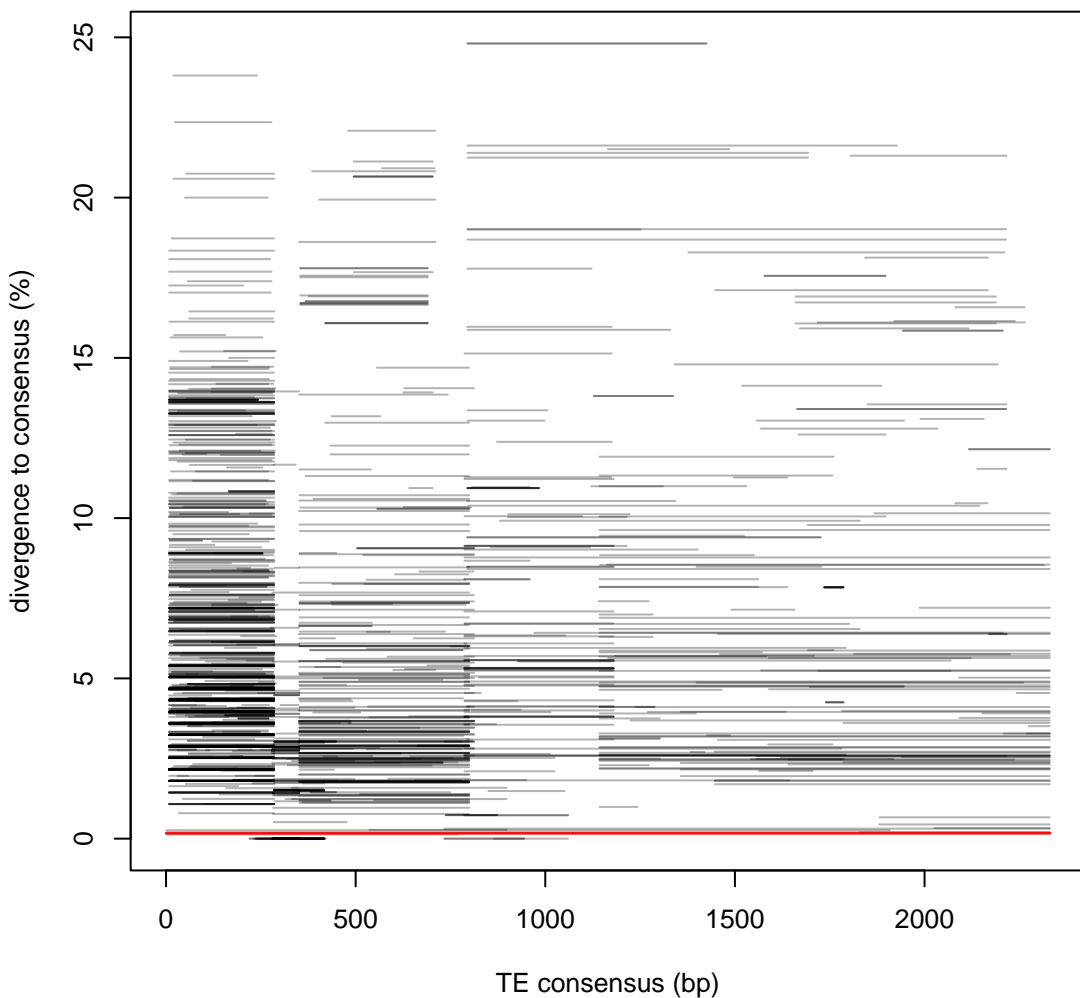
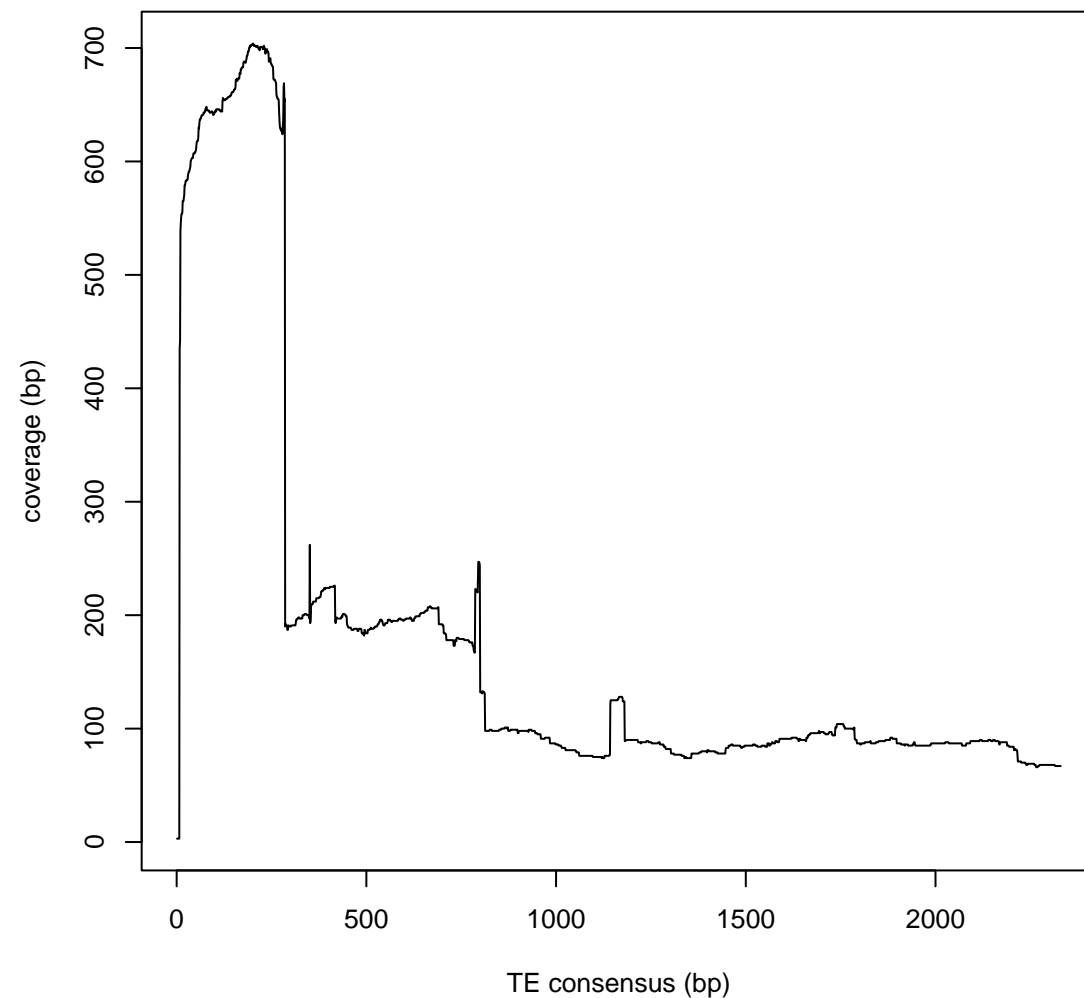


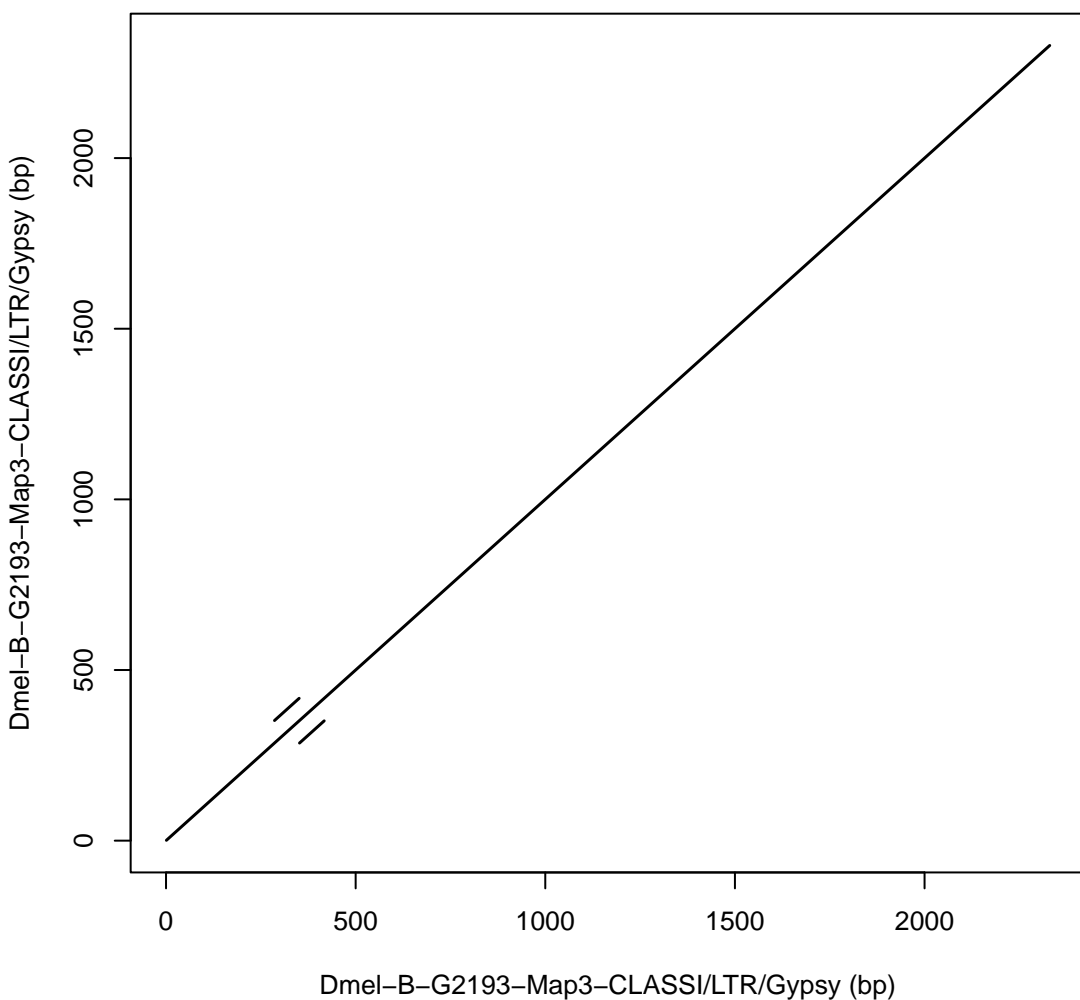
TE: Dmel-B-G2193-Map3-CLASSI/LTR/Gypsy
consensus size: 2330bp; fragments: 1454; full length: 1 (>=2097bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

