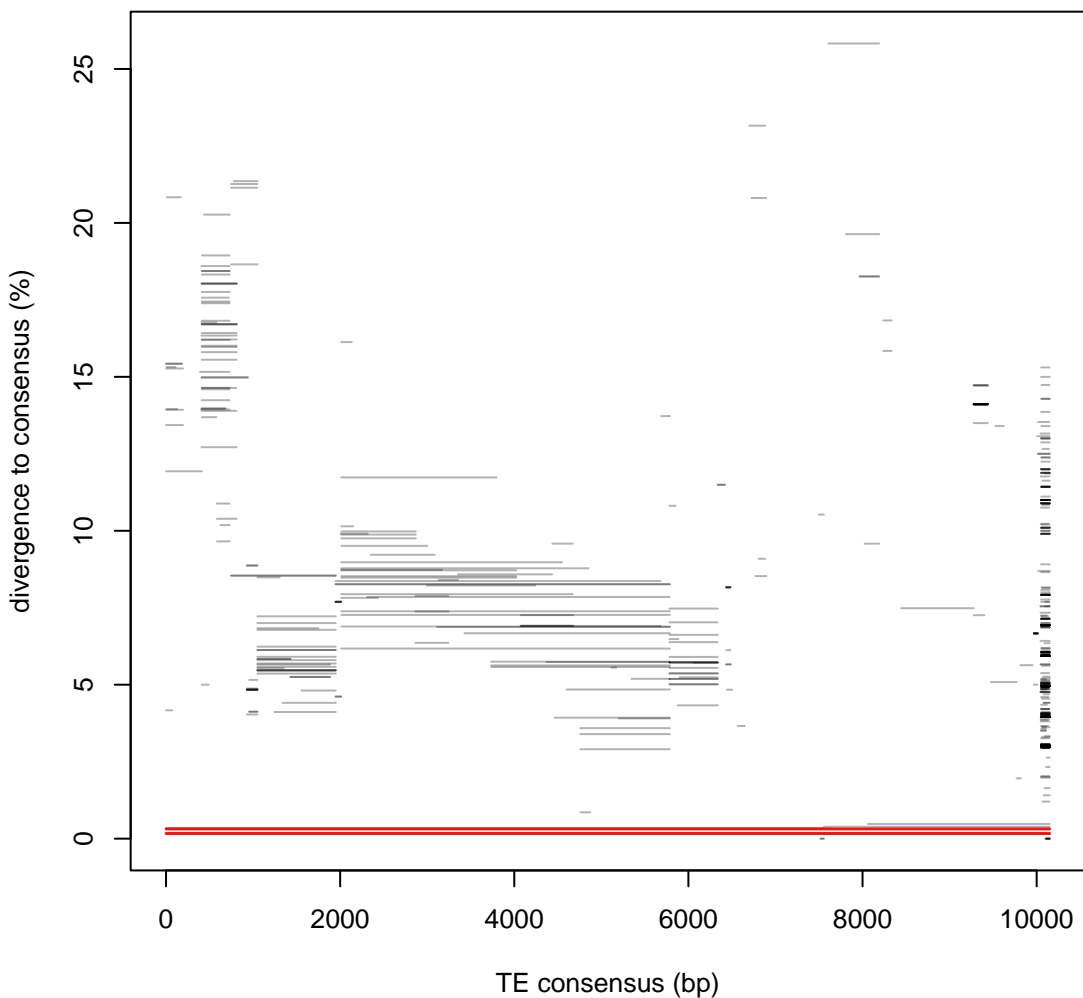
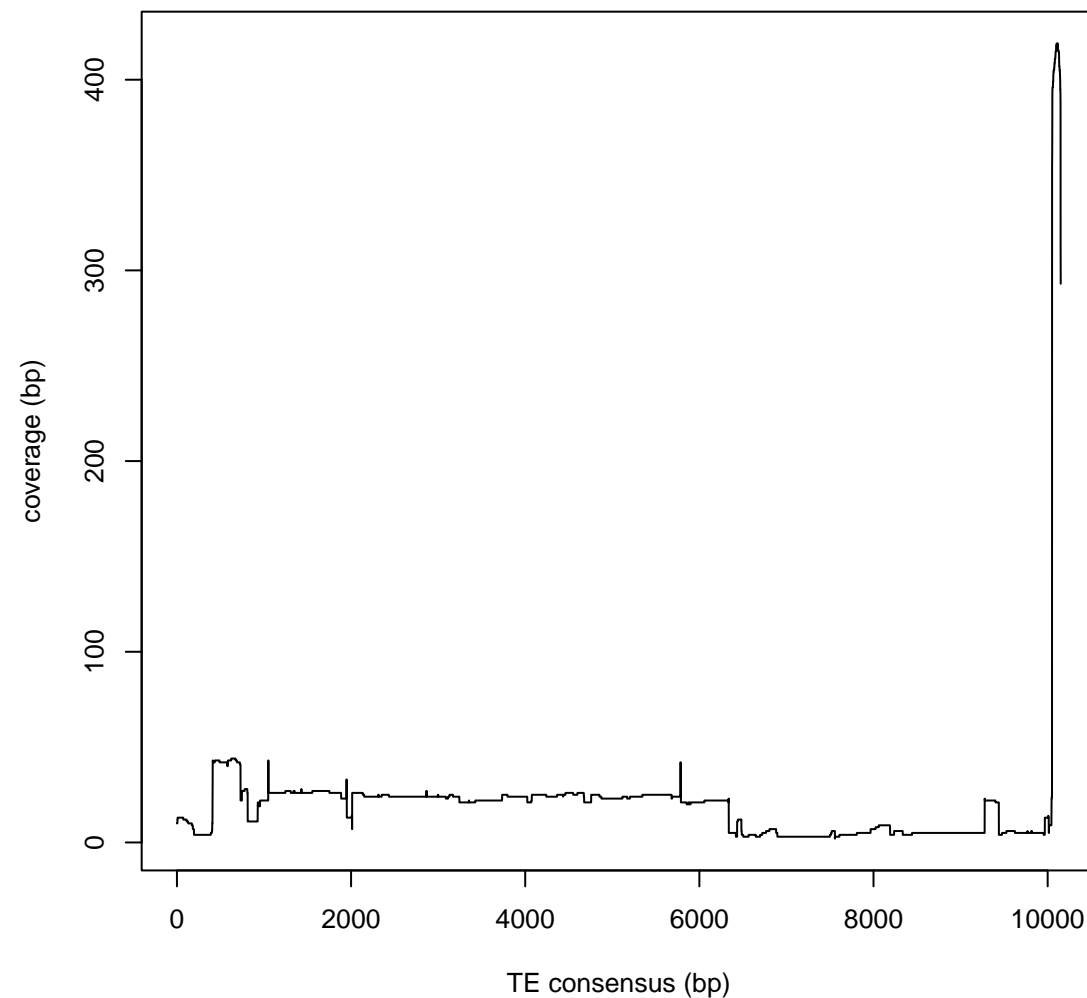


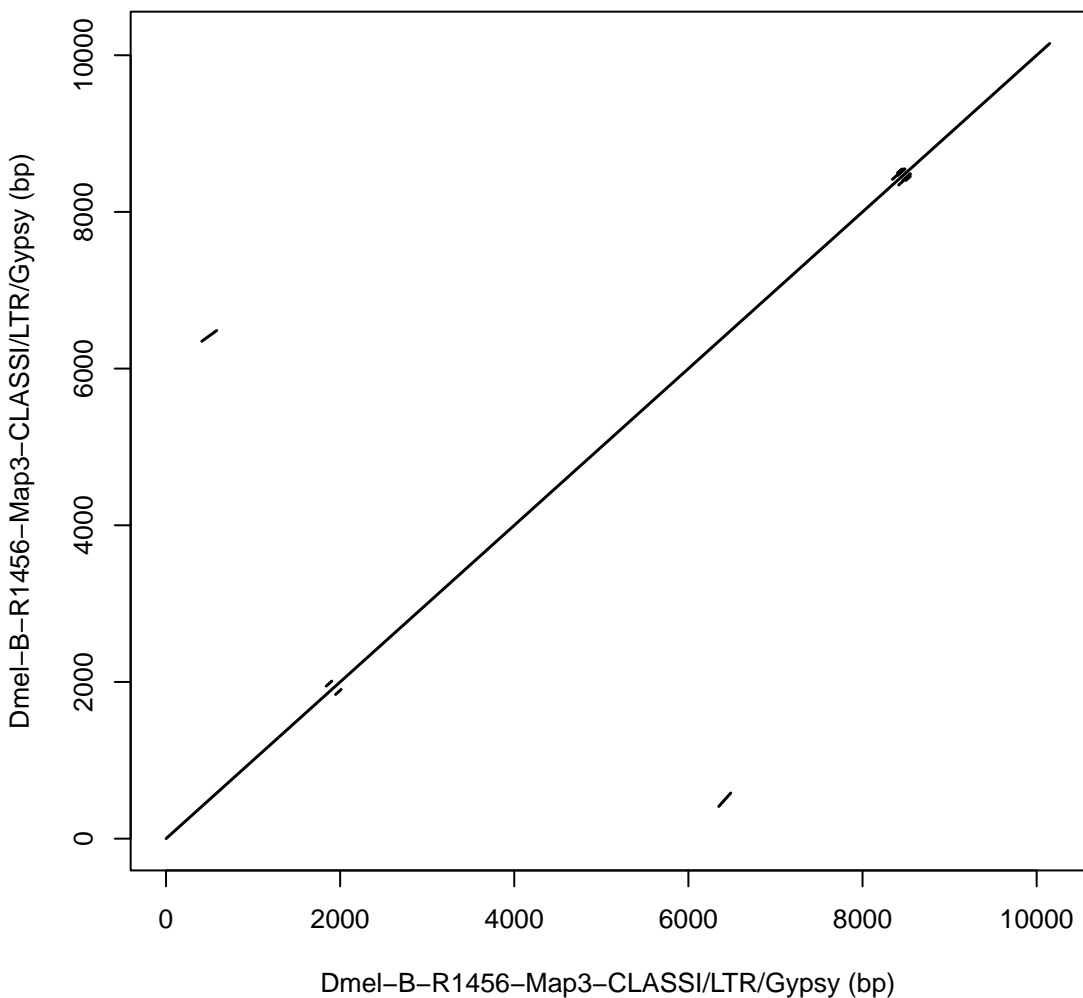
TE: Dmel-B-R1456-Map3-CLASSI/LTR/Gypsy  
consensus size: 10150bp; fragments: 663; full length: 2 (>=9135bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

