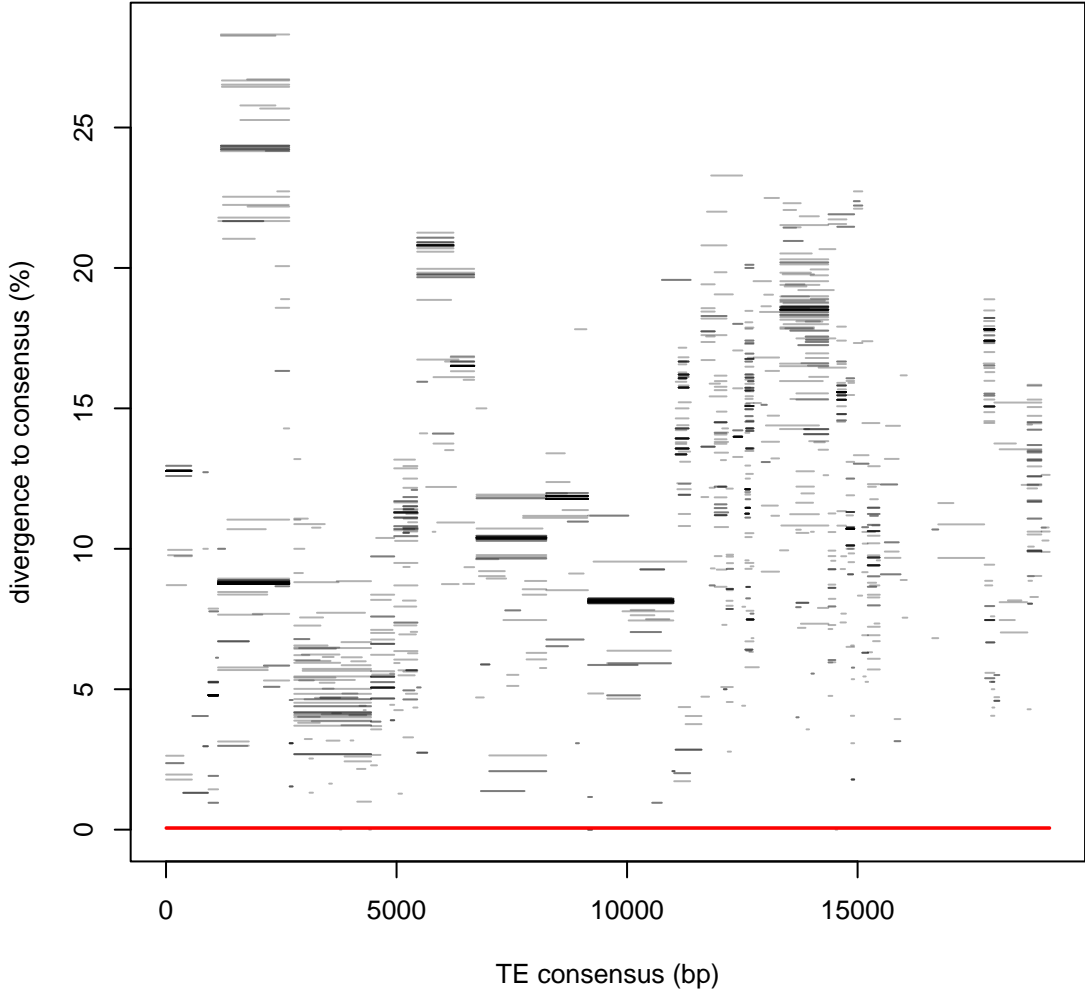
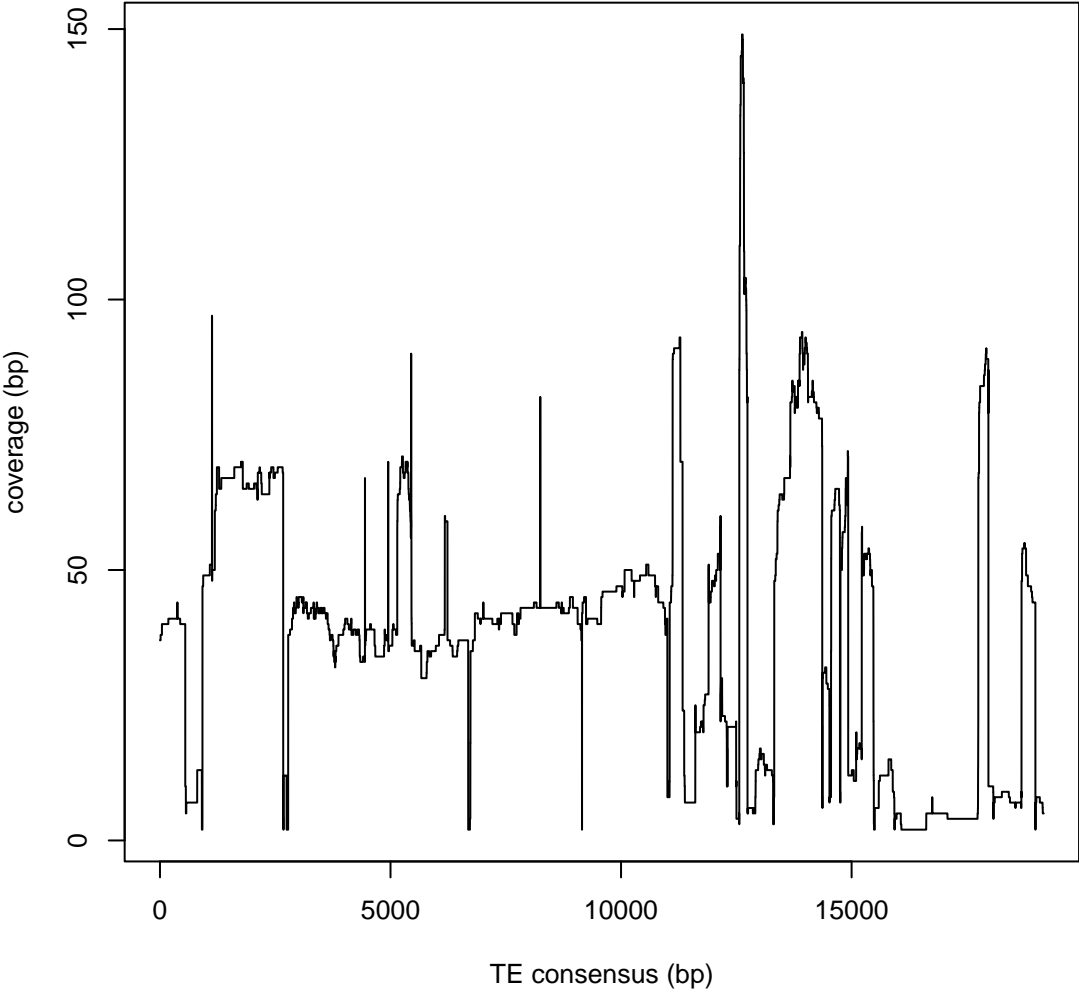


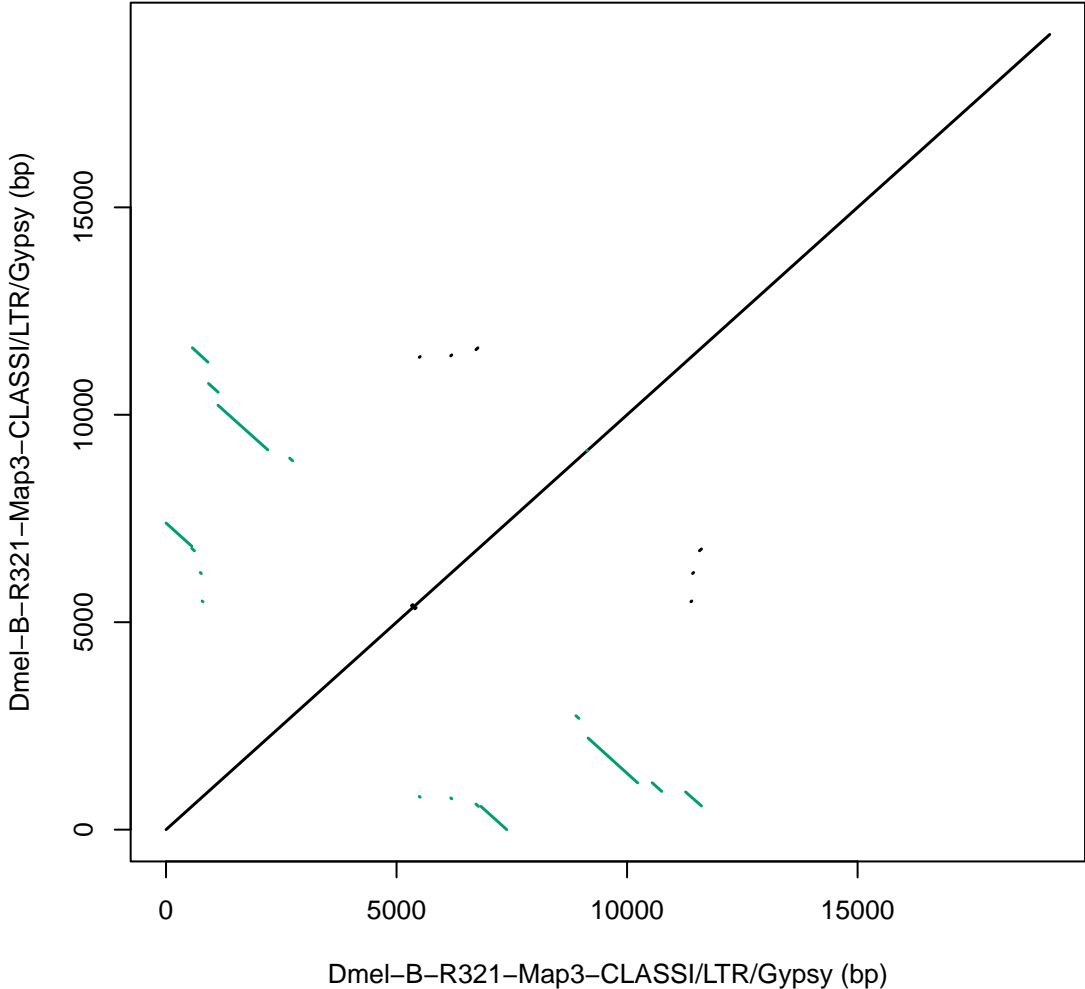
TE: Dmel-B-R321-Map3-CLASSI/LTR/Gypsy  
consensus size: 19167bp; fragments: 1530; full length: 2 (>=17250.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

