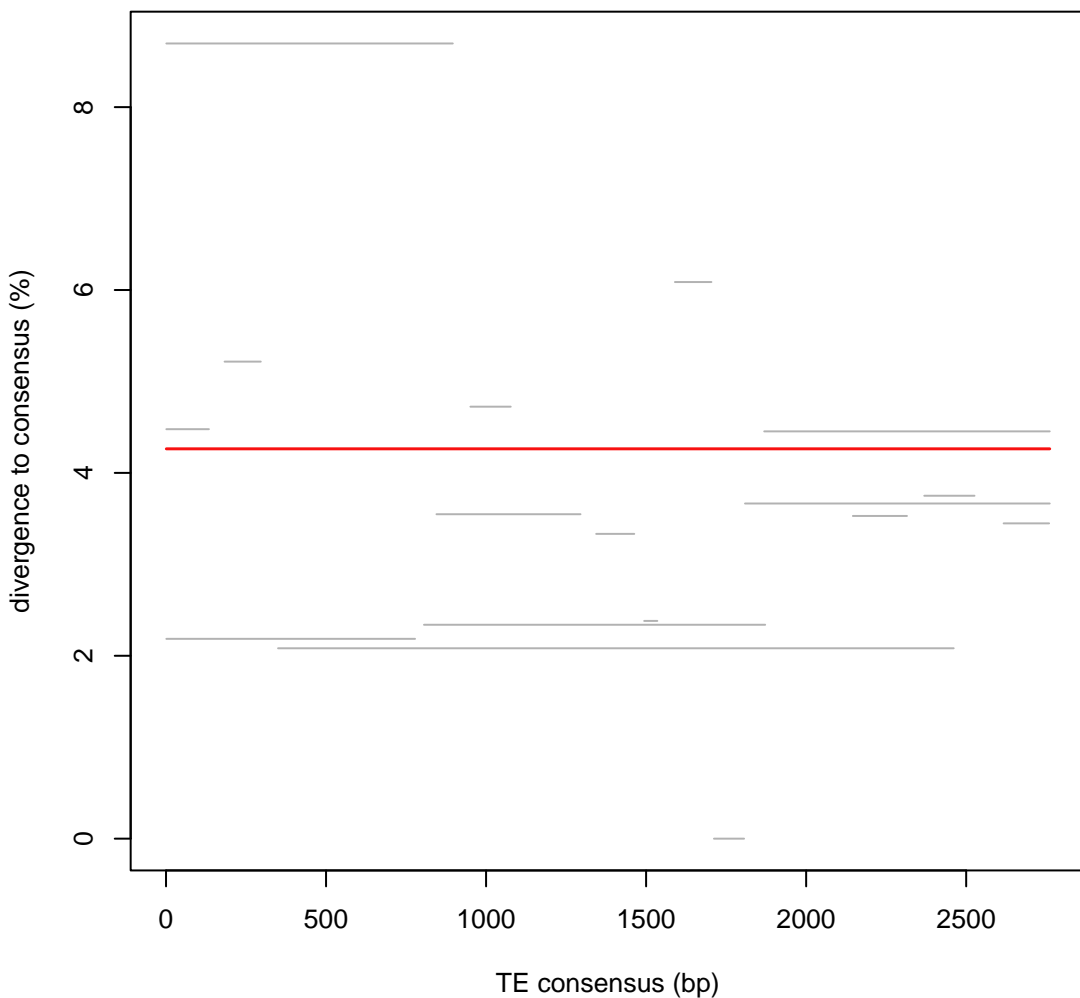
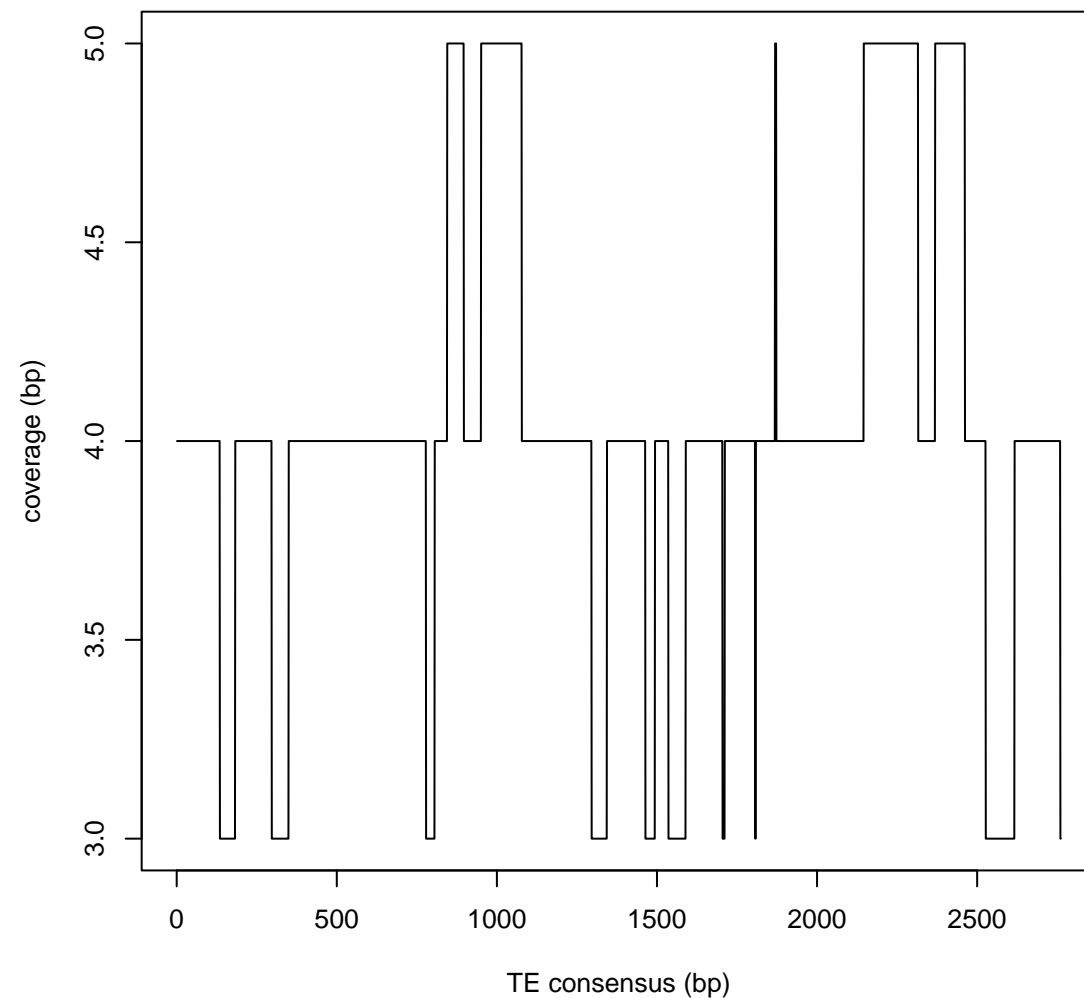


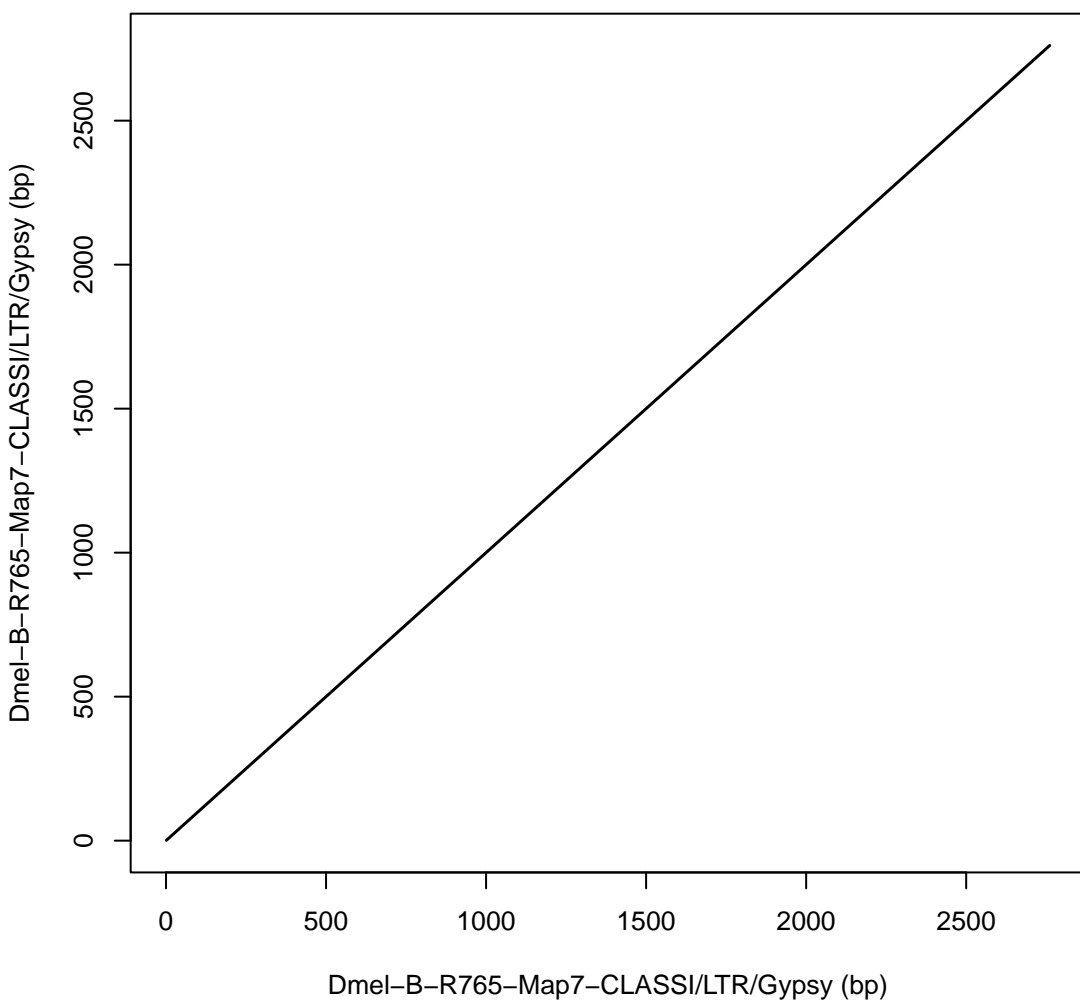
TE: Dmel-B-R765-Map7-CLASSI/LTR/Gypsy
consensus size: 2761bp; fragments: 18; full length: 1 (>=2484.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

