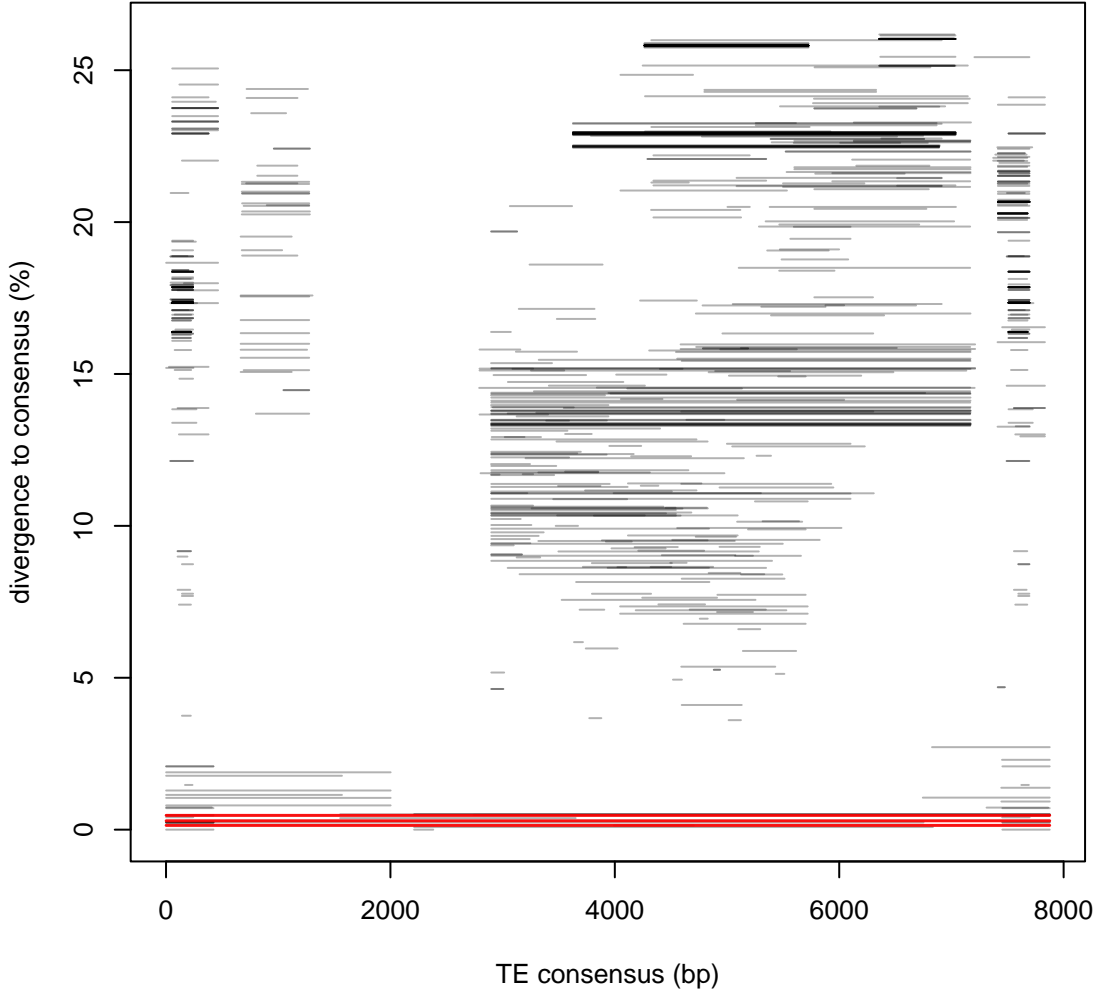
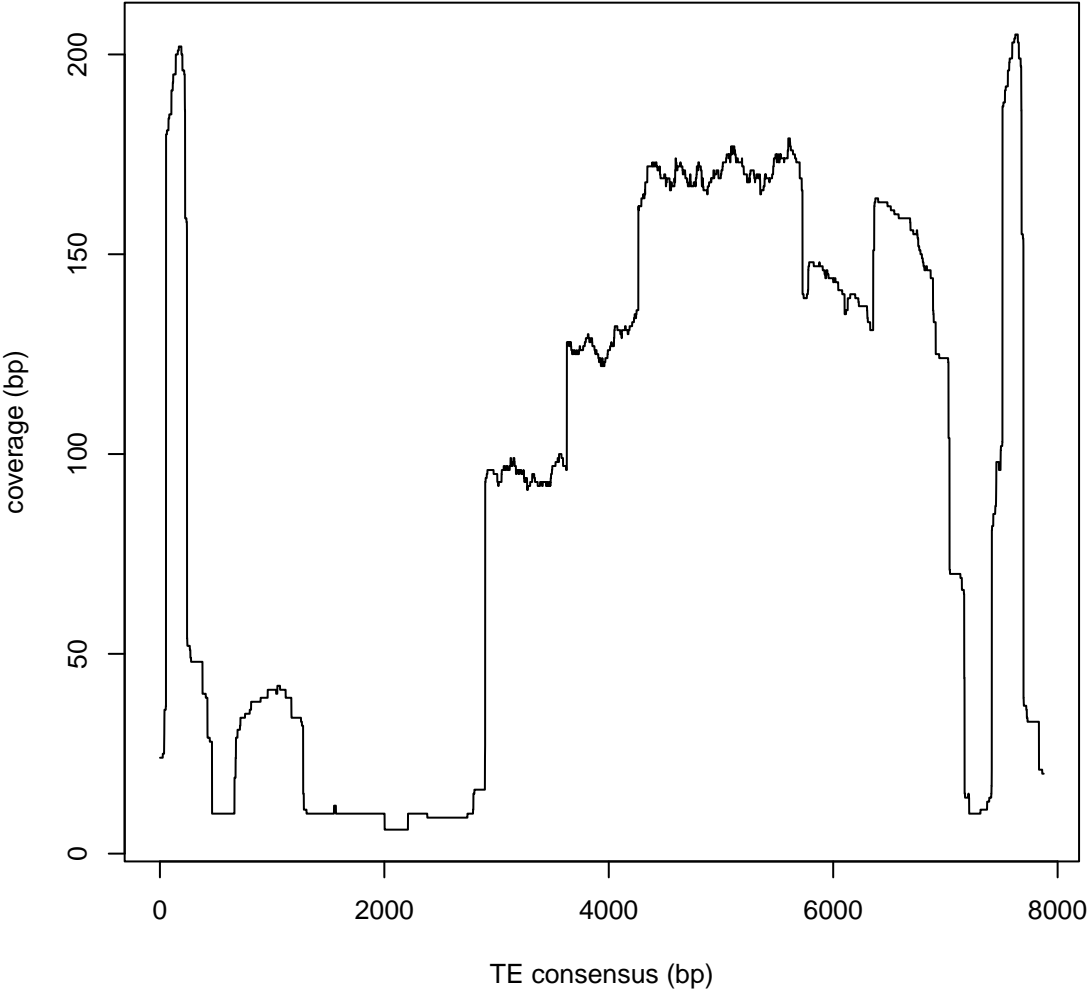


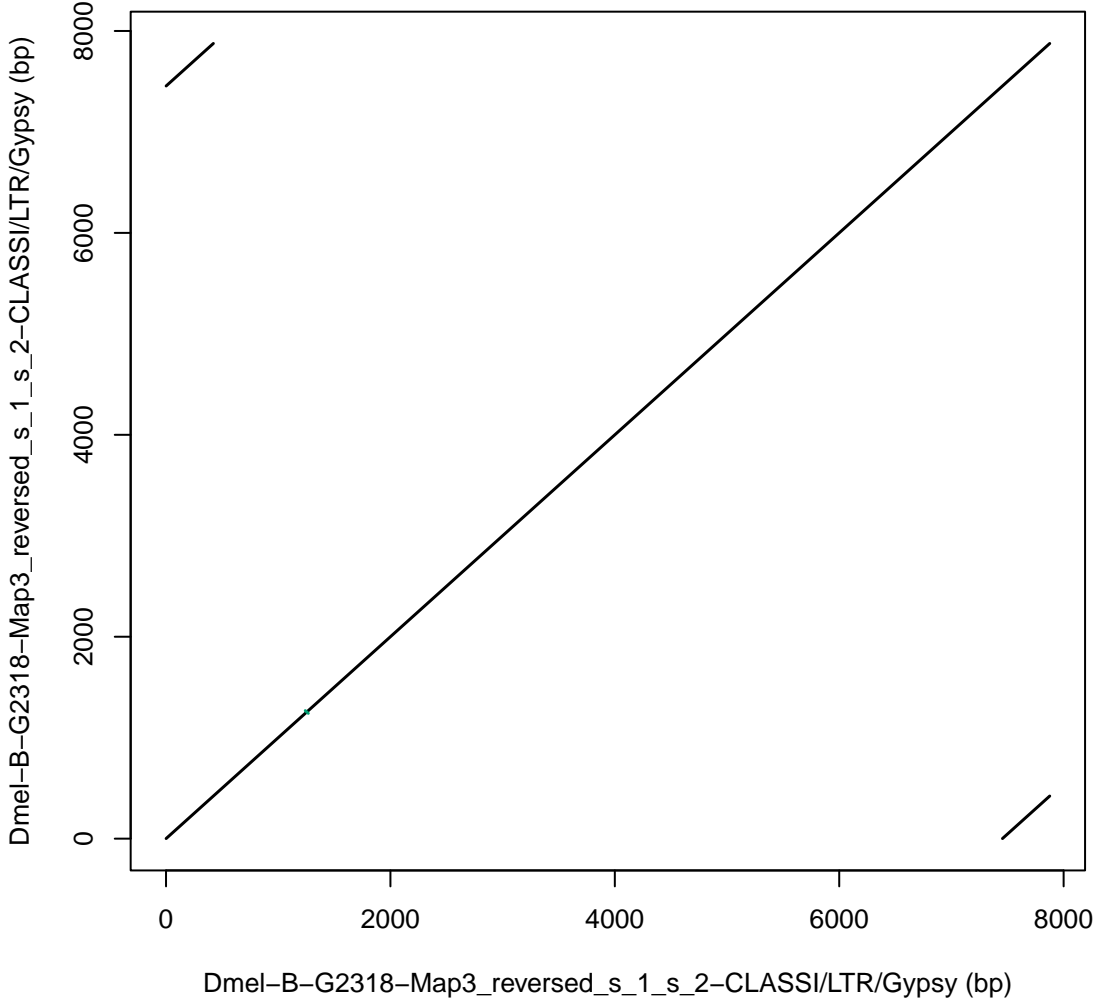
TE: Dmel-B-G2318-Map3\_reversed\_s\_1\_s\_2-CLASSI/LTR/Gypsy  
consensus size: 7876bp; fragments: 818; full length: 3 (>=7088.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

