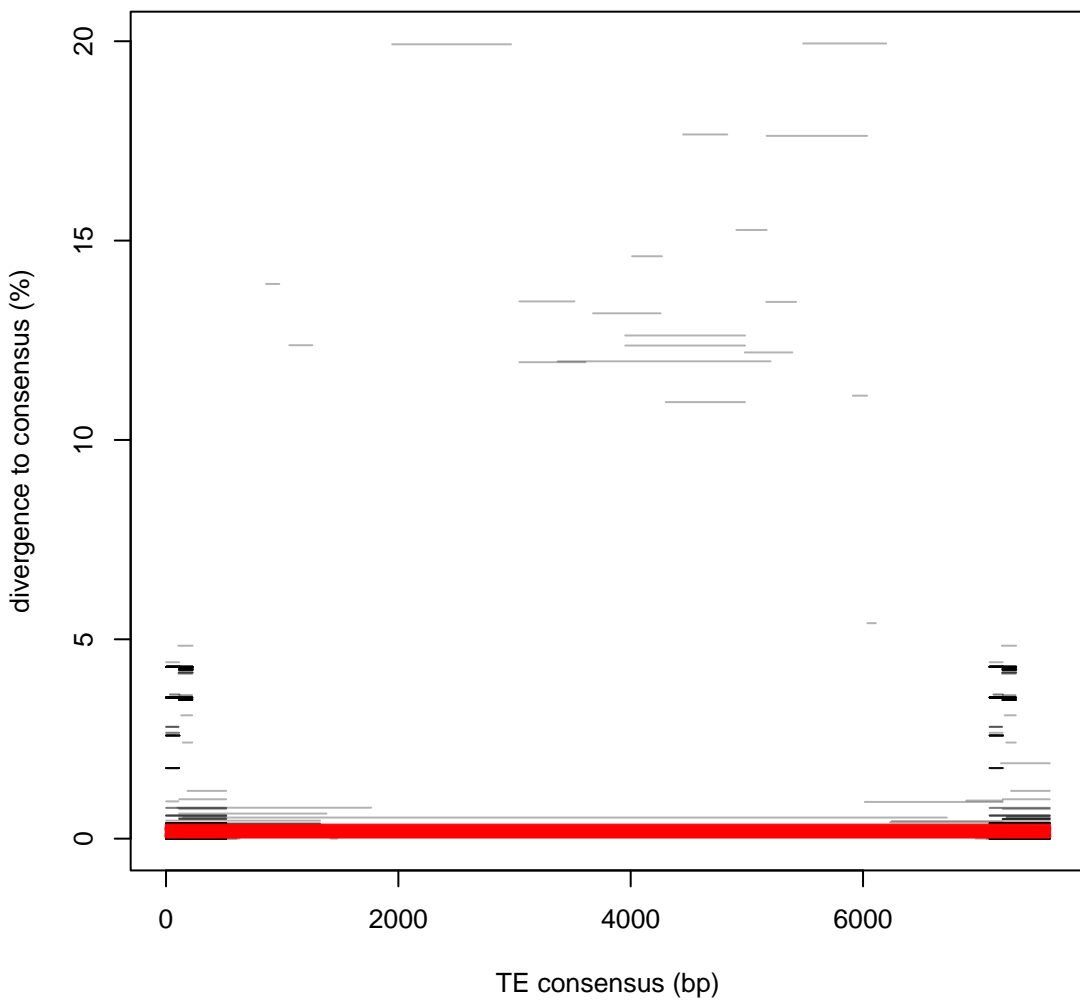
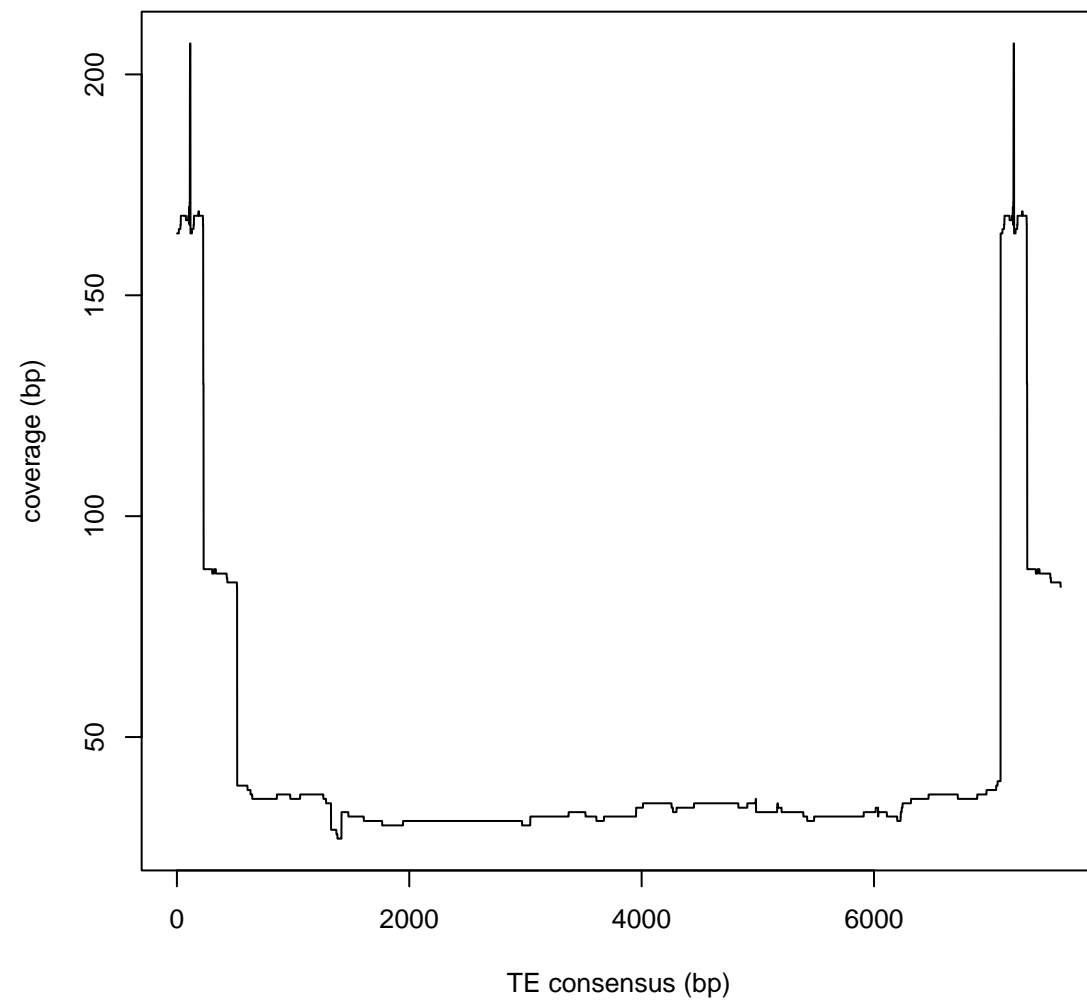


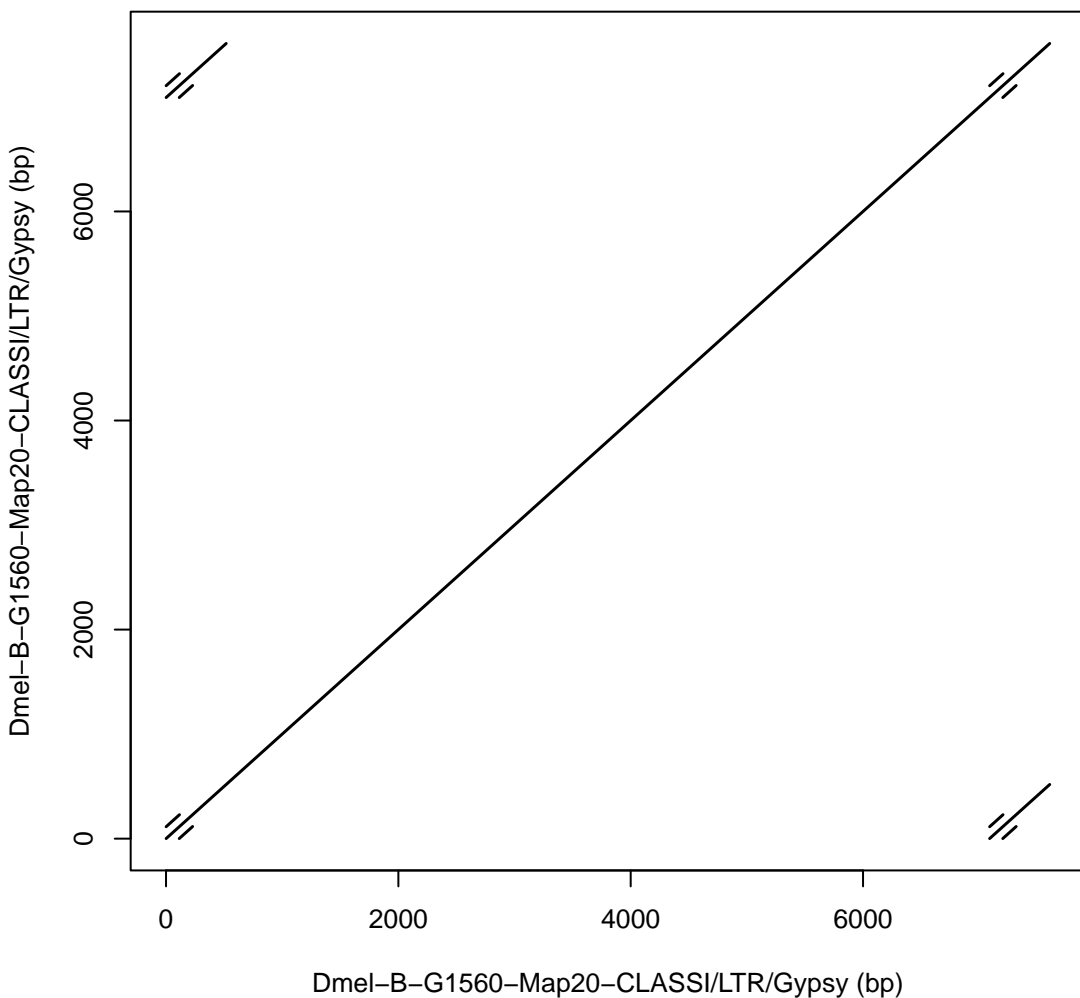
TE: Dmel-B-G1560-Map20-CLASSI/LTR/Gypsy
consensus size: 7607bp; fragments: 516; full length: 24 (>=6846.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

