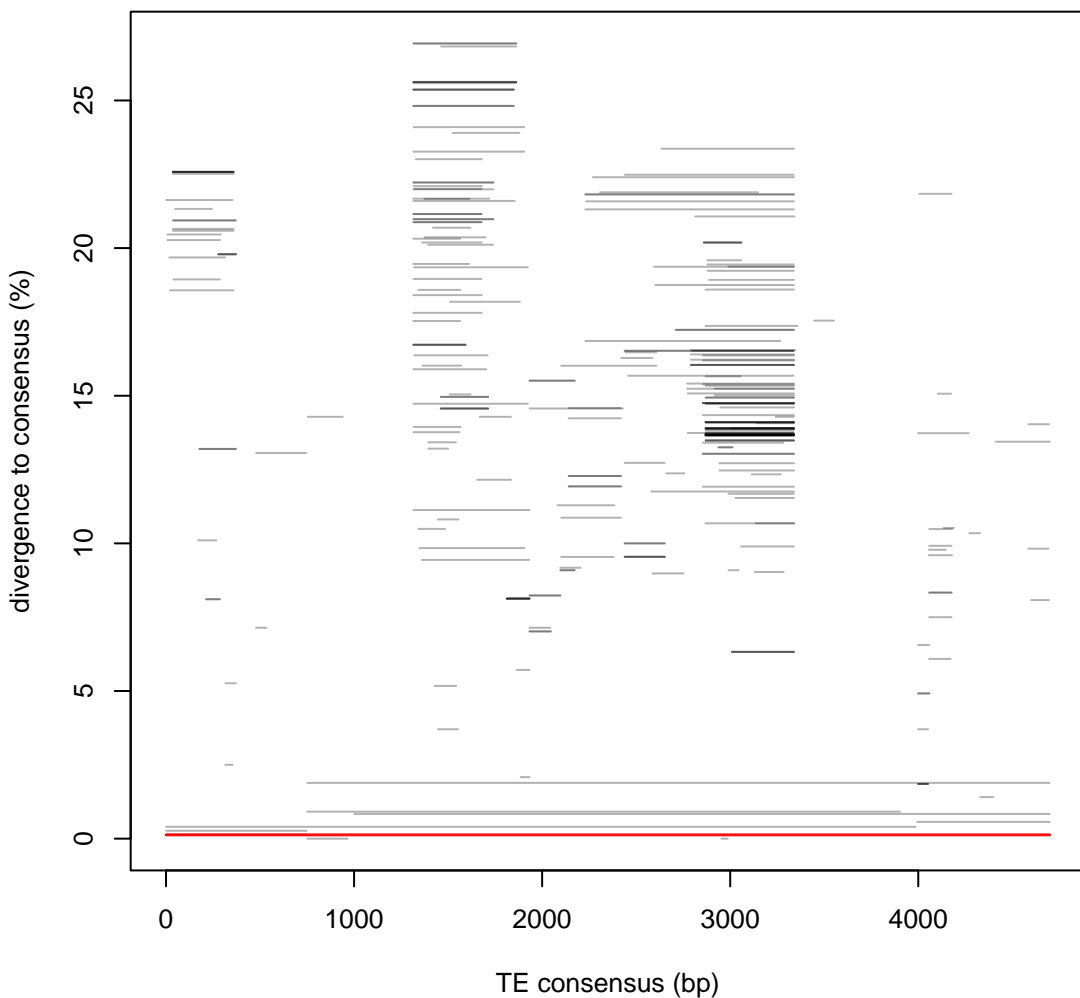
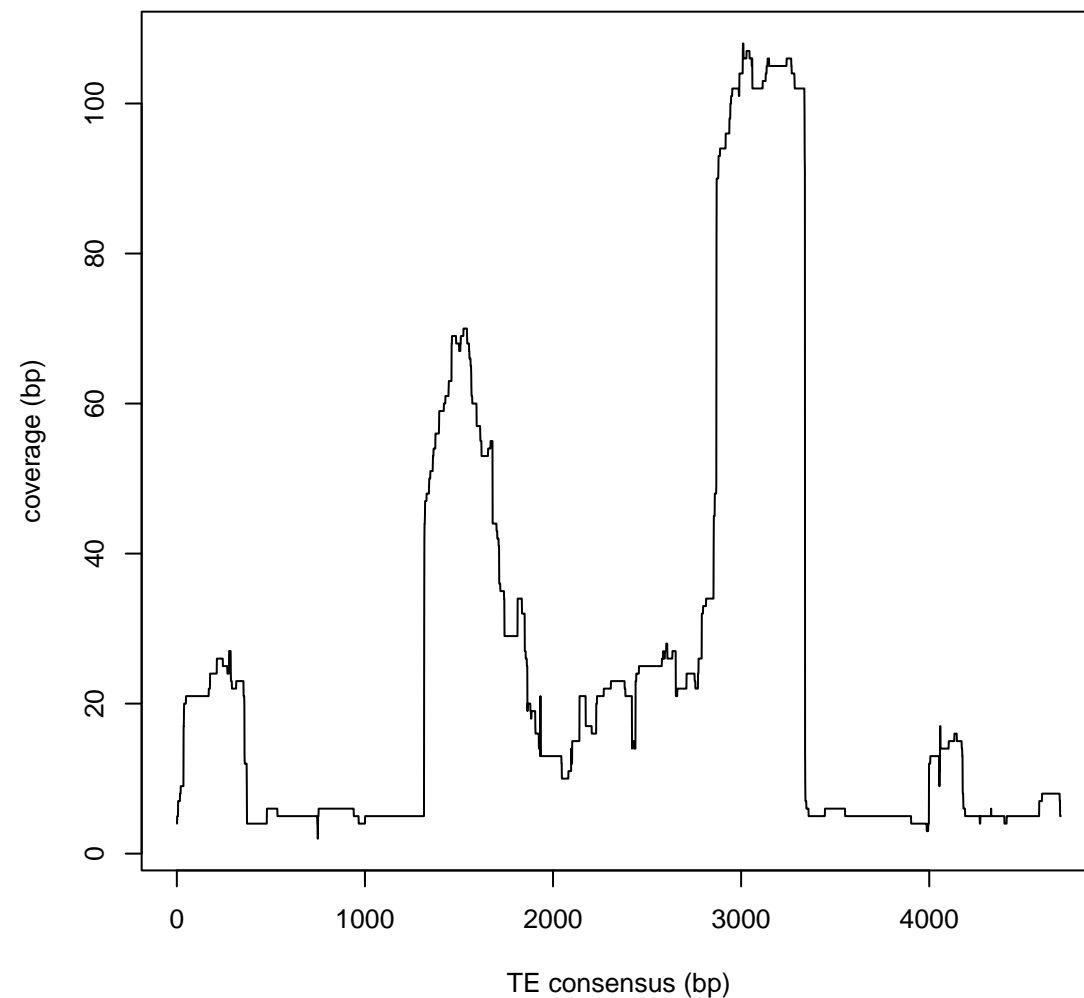


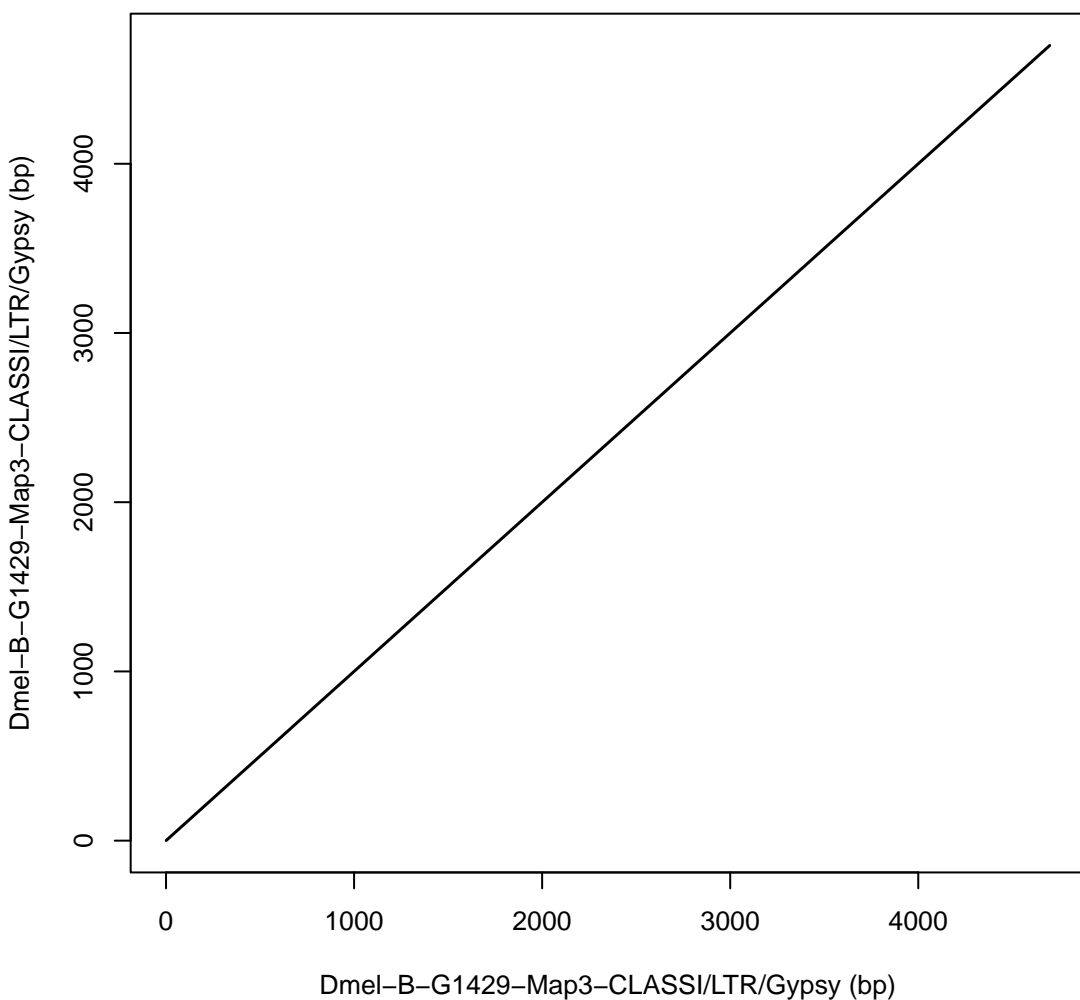
TE: Dmel-B-G1429-Map3-CLASSI/LTR/Gypsy
consensus size: 4699bp; fragments: 283; full length: 1 (>=4229.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

