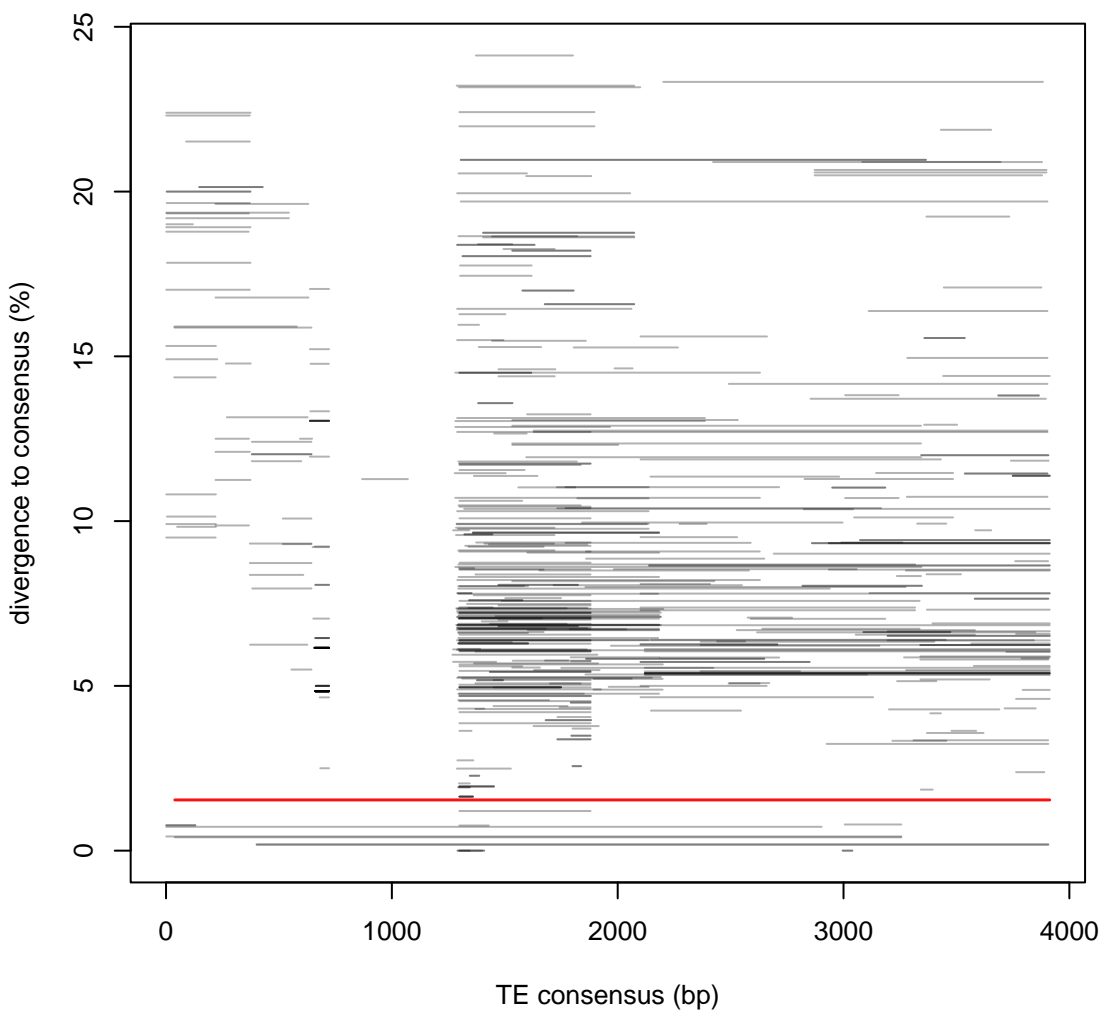
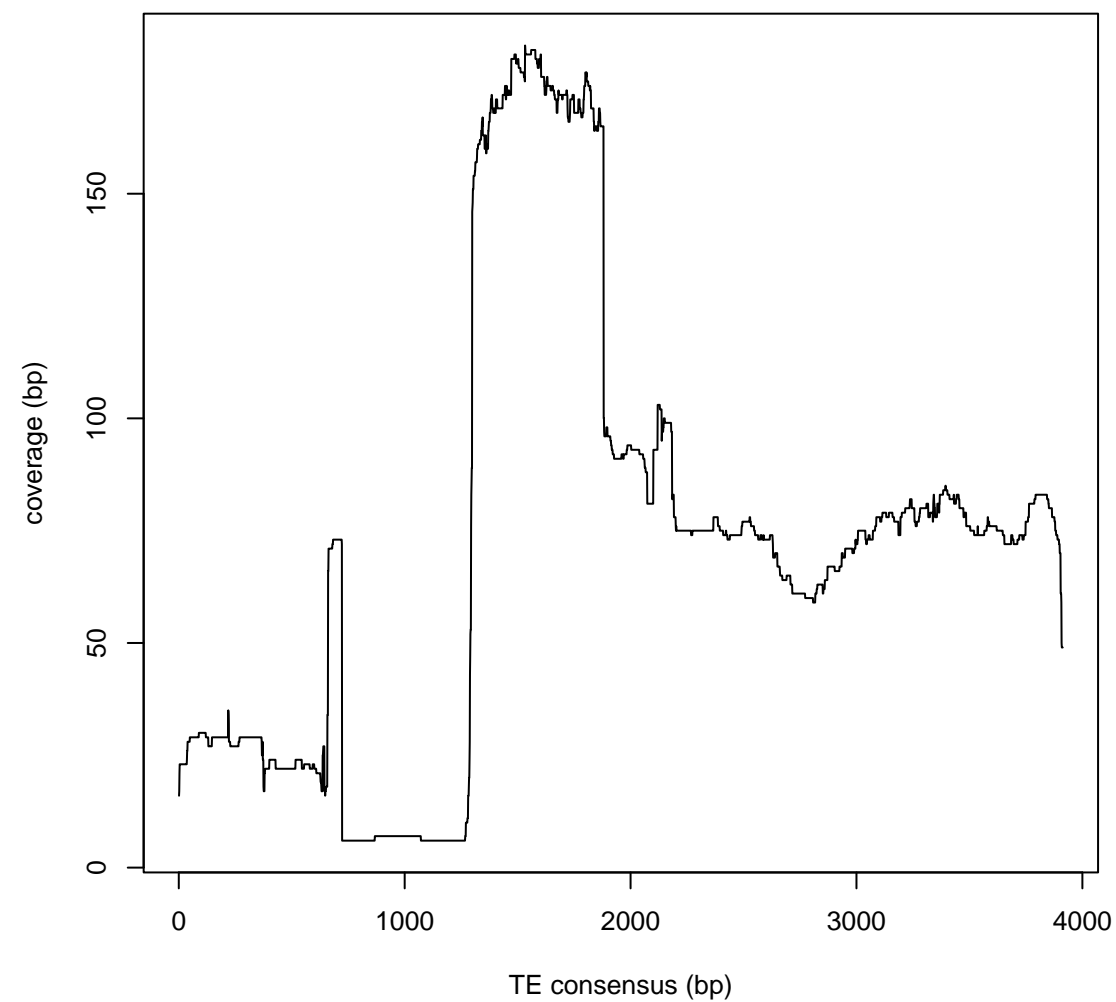


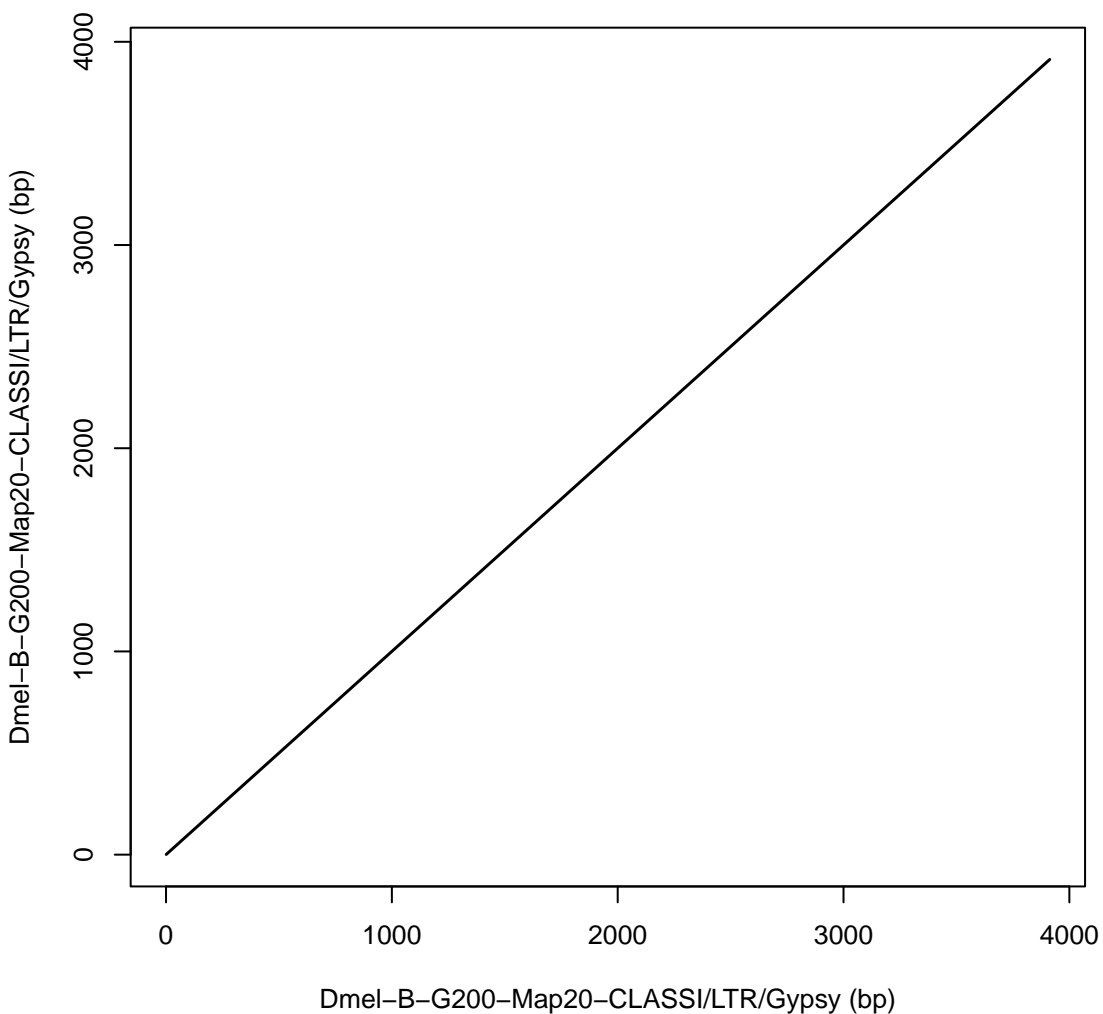
TE: Dmel-B-G200-Map20-CLASSI/LTR/Gypsy
consensus size: 3913bp; fragments: 528; full length: 1 (>=3521.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

