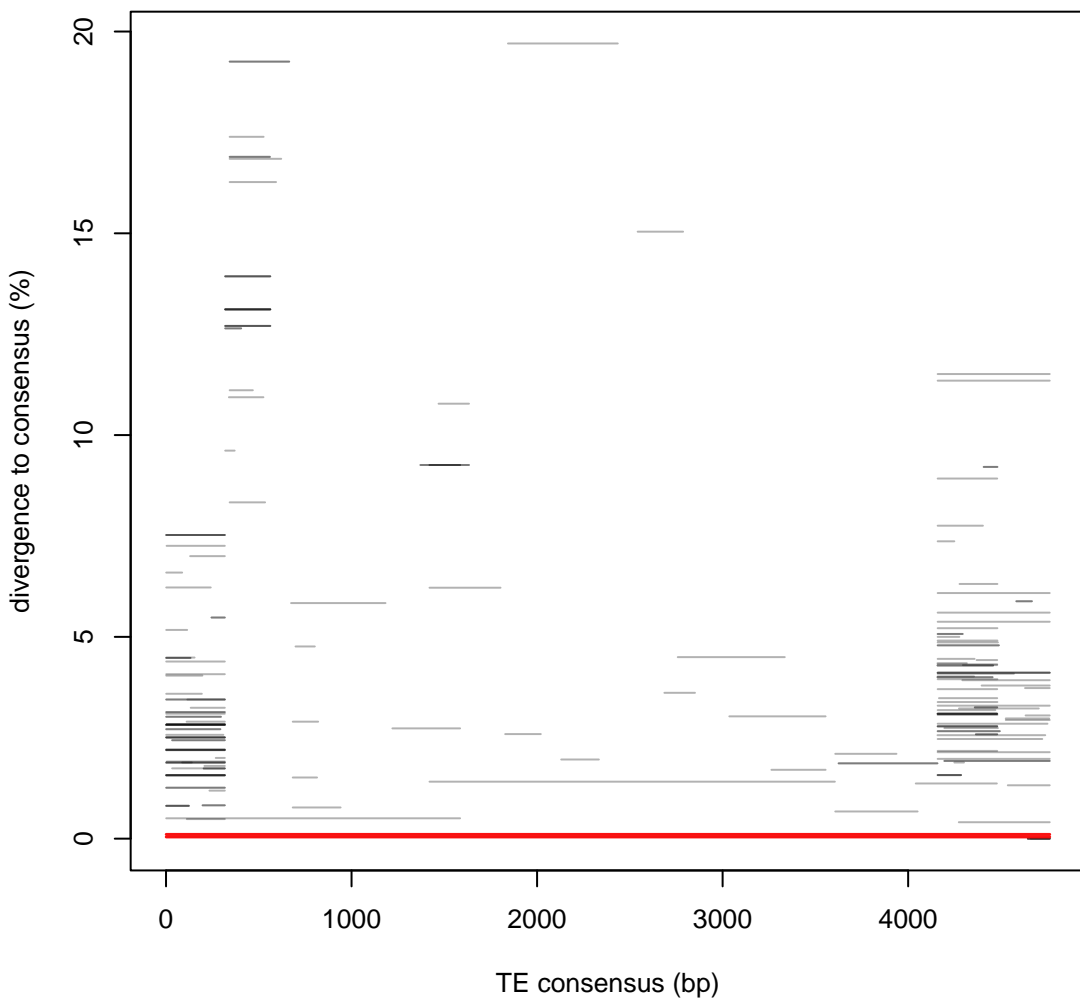
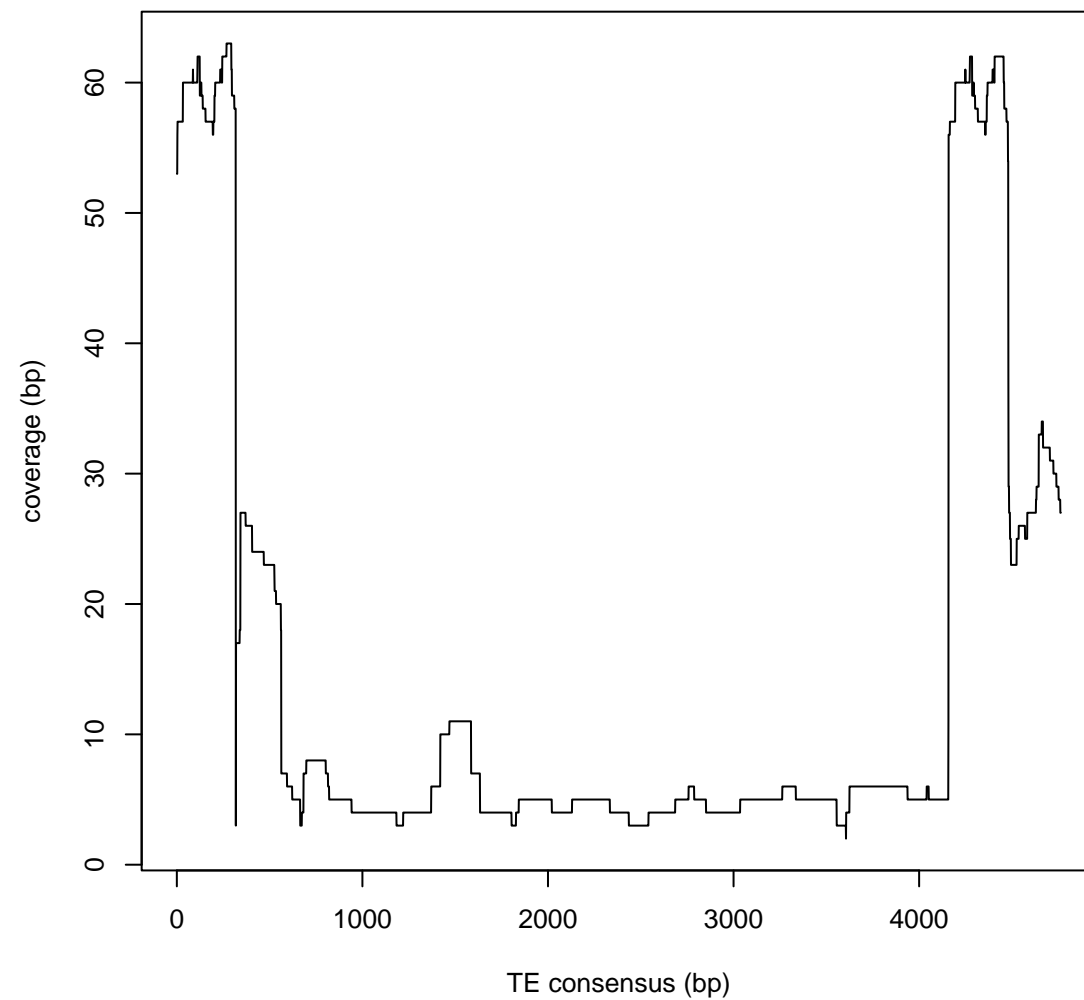


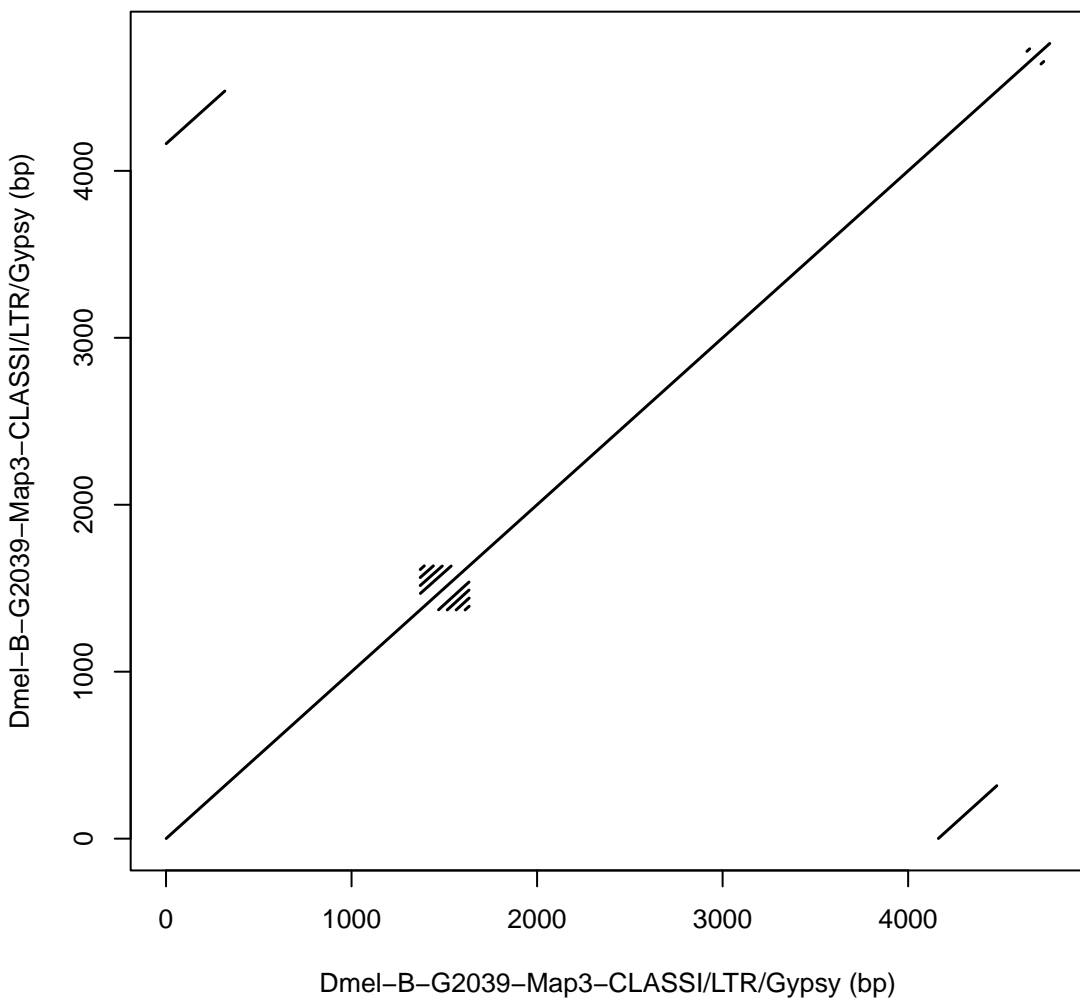
TE: Dmel-B-G2039-Map3-CLASSI/LTR/Gypsy
consensus size: 4763bp; fragments: 208; full length: 2 (≥ 4286.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

