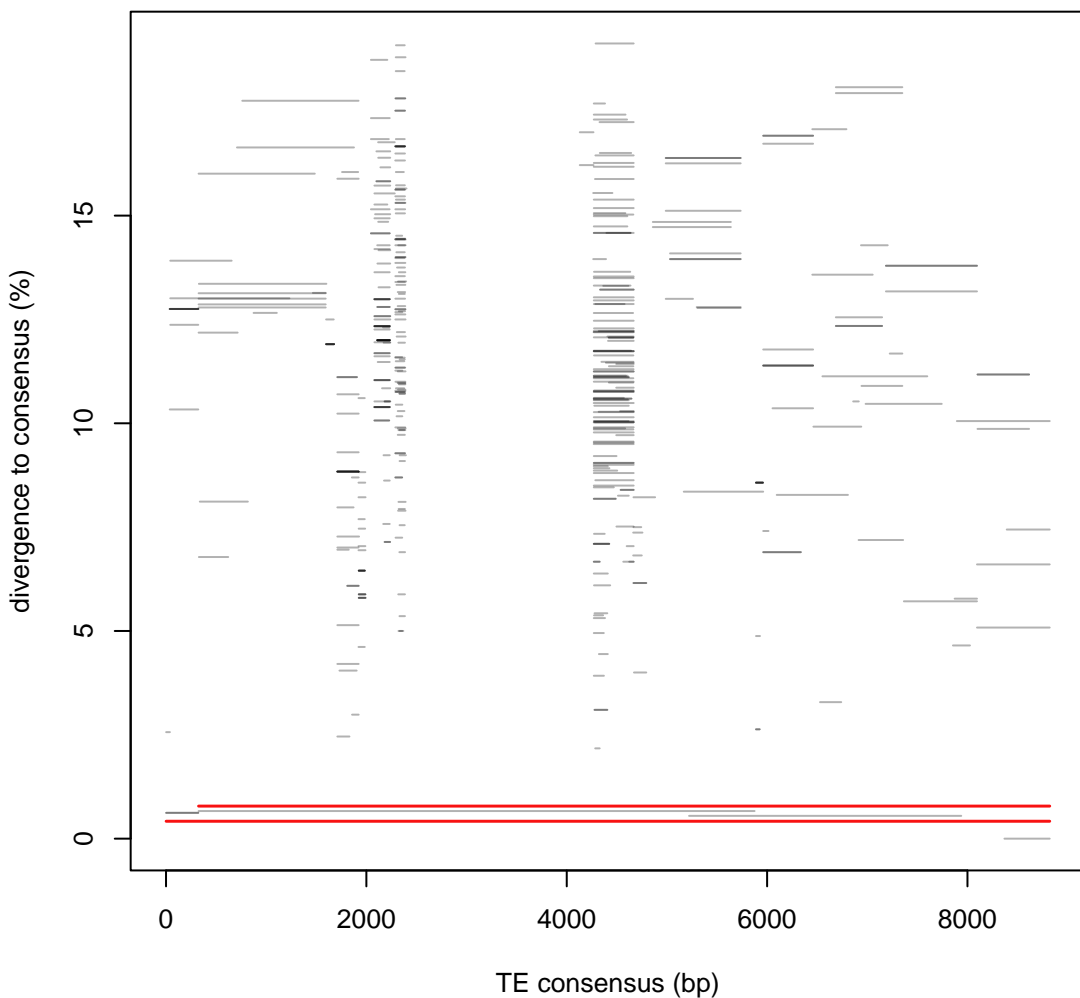
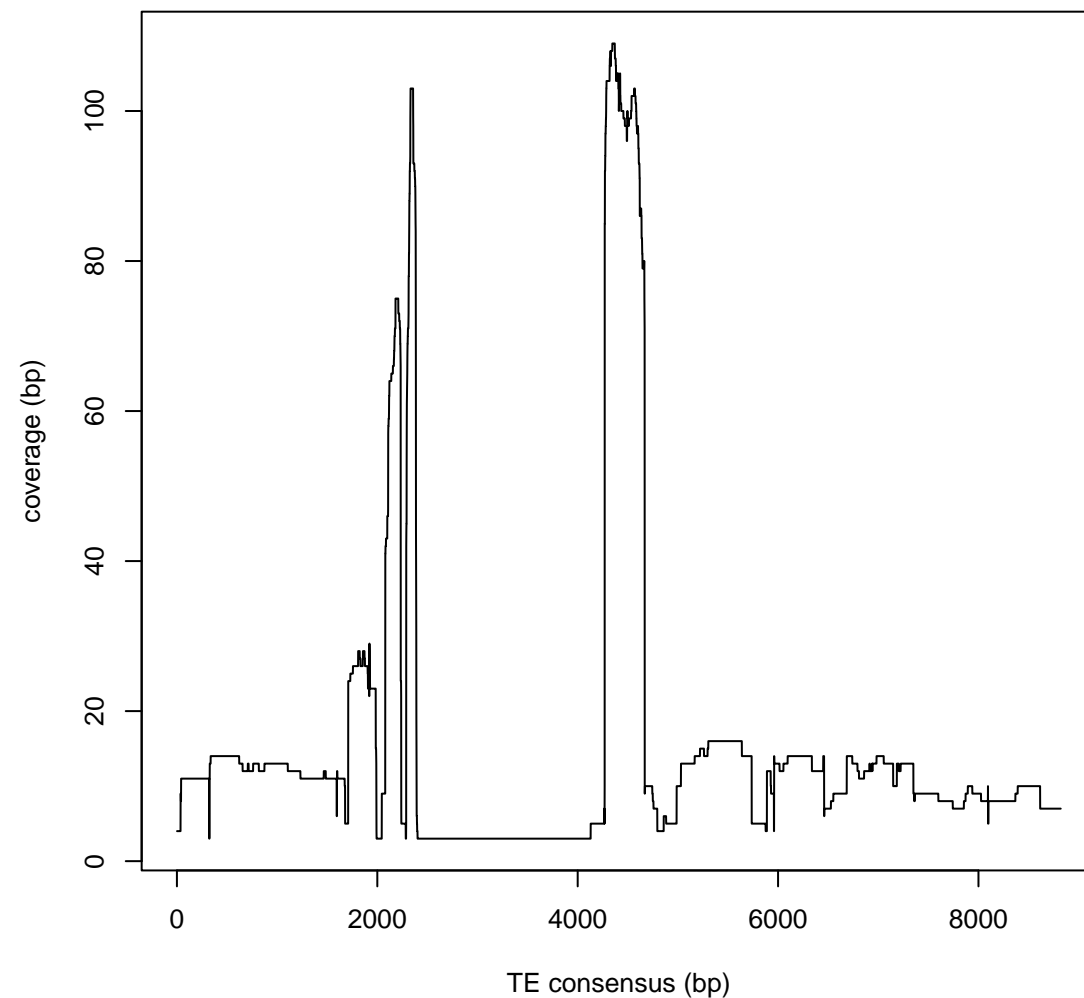


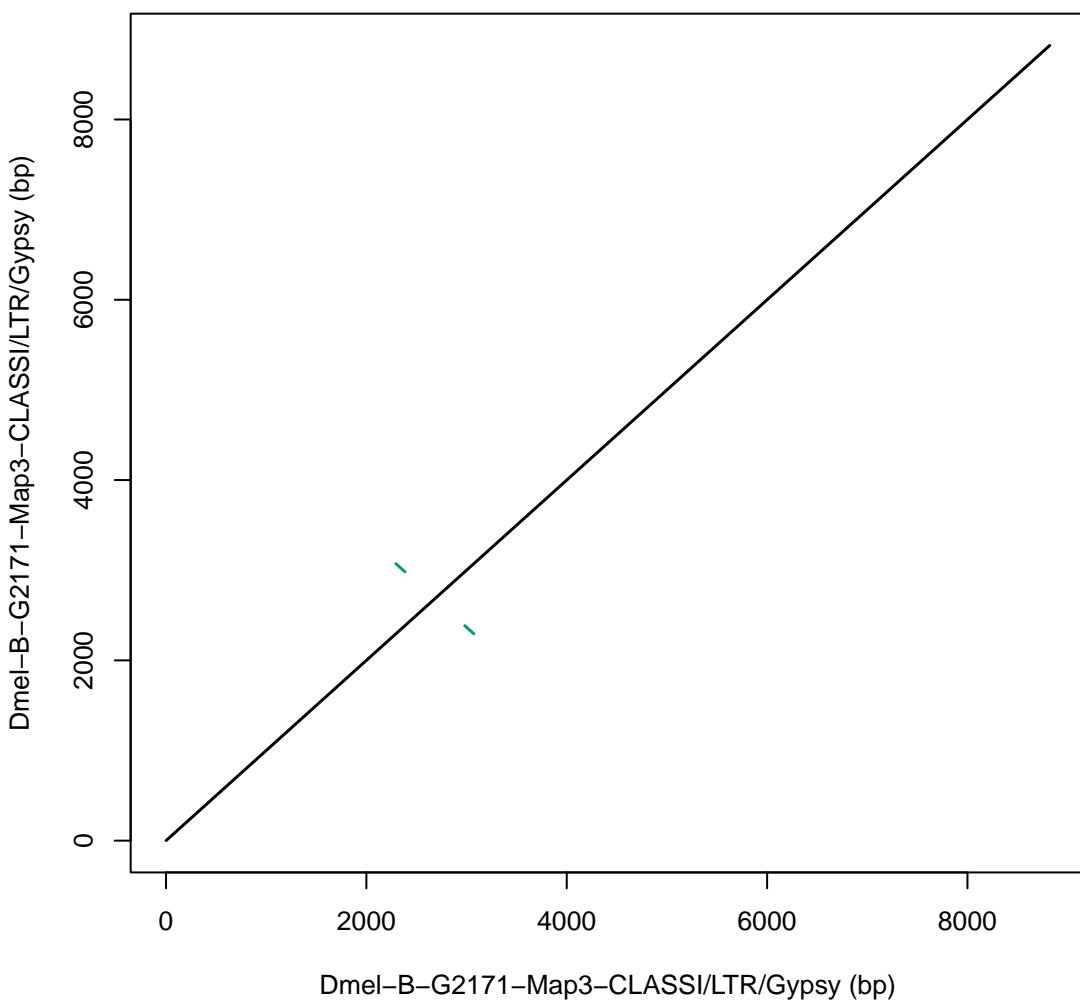
TE: Dmel-B-G2171-Map3-CLASSI/LTR/Gypsy  
consensus size: 8821bp; fragments: 452; full length: 2 (>=7938.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

