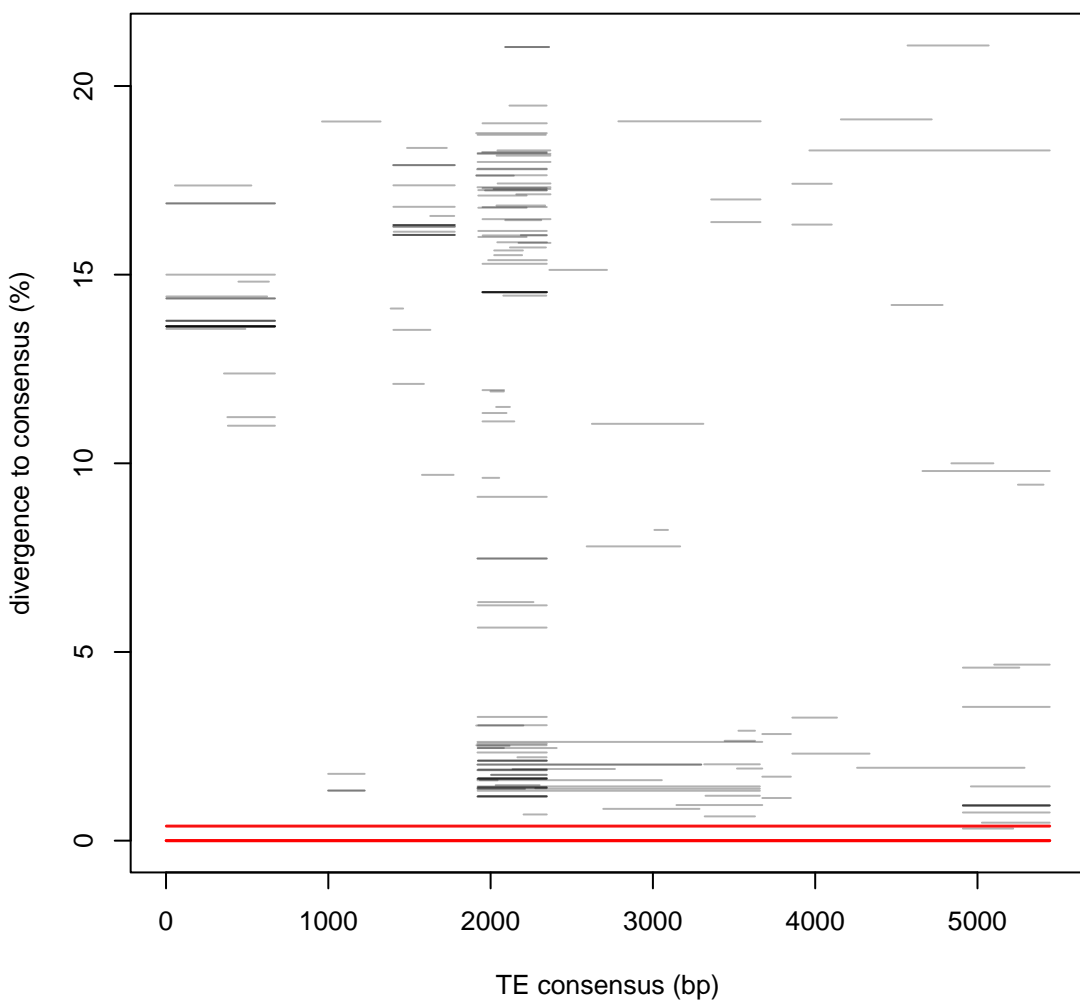
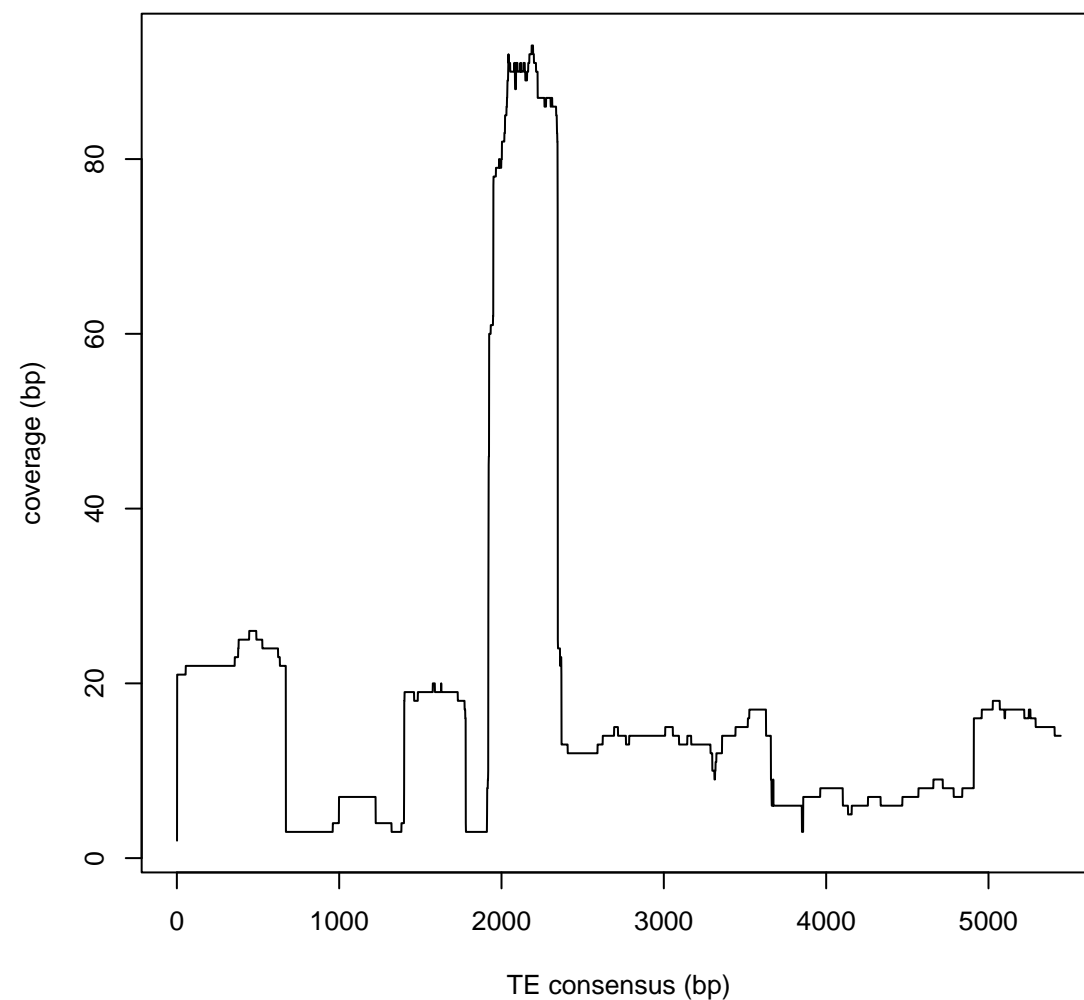


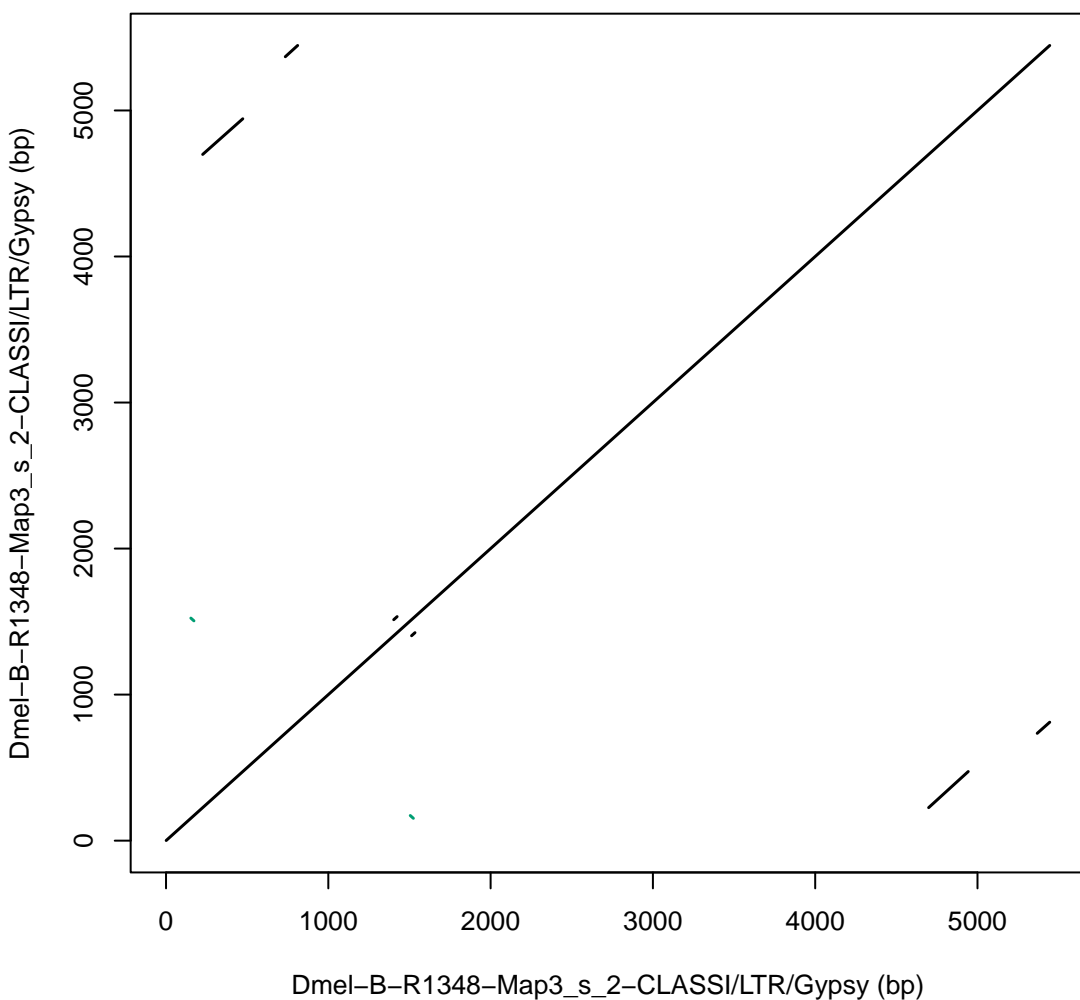
TE: Dmel-B-R1348-Map3\_s\_2-CLASSI/LTR/Gypsy  
consensus size: 5445bp; fragments: 194; full length: 3 (>=4900.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

