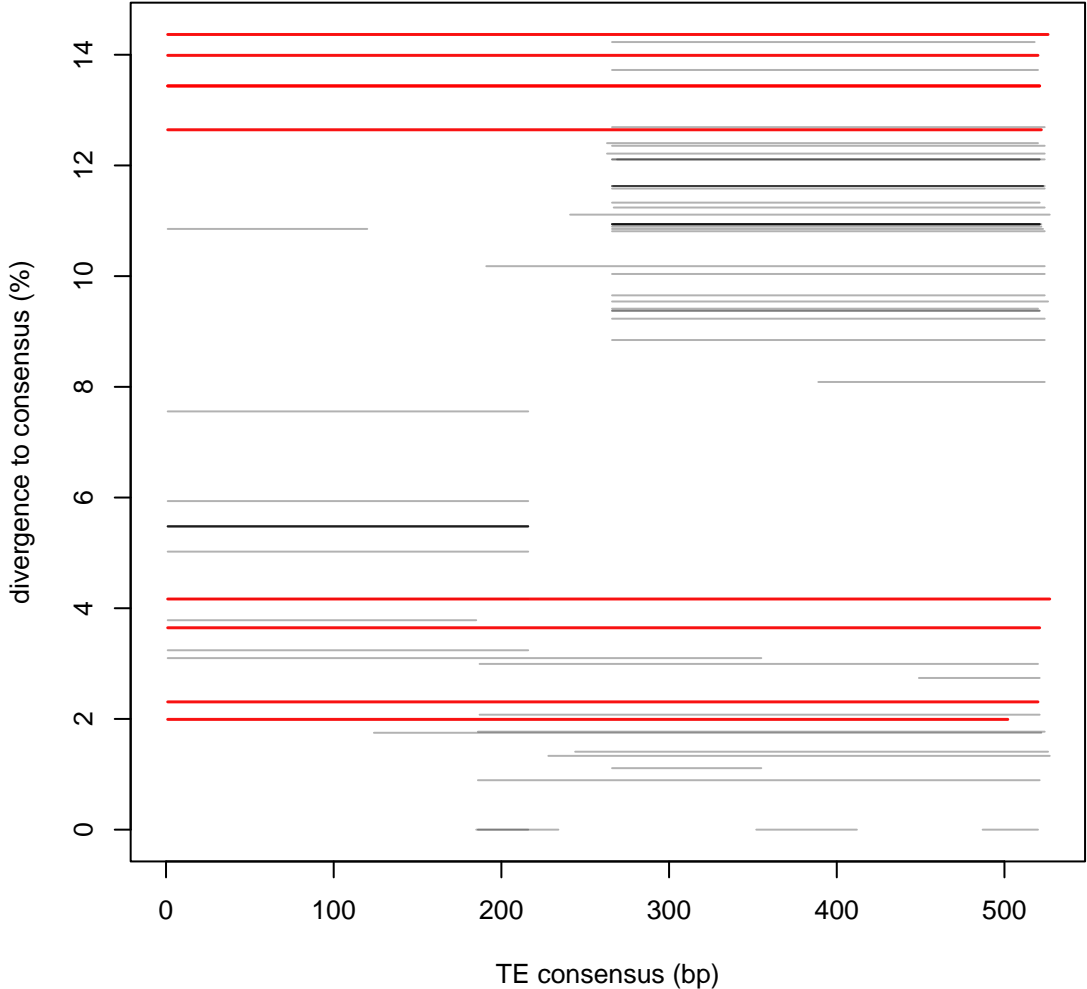
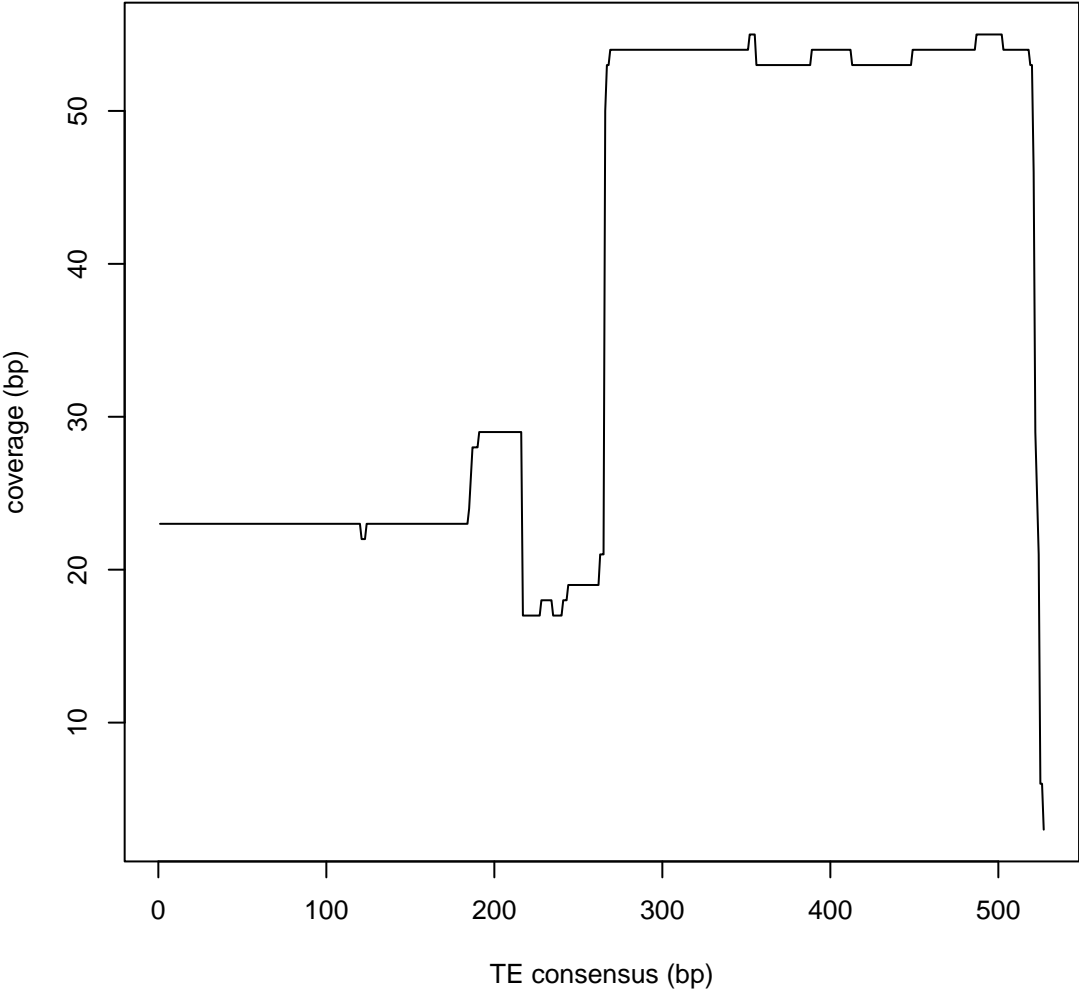


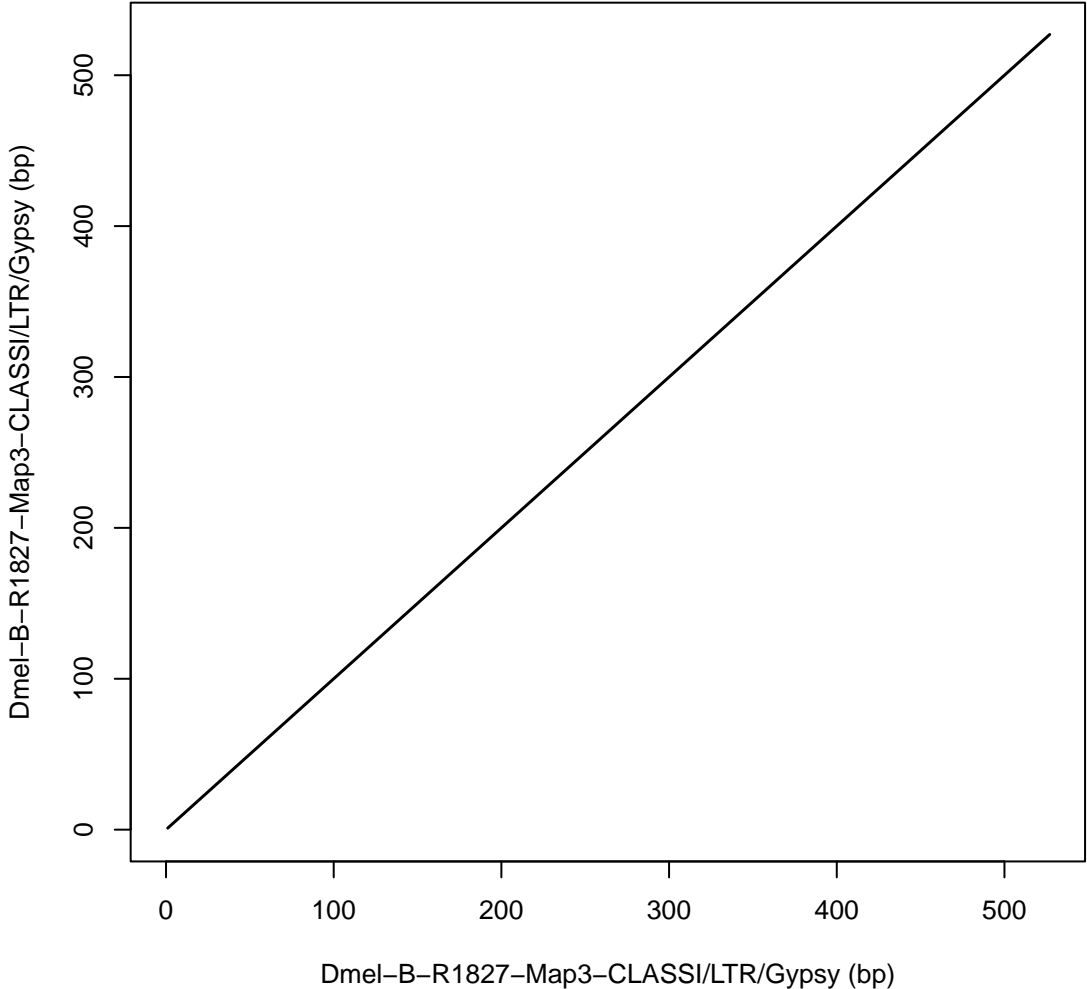
TE: Dmel-B-R1827-Map3-CLASSI/LTR/Gypsy
consensus size: 527bp; fragments: 73; full length: 9 (>=474.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

