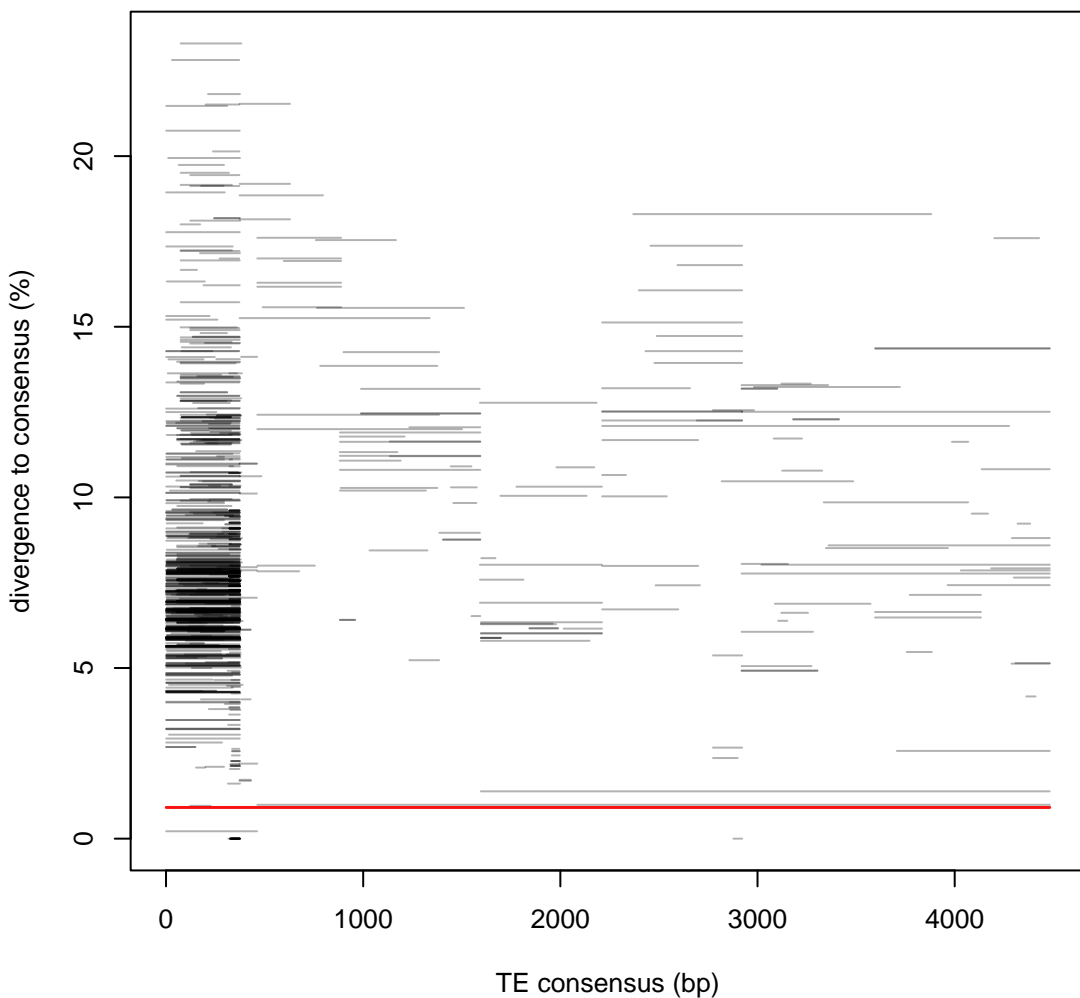
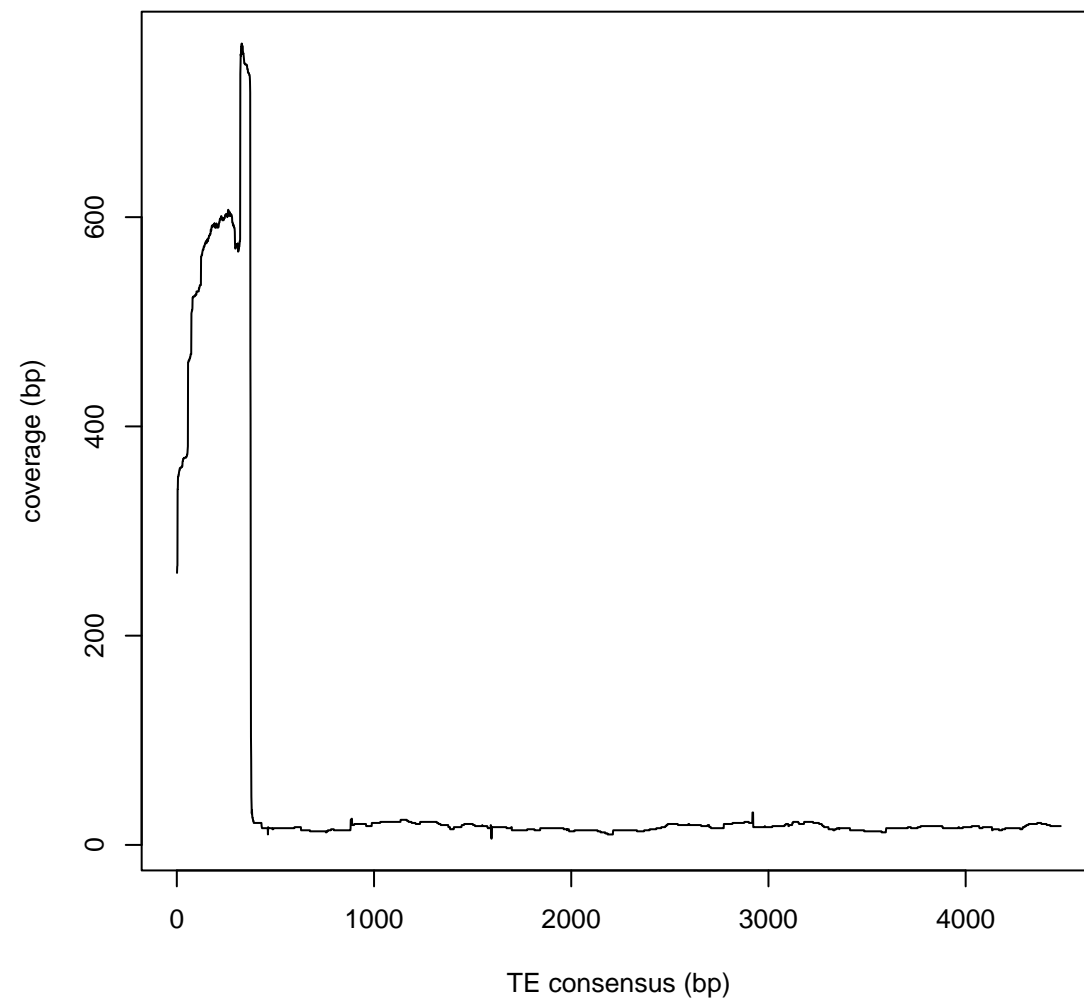


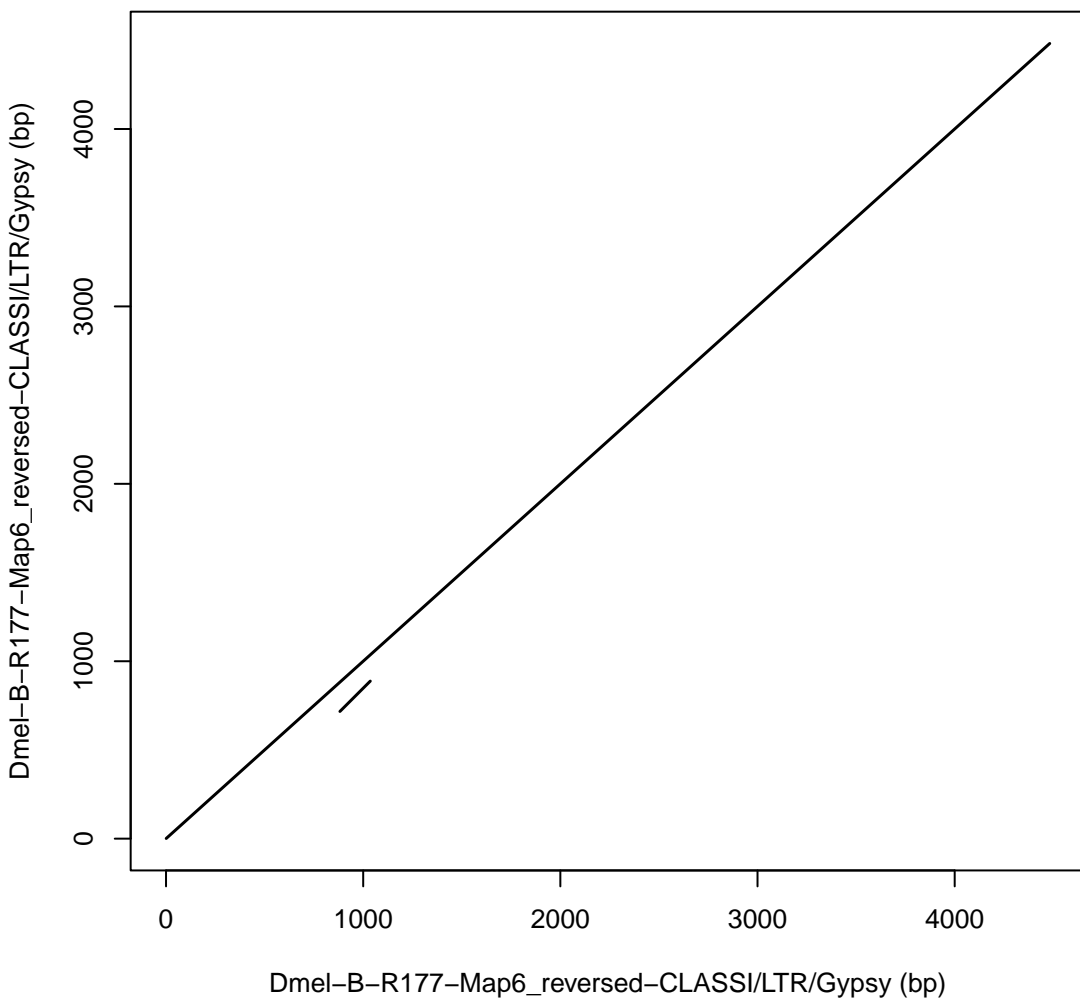
TE: Dmel-B-R177-Map6_reversed-CLASSI/LTR/Gypsy
consensus size: 4482bp; fragments: 1033; full length: 1 (>=4033.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

