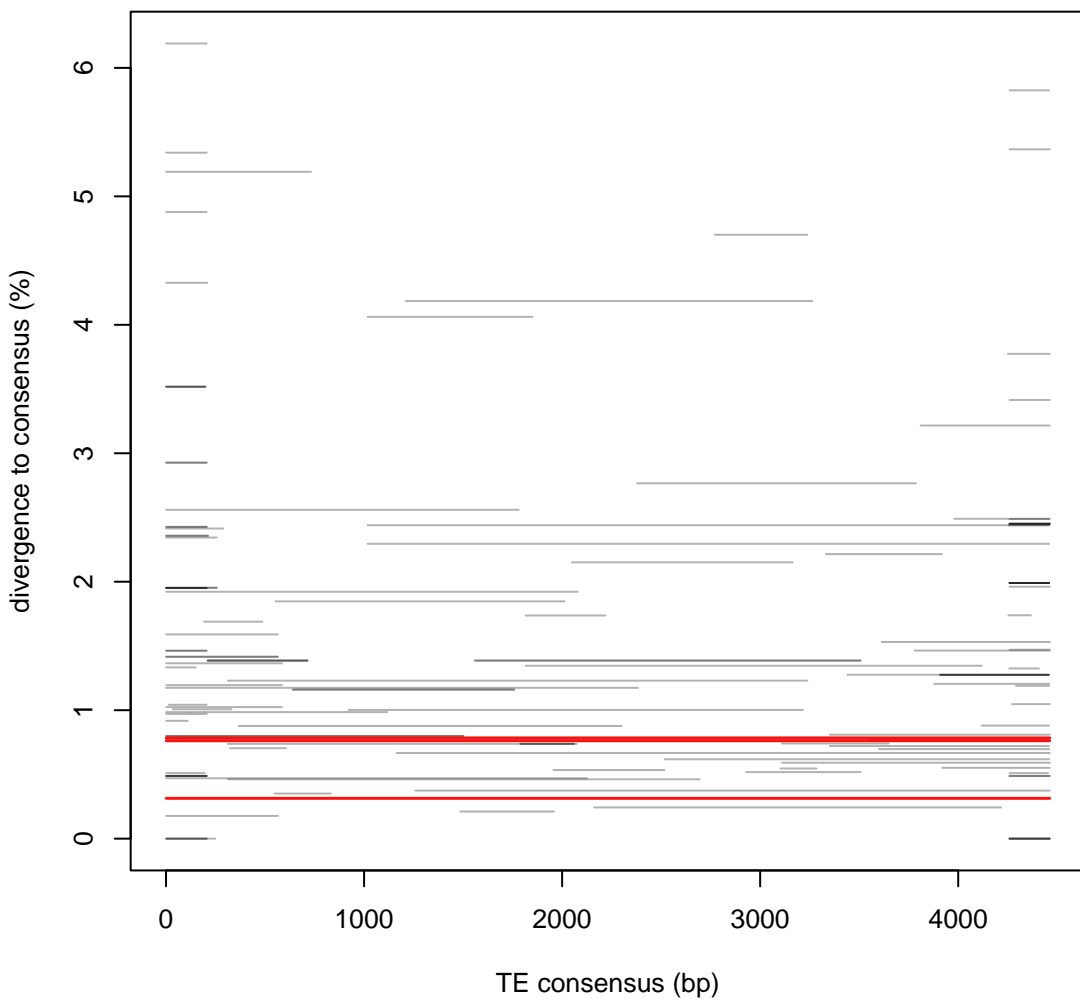
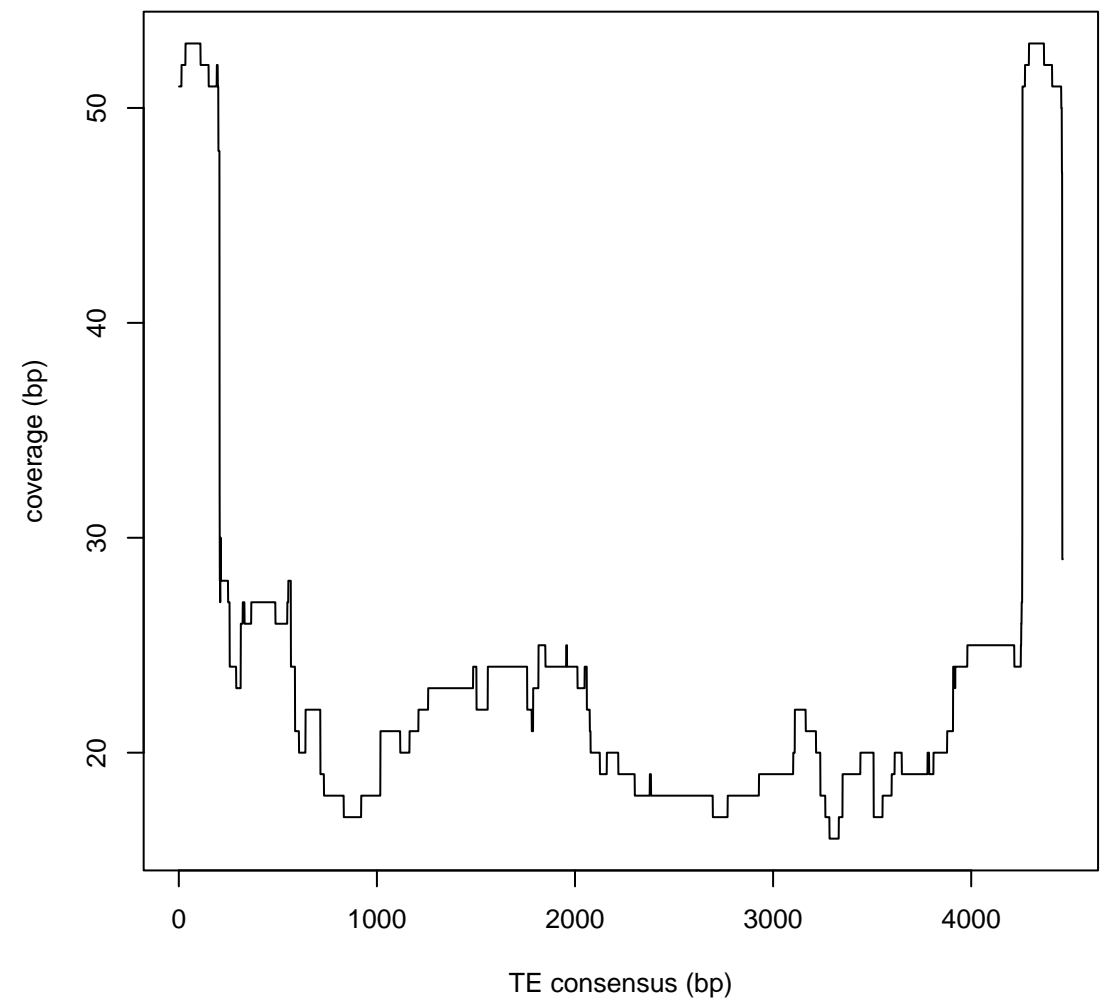


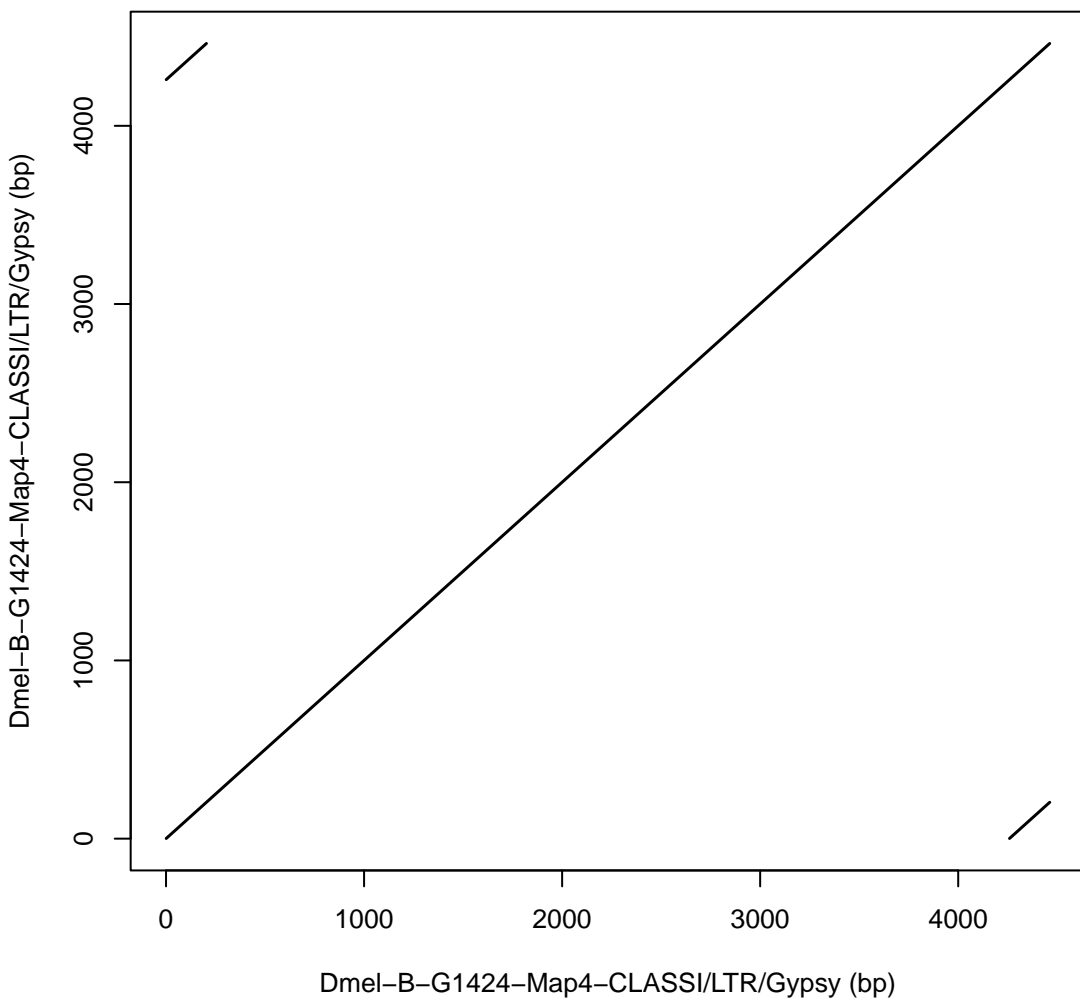
TE: Dmel-B-G1424-Map4-CLASSI/LTR/Gypsy
consensus size: 4462bp; fragments: 135; full length: 3 (>=4015.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

