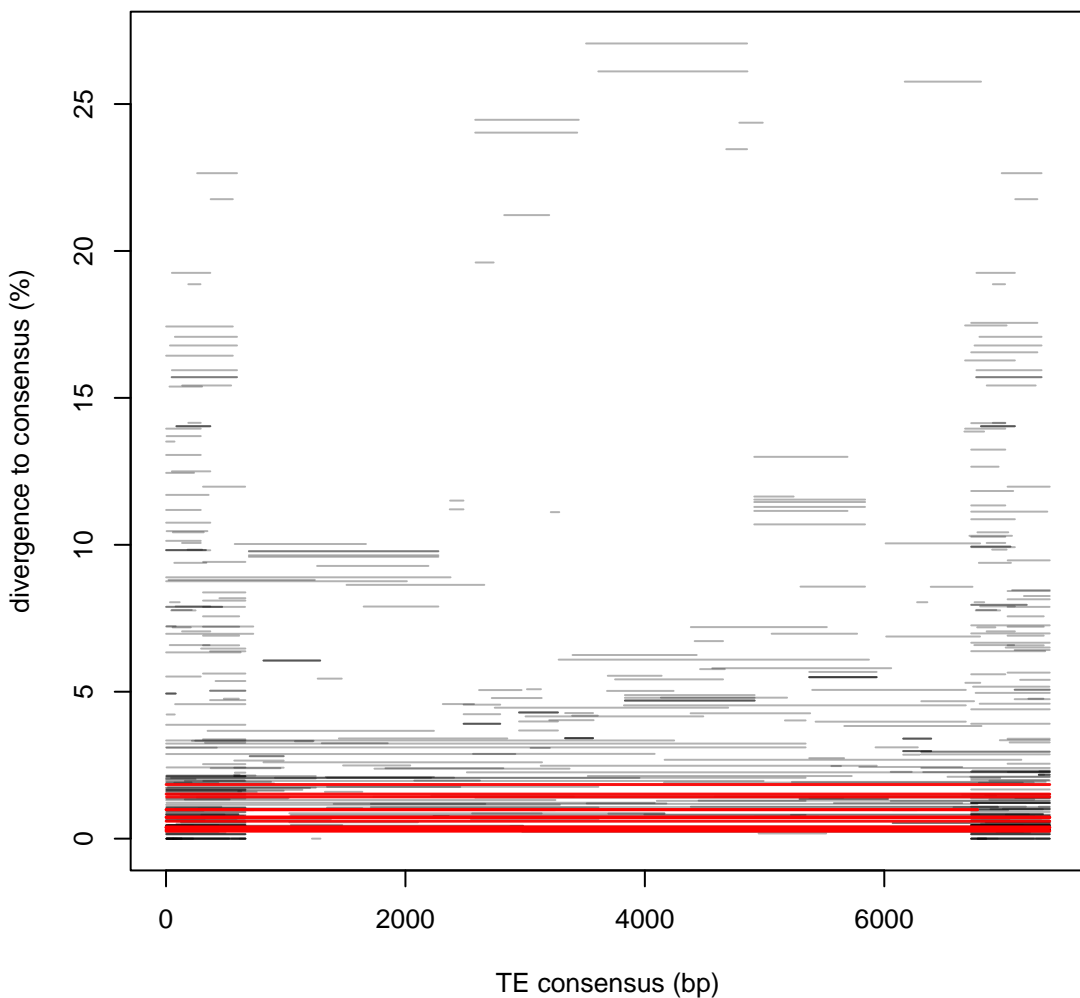
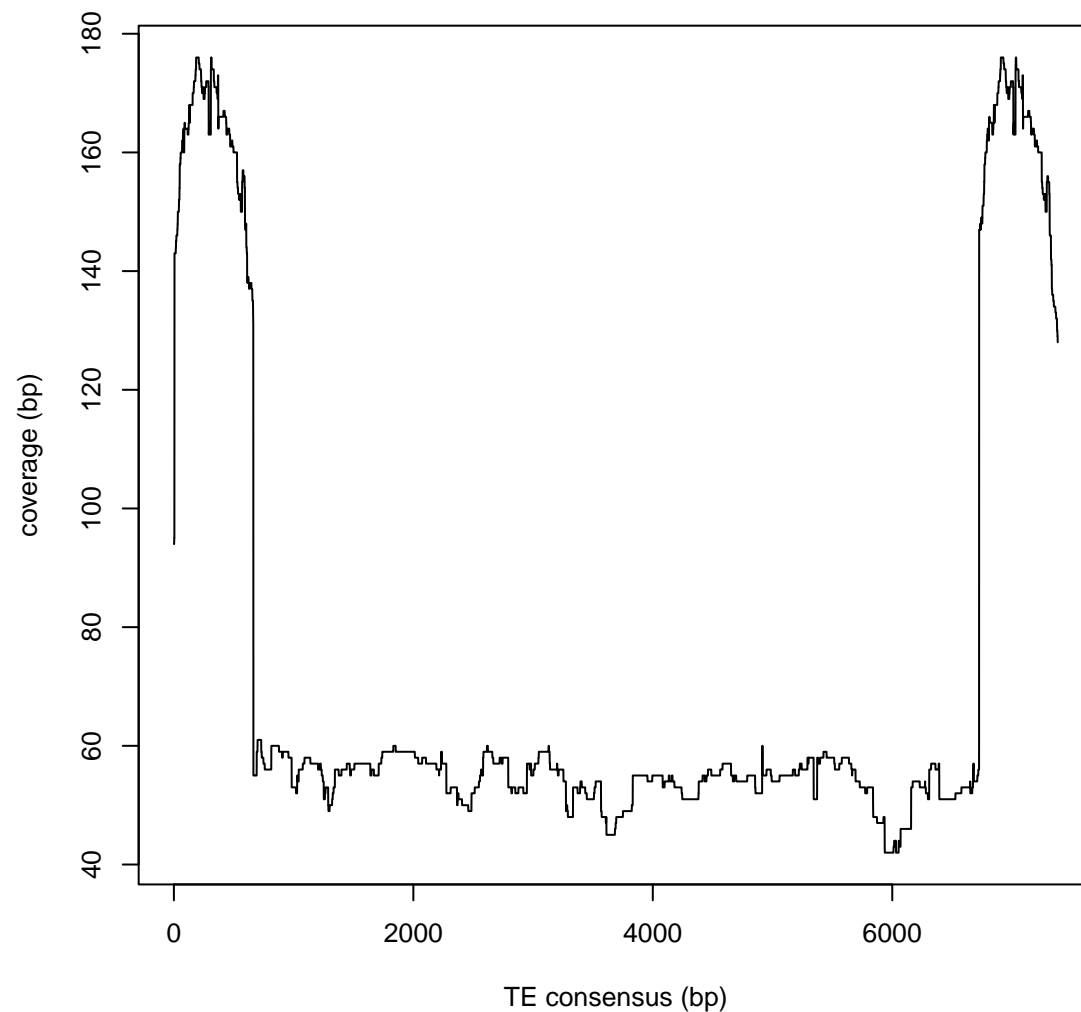


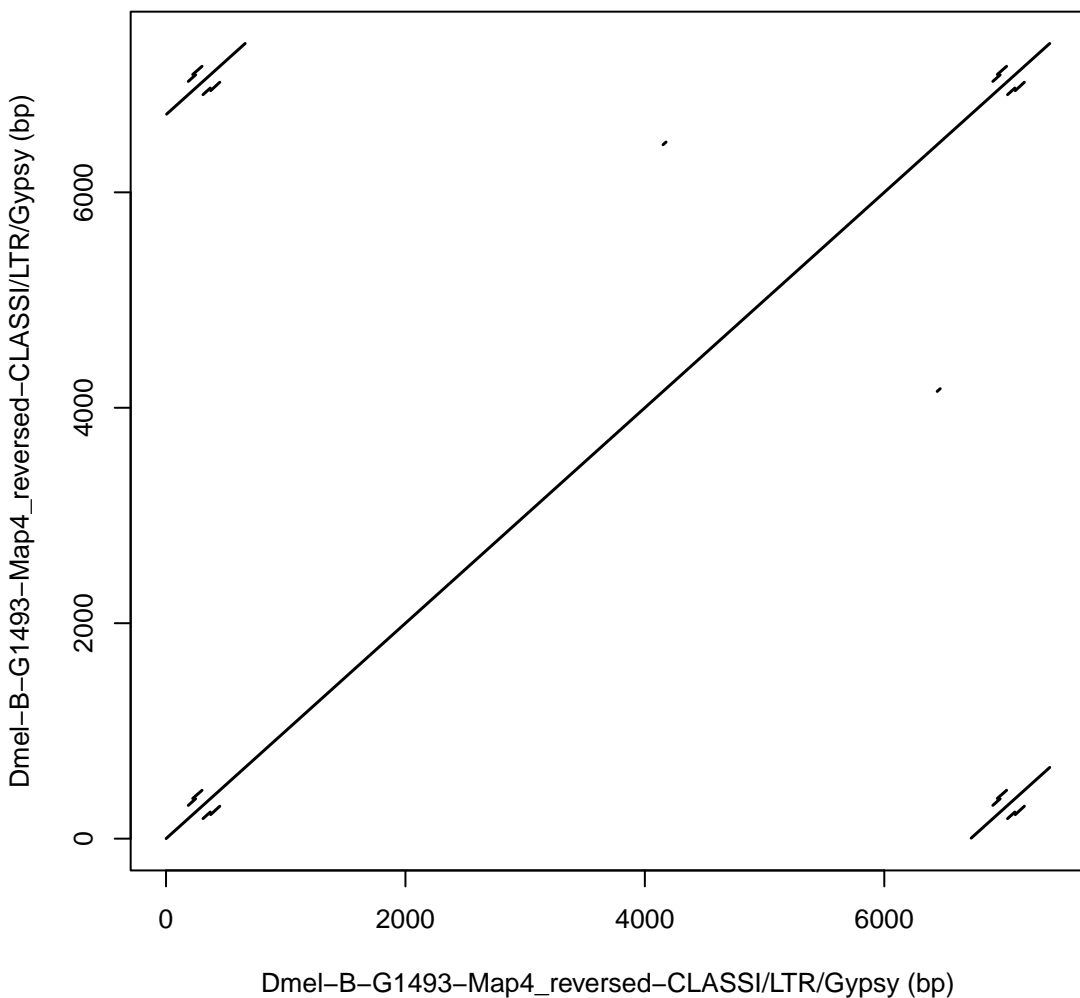
TE: Dmel-B-G1493-Map4_reversed-CLASSI/LTR/Gypsy
consensus size: 7382bp; fragments: 583; full length: 16 (>=6643.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

