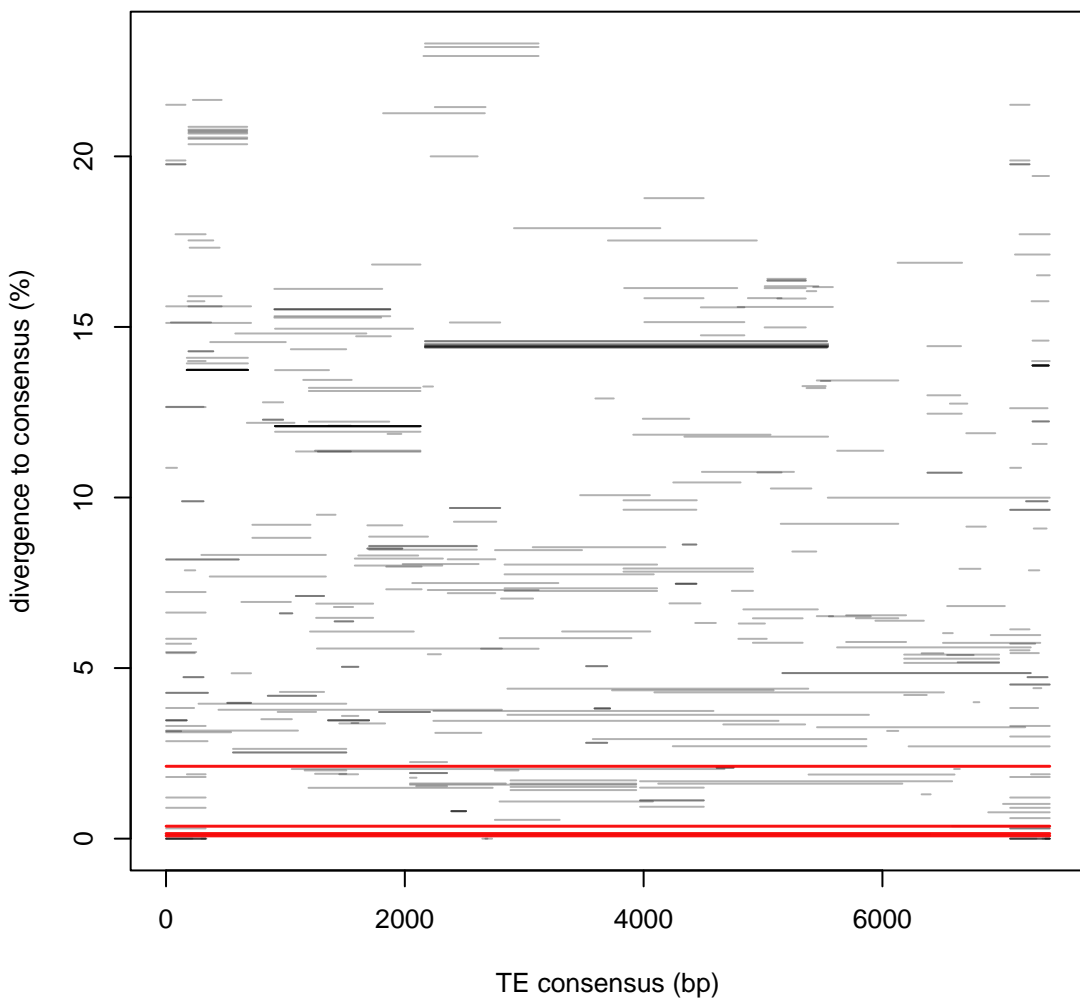
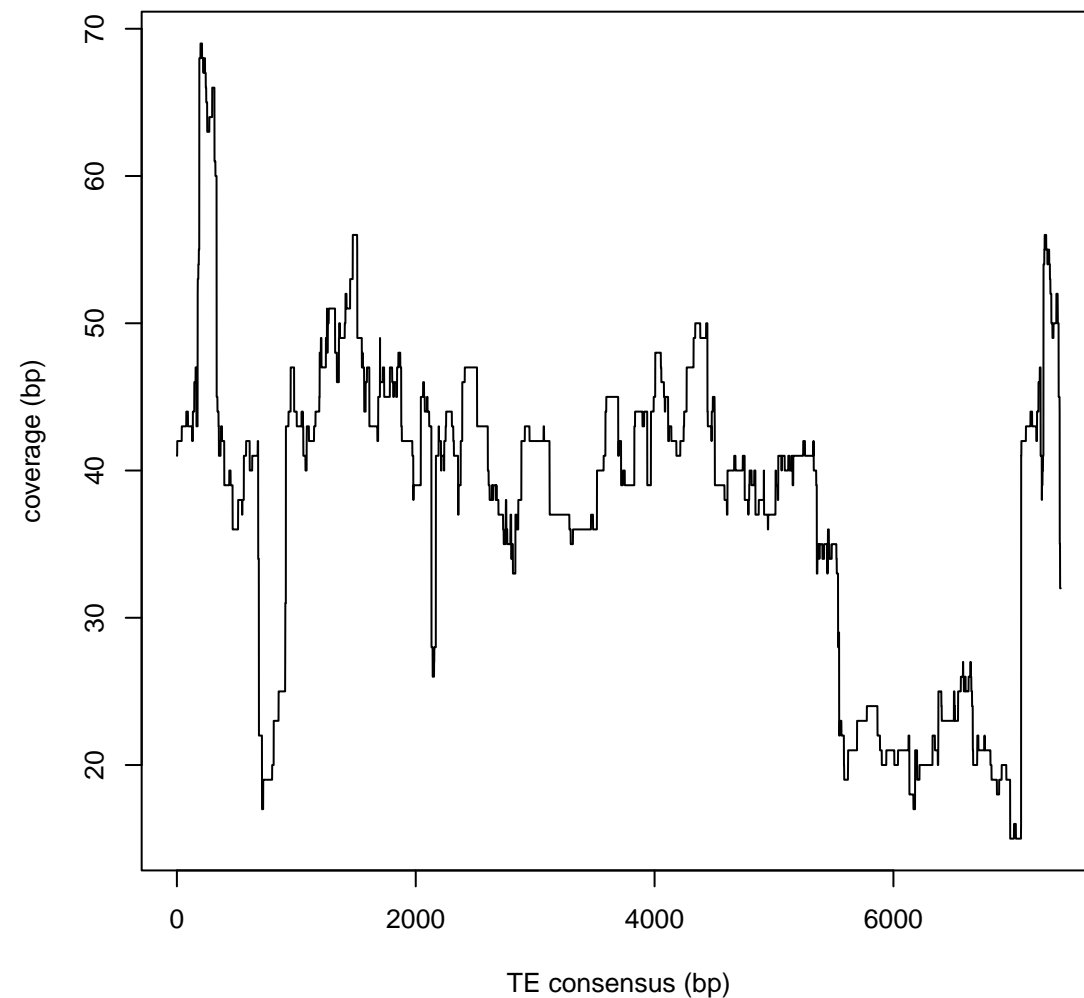


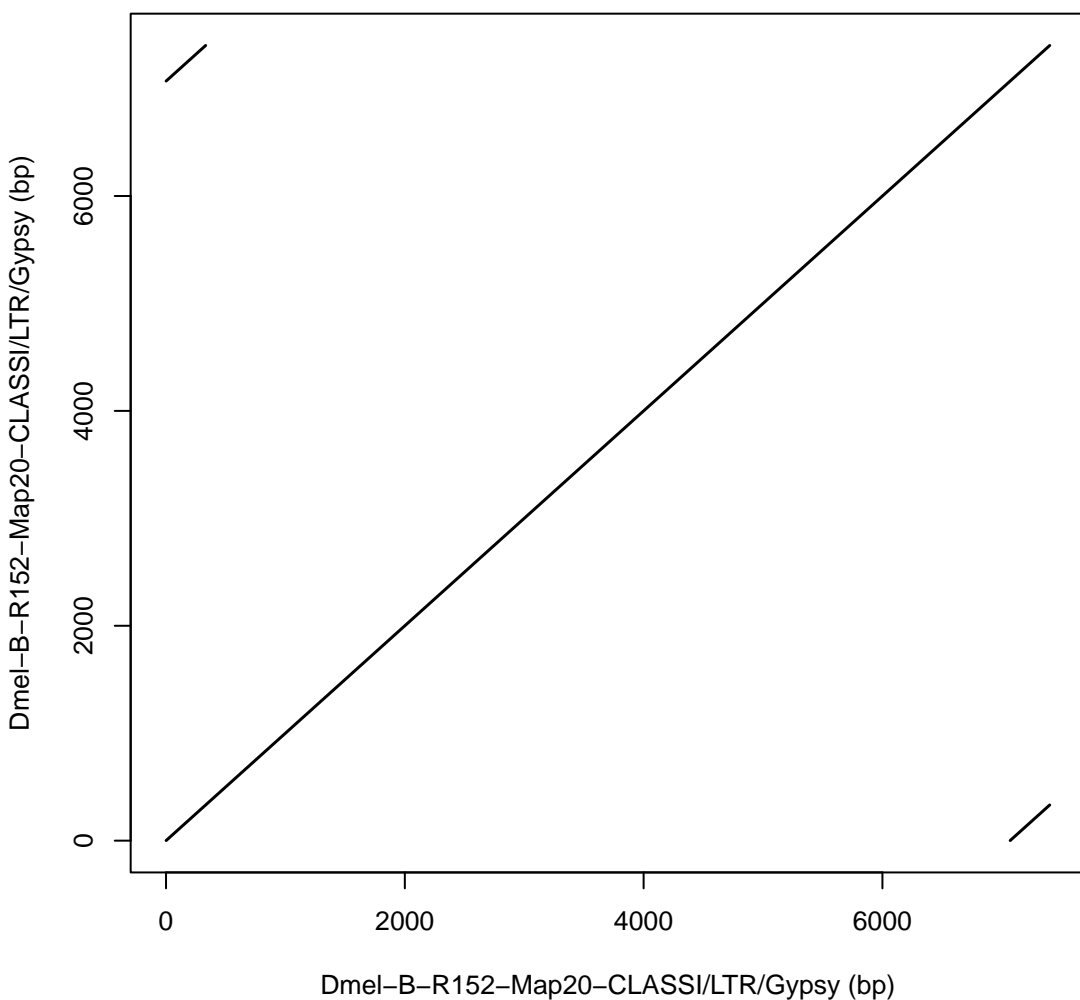
TE: Dmel-B-R152-Map20-CLASSI/LTR/Gypsy
consensus size: 7401bp; fragments: 406; full length: 4 (>=6660.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

