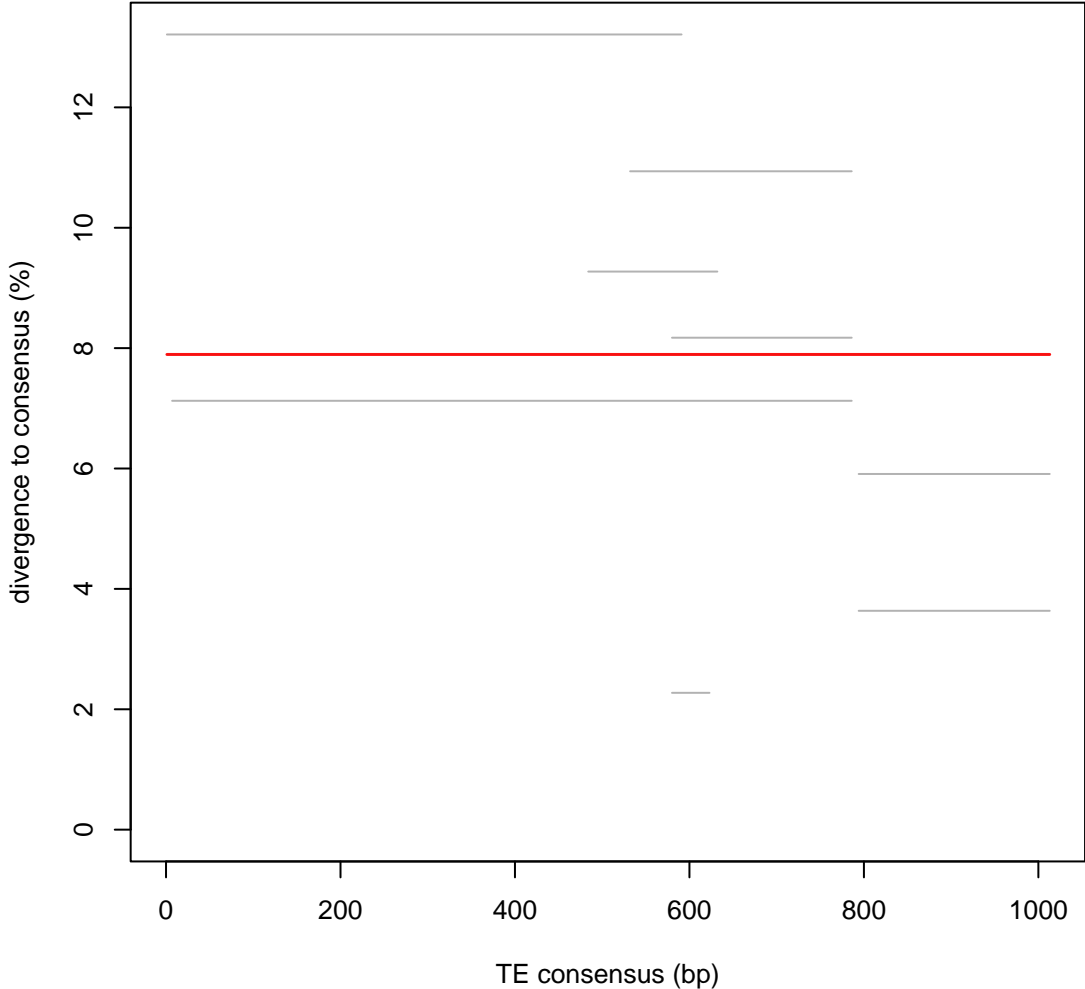
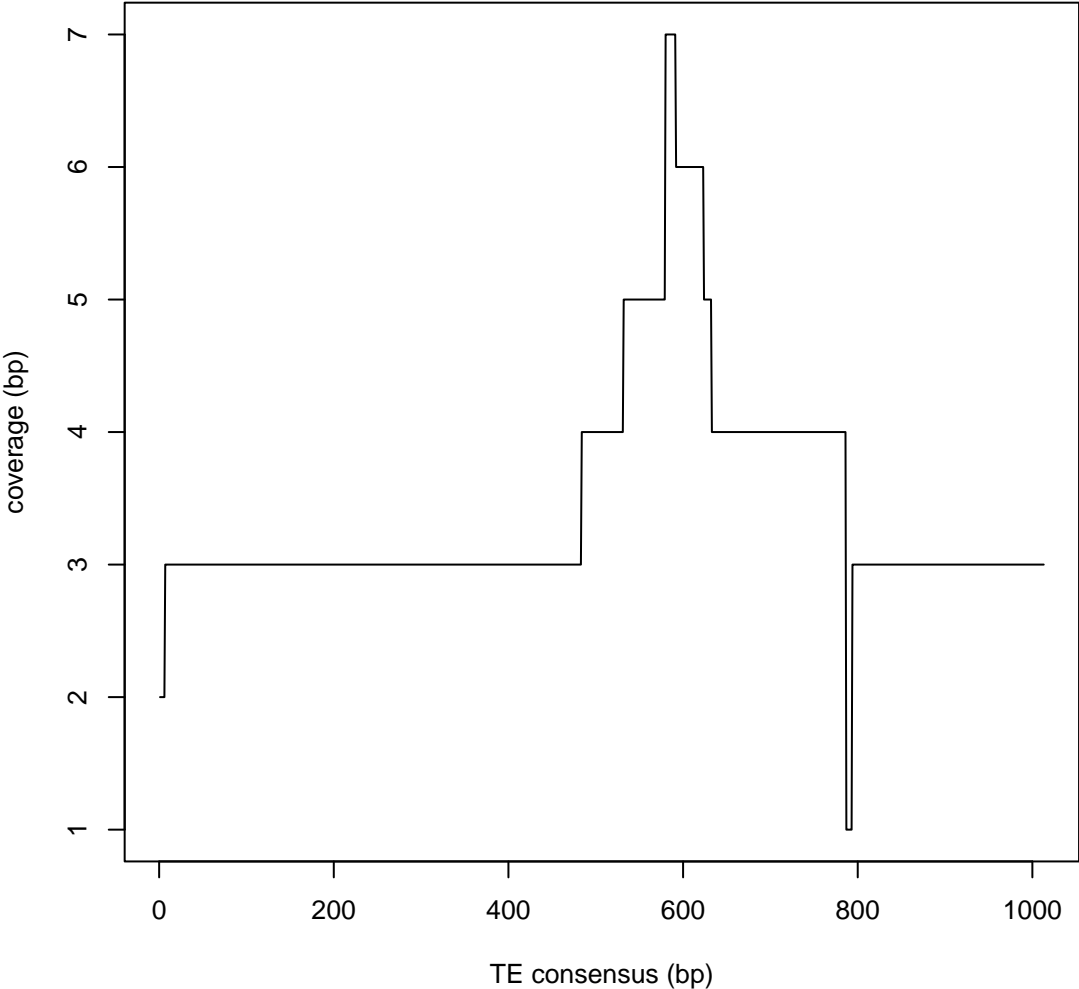


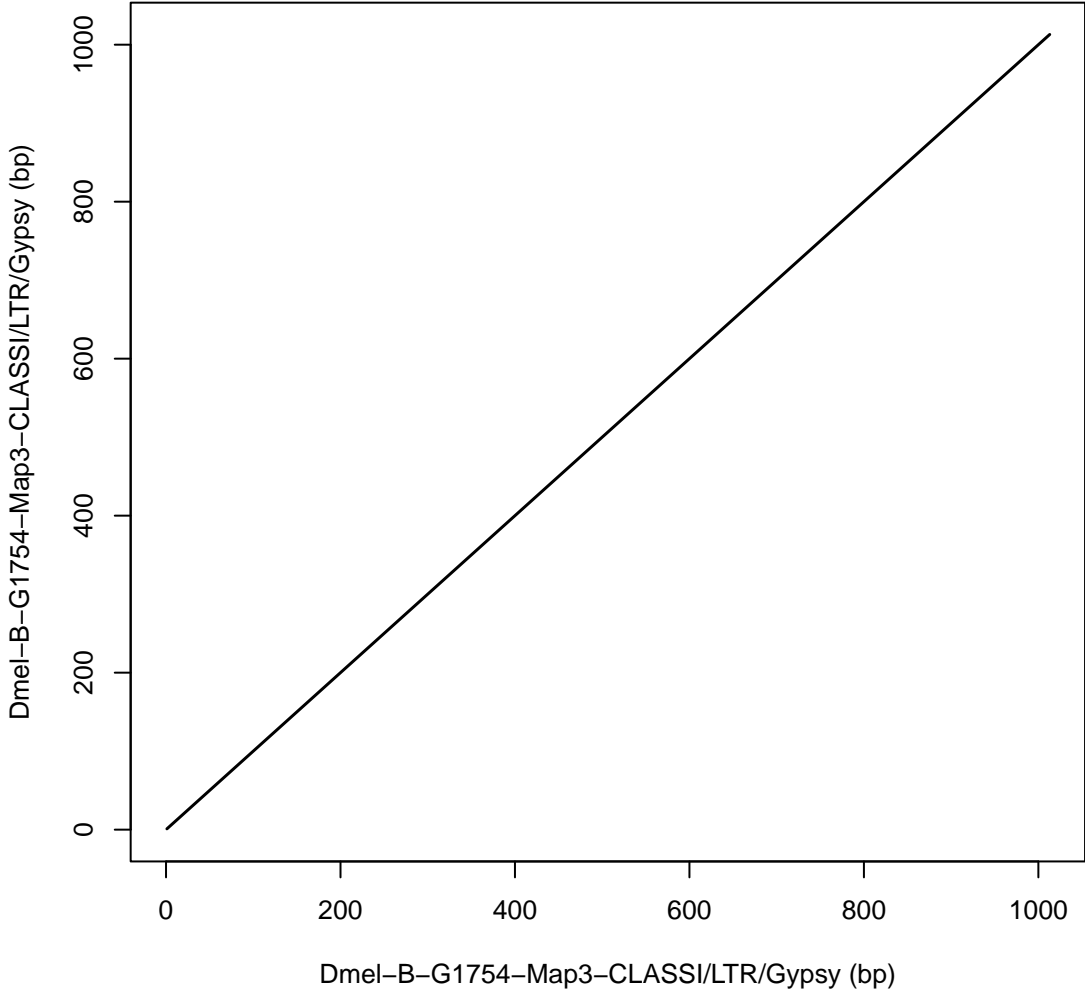
TE: Dmel-B-G1754-Map3-CLASSI/LTR/Gypsy
consensus size: 1013bp; fragments: 9; full length: 1 (>=911.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

