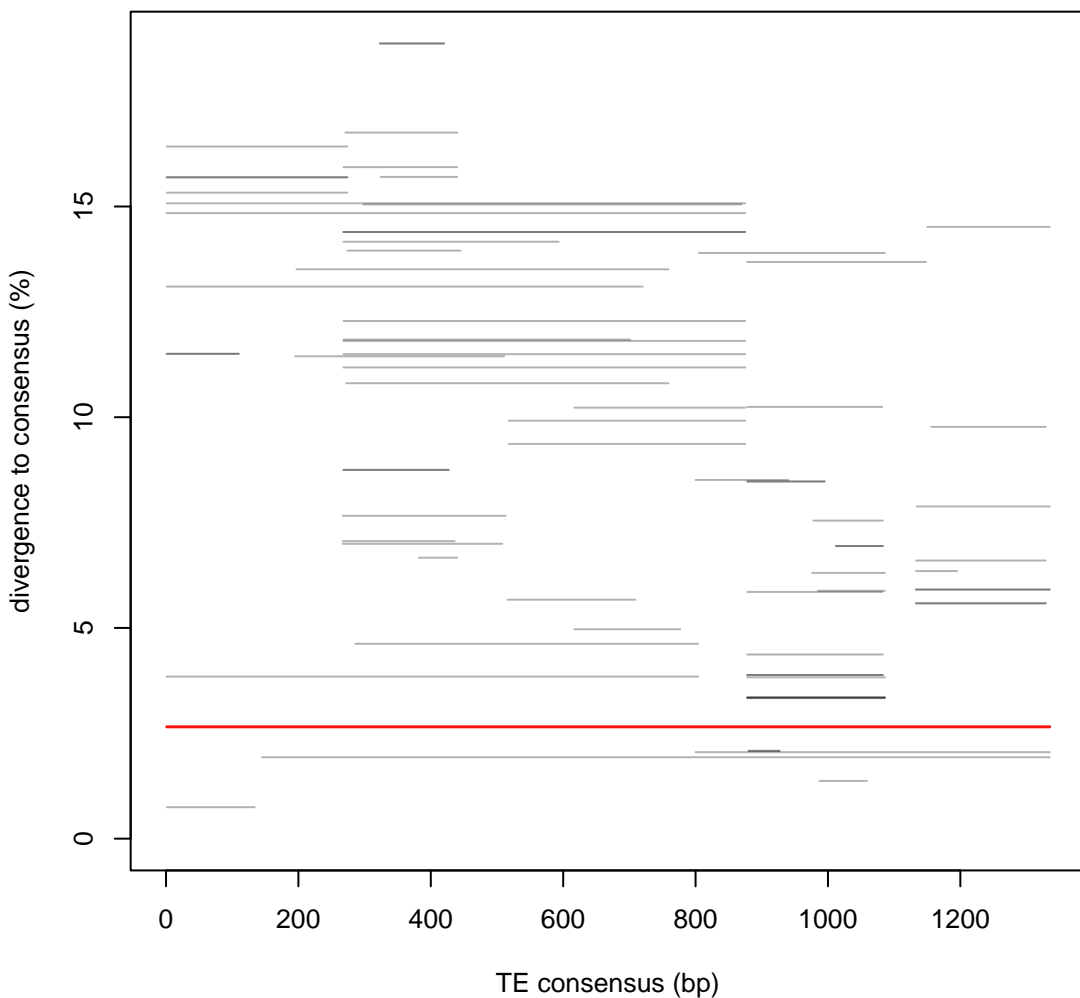
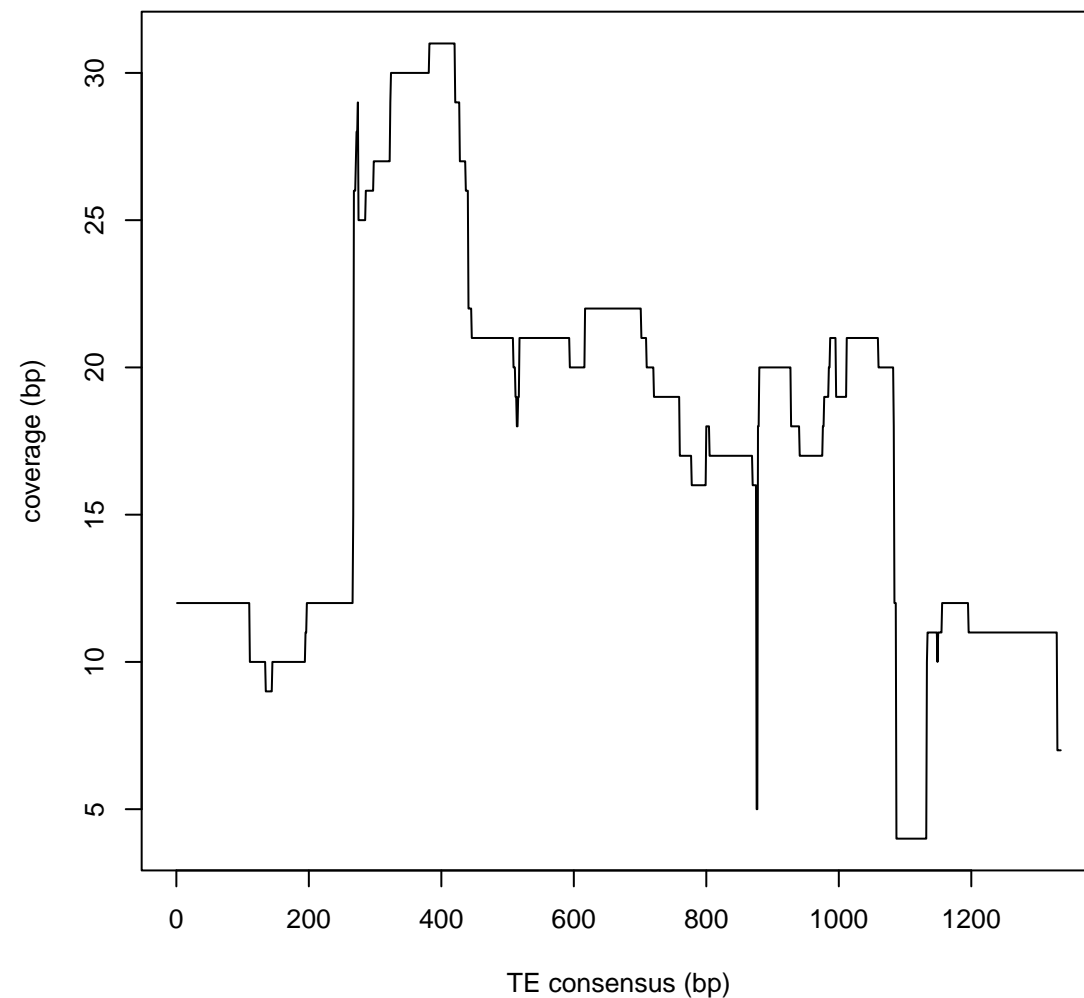


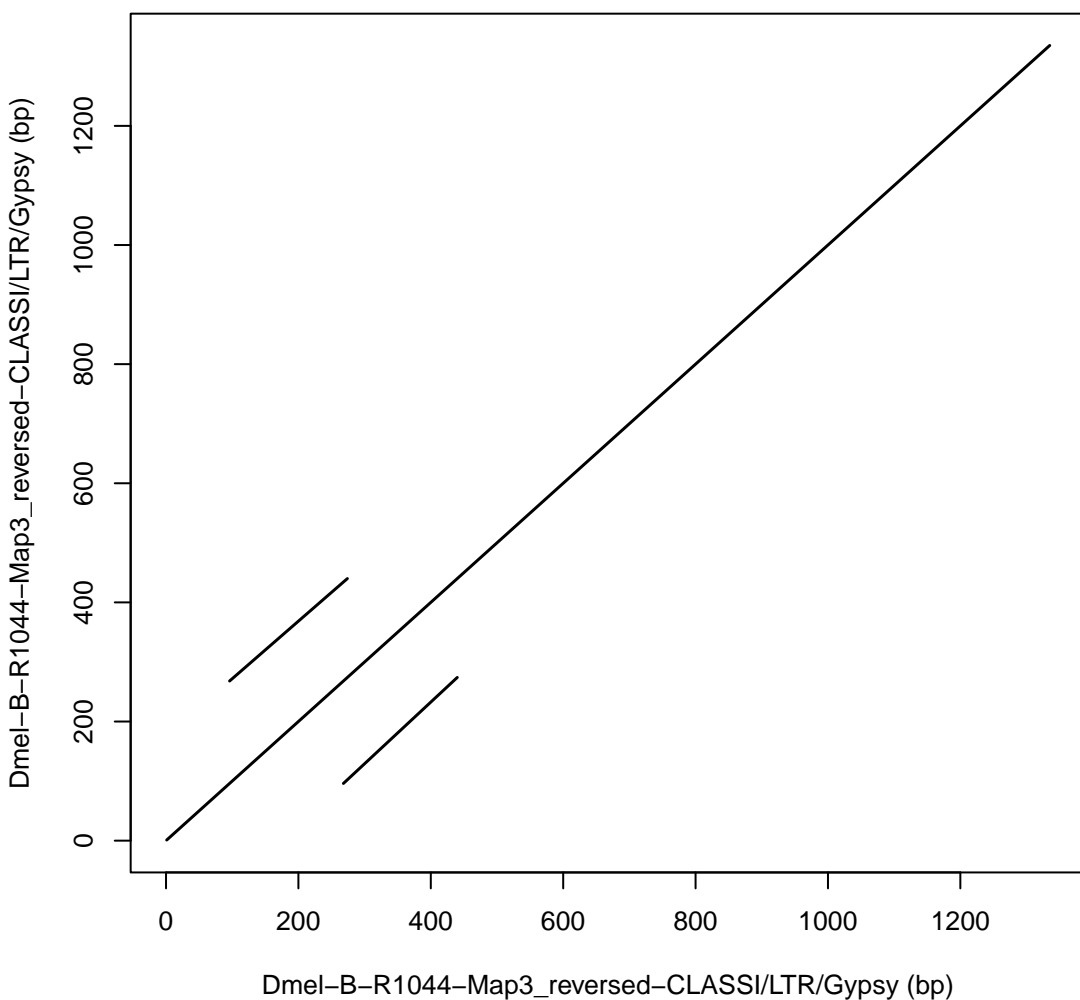
TE: Dmel-B-R1044-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 1335bp; fragments: 76; full length: 1 (>=1201.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

