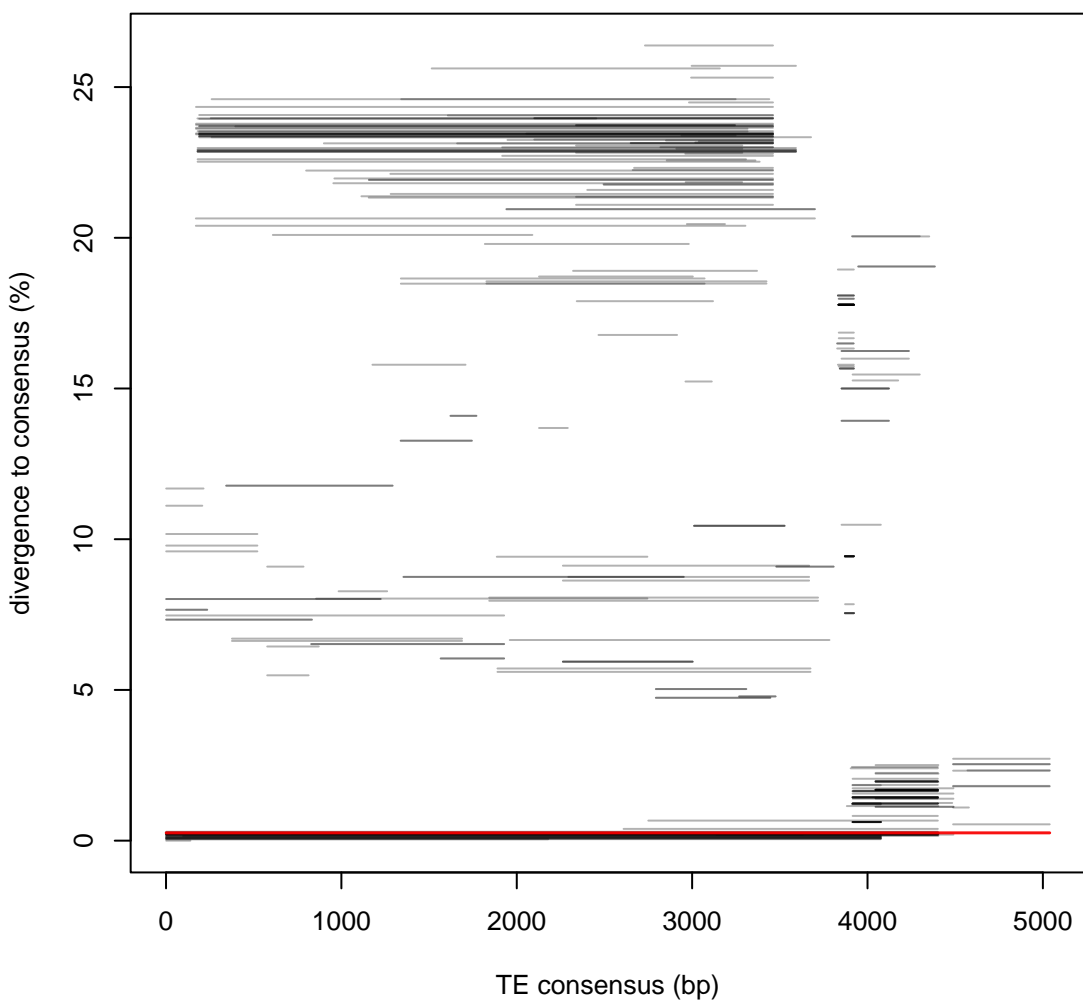
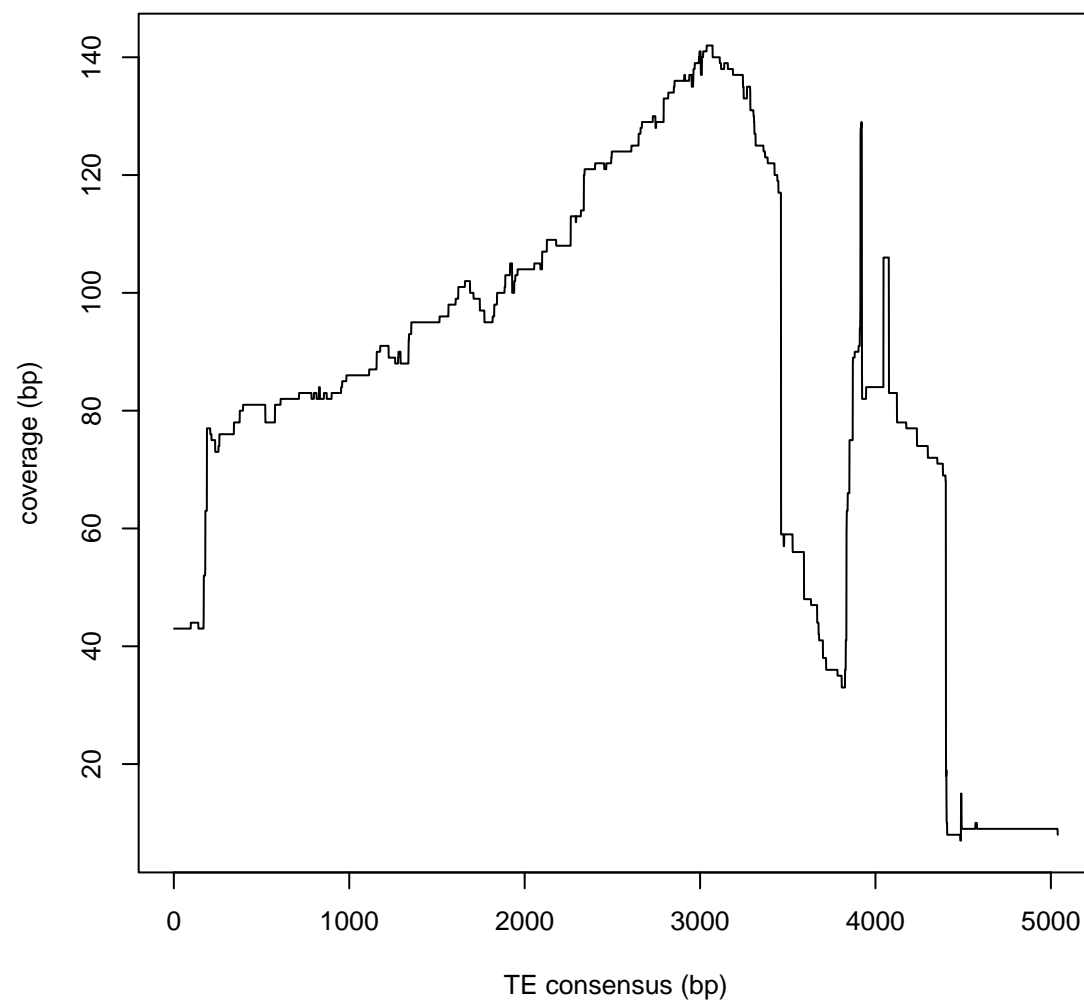


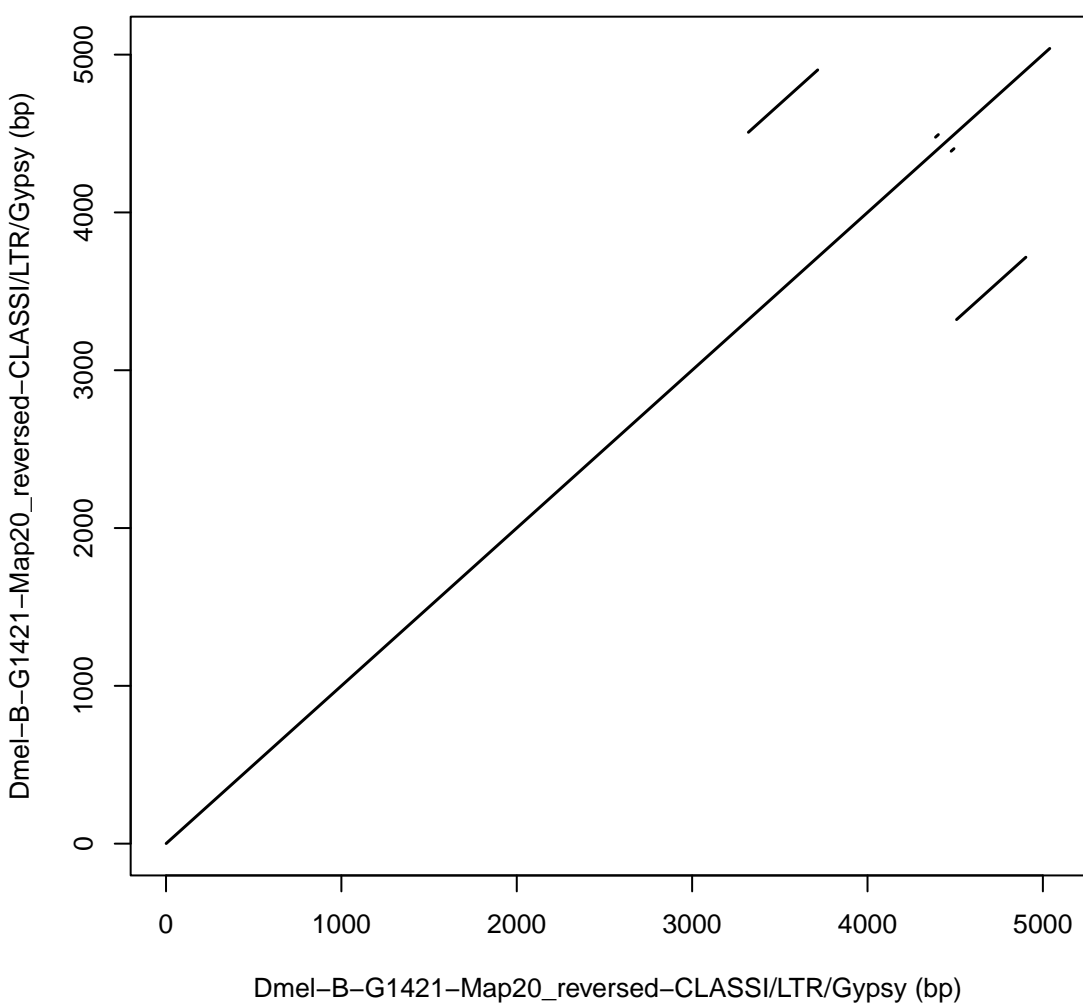
TE: Dmel-B-G1421-Map20_reversed-CLASSI/LTR/Gypsy
consensus size: 5039bp; fragments: 321; full length: 1 (>=4535.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

