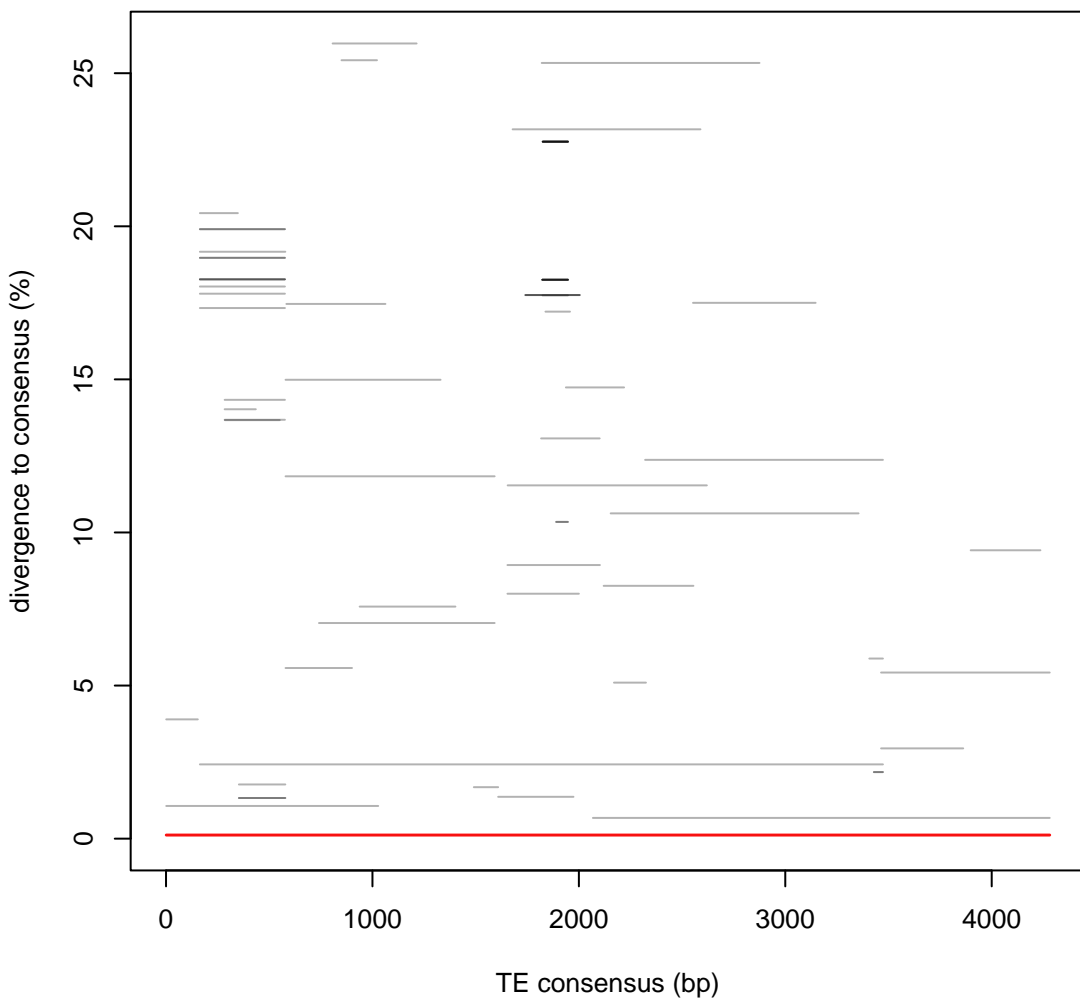
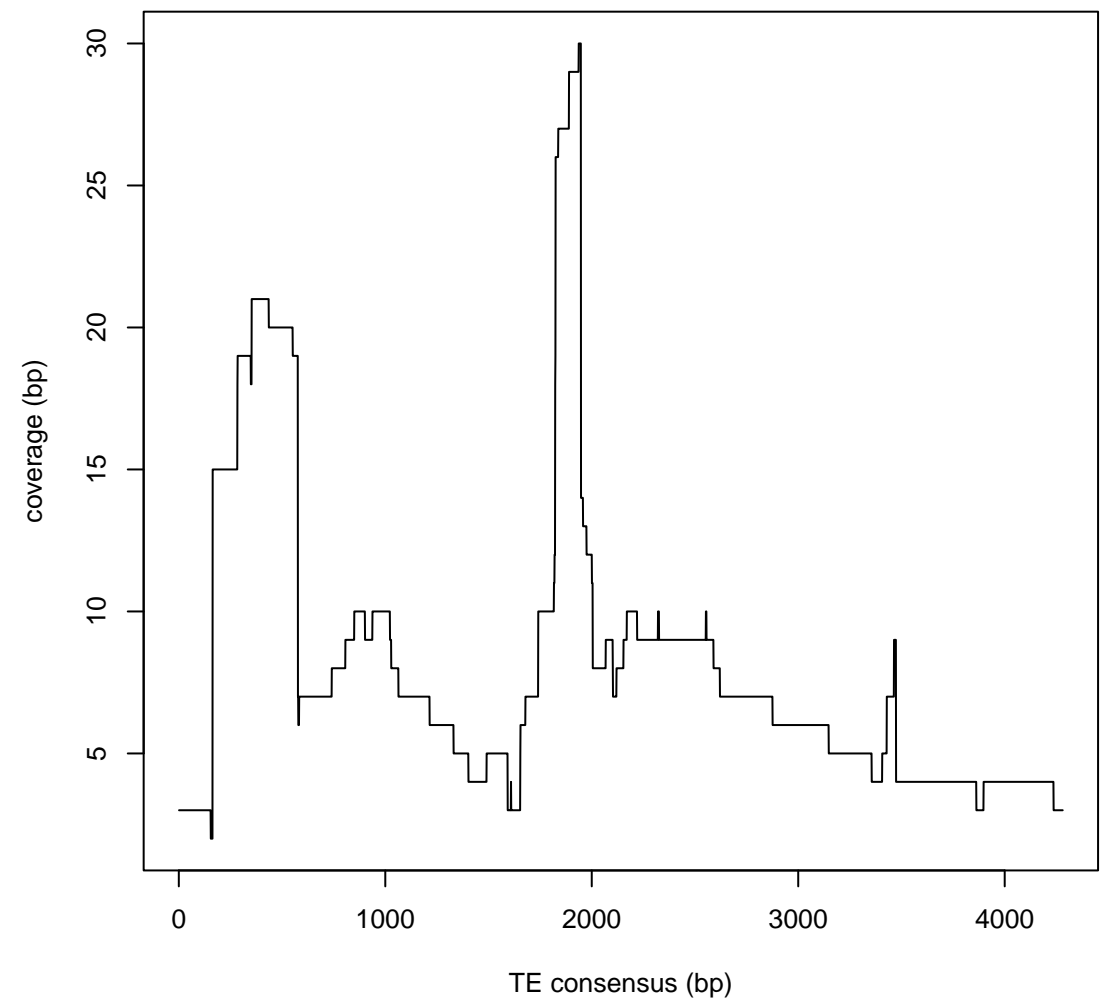


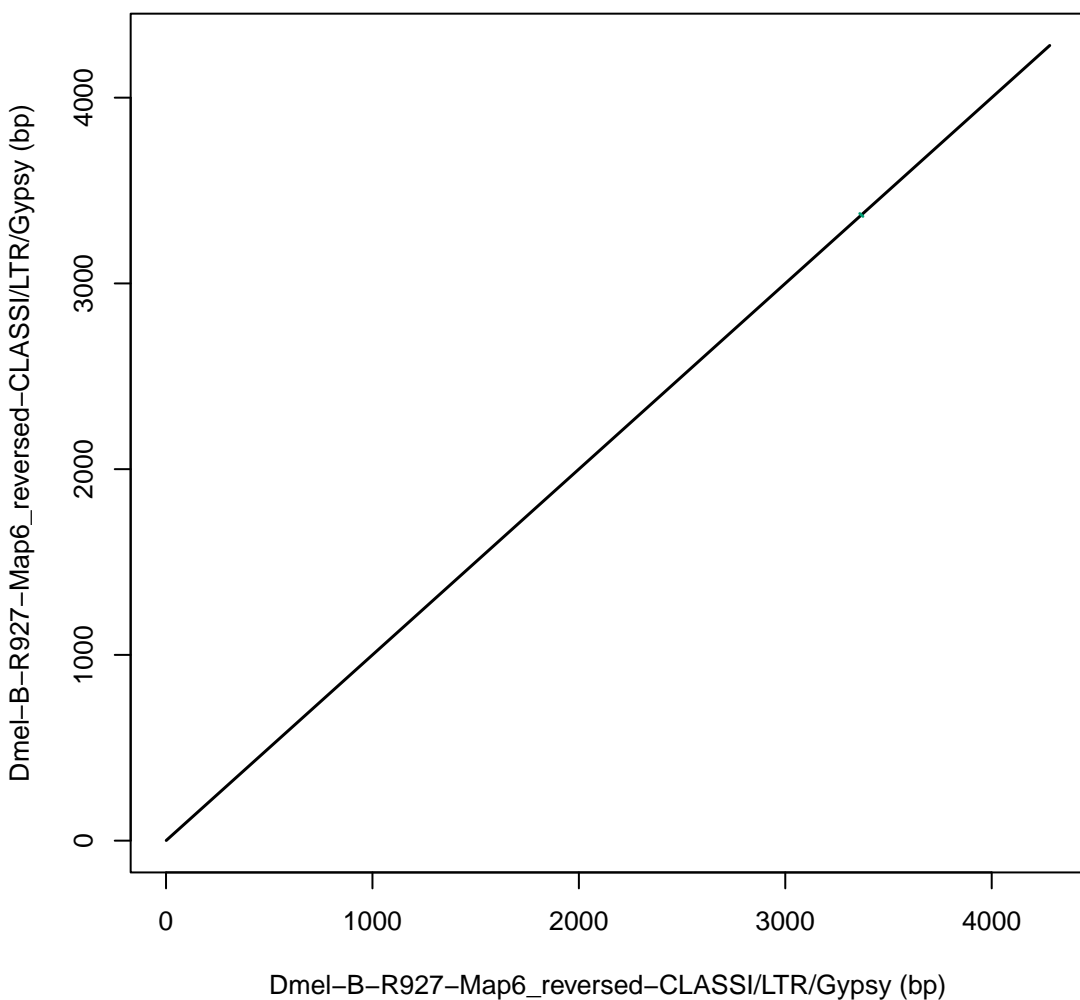
TE: Dmel-B-R927-Map6_reversed-CLASSI/LTR/Gypsy
consensus size: 4281bp; fragments: 72; full length: 1 (>=3852.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

