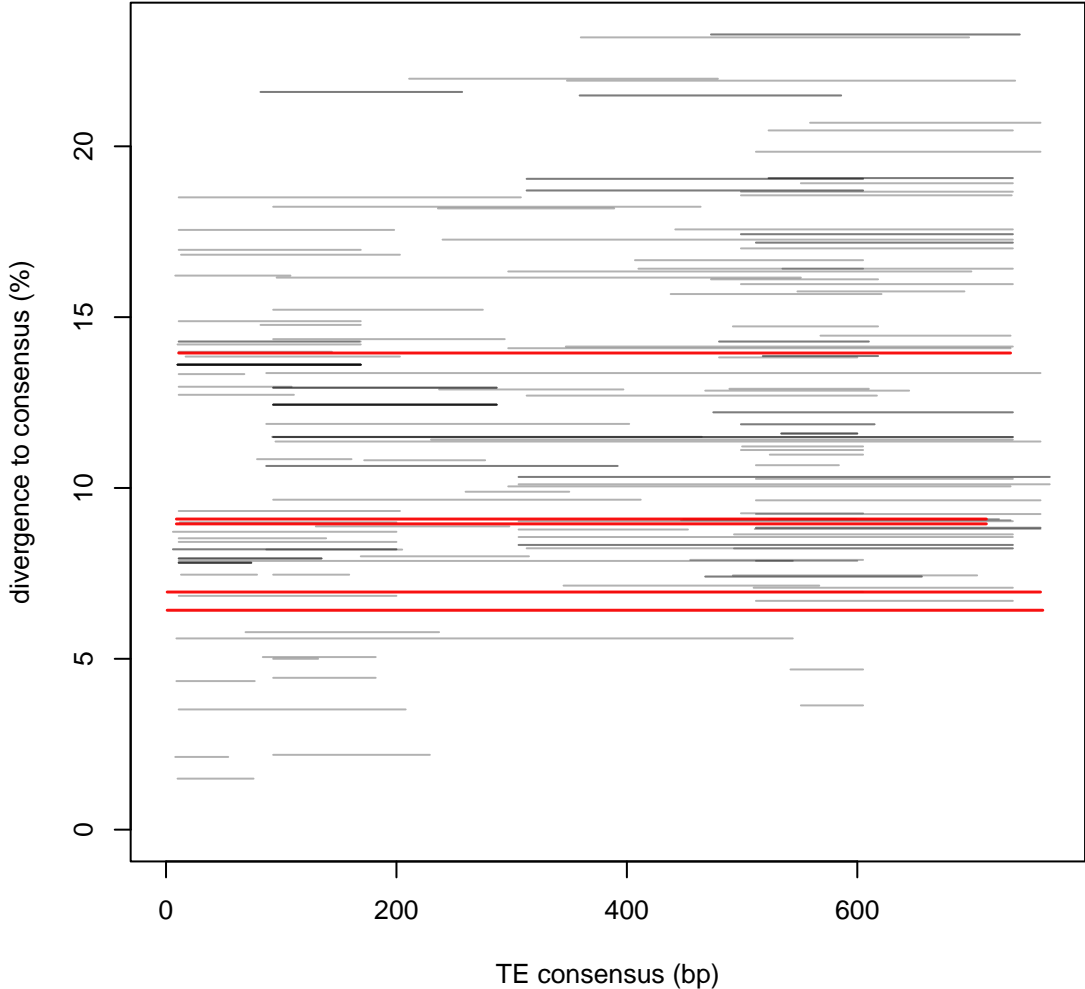
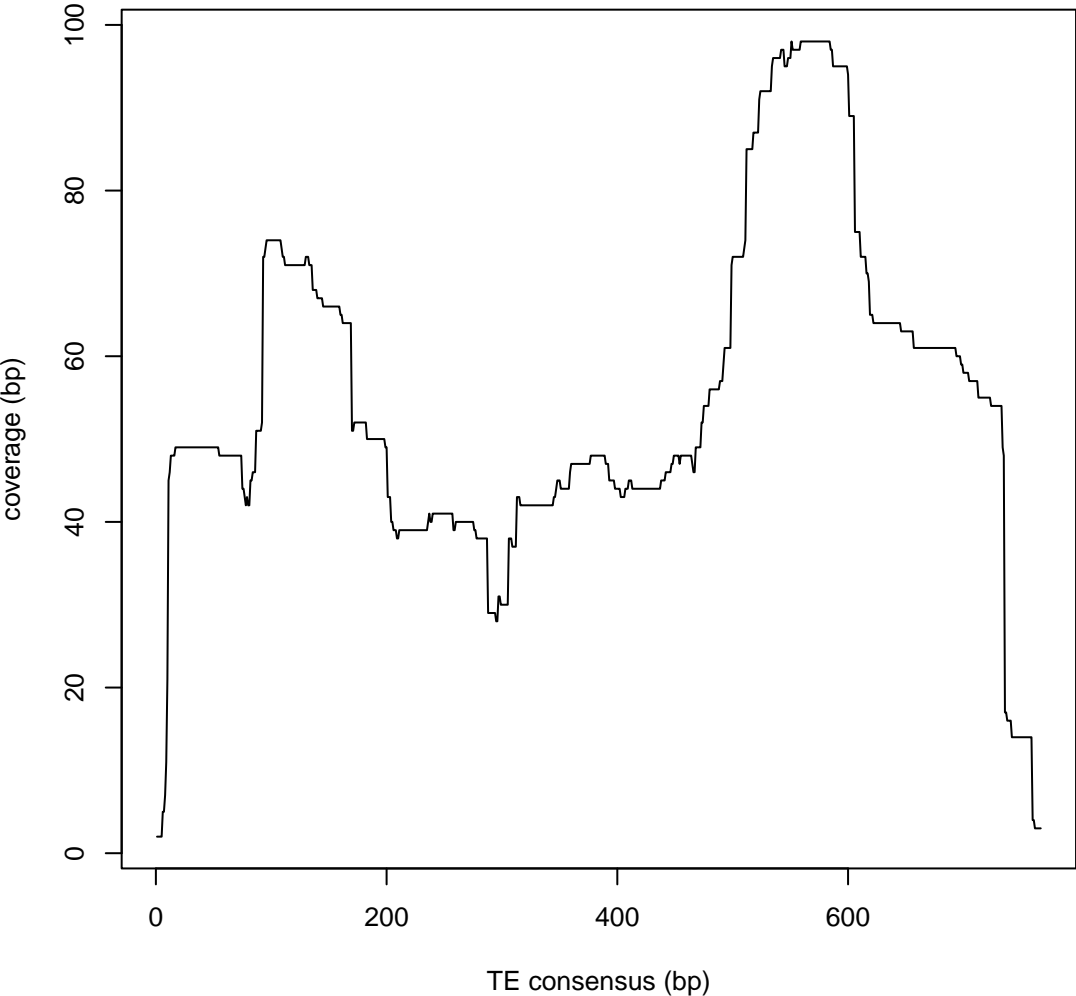


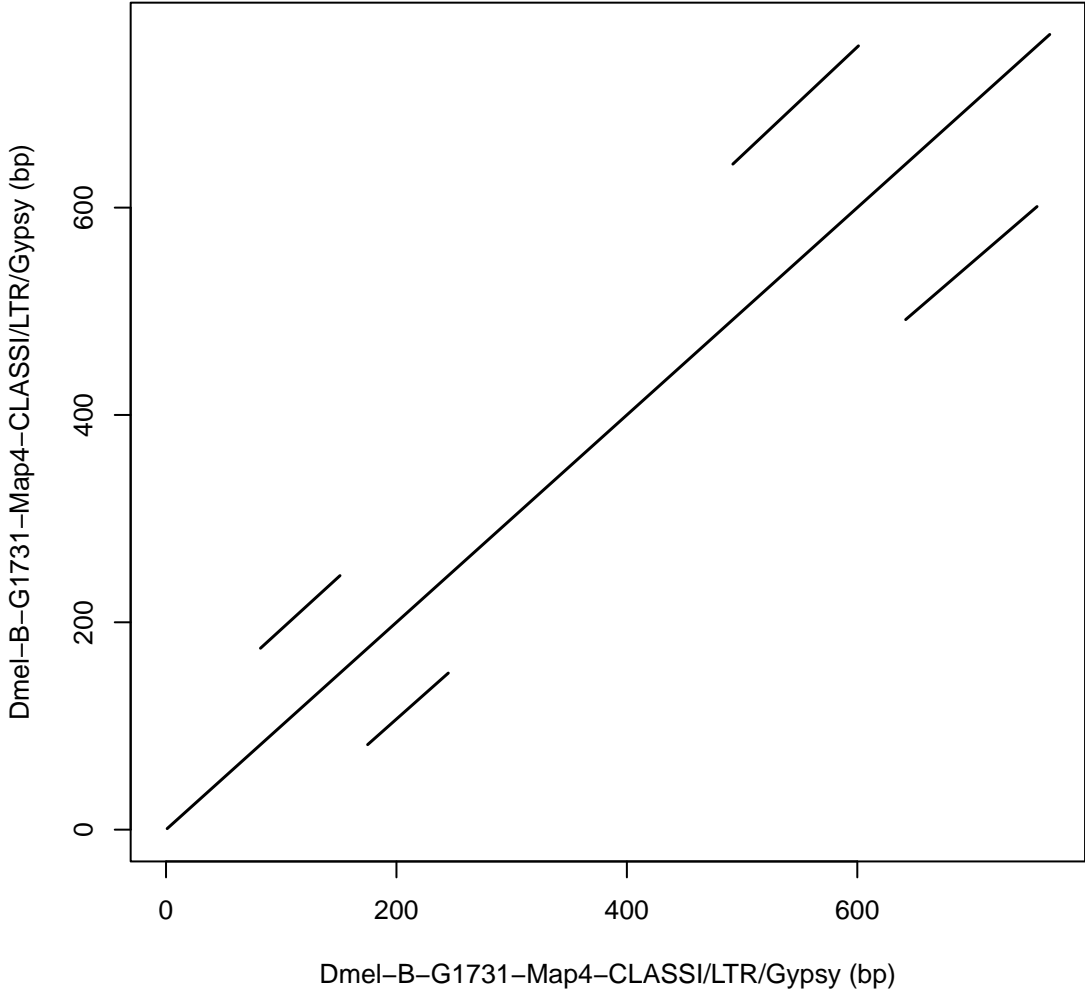
TE: Dmel-B-G1731-Map4-CLASSI/LTR/Gypsy  
consensus size: 767bp; fragments: 180; full length: 5 (>=690.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

