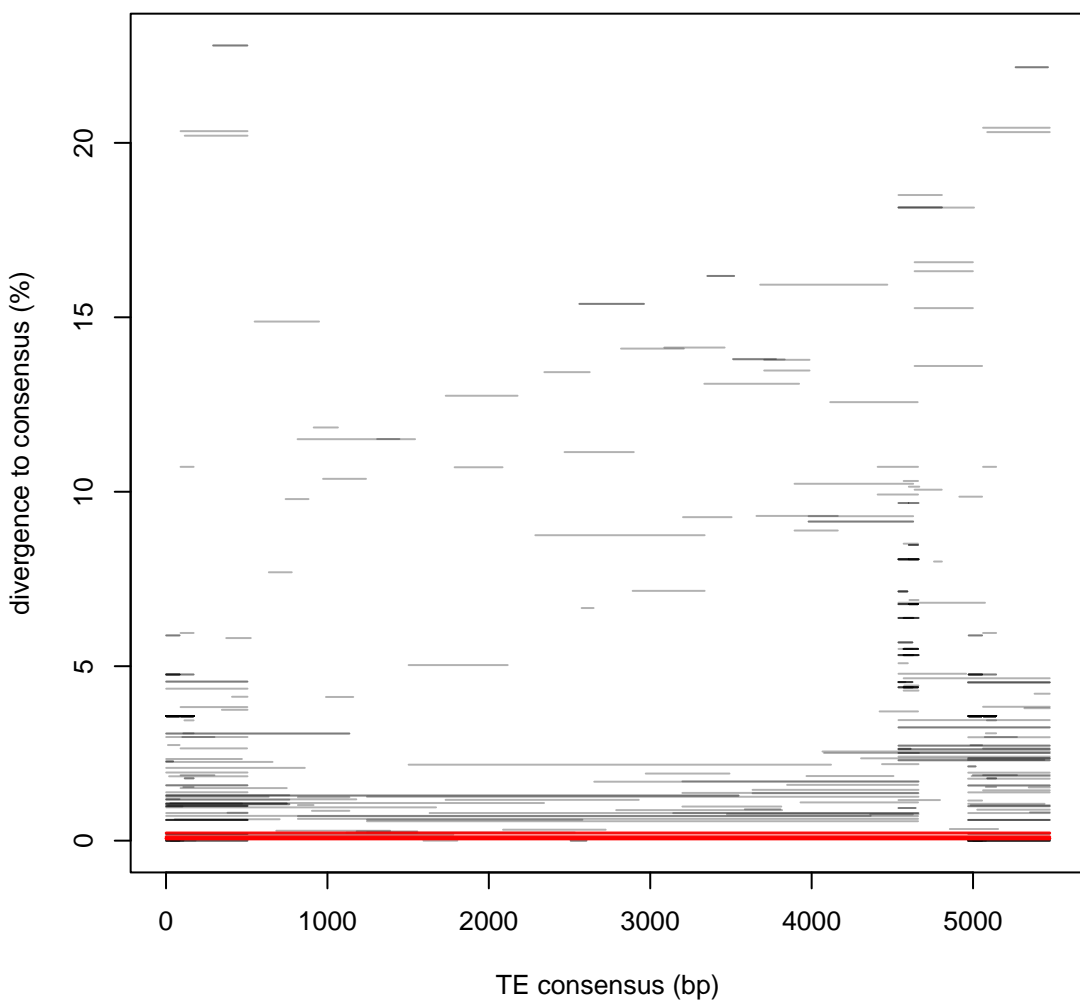
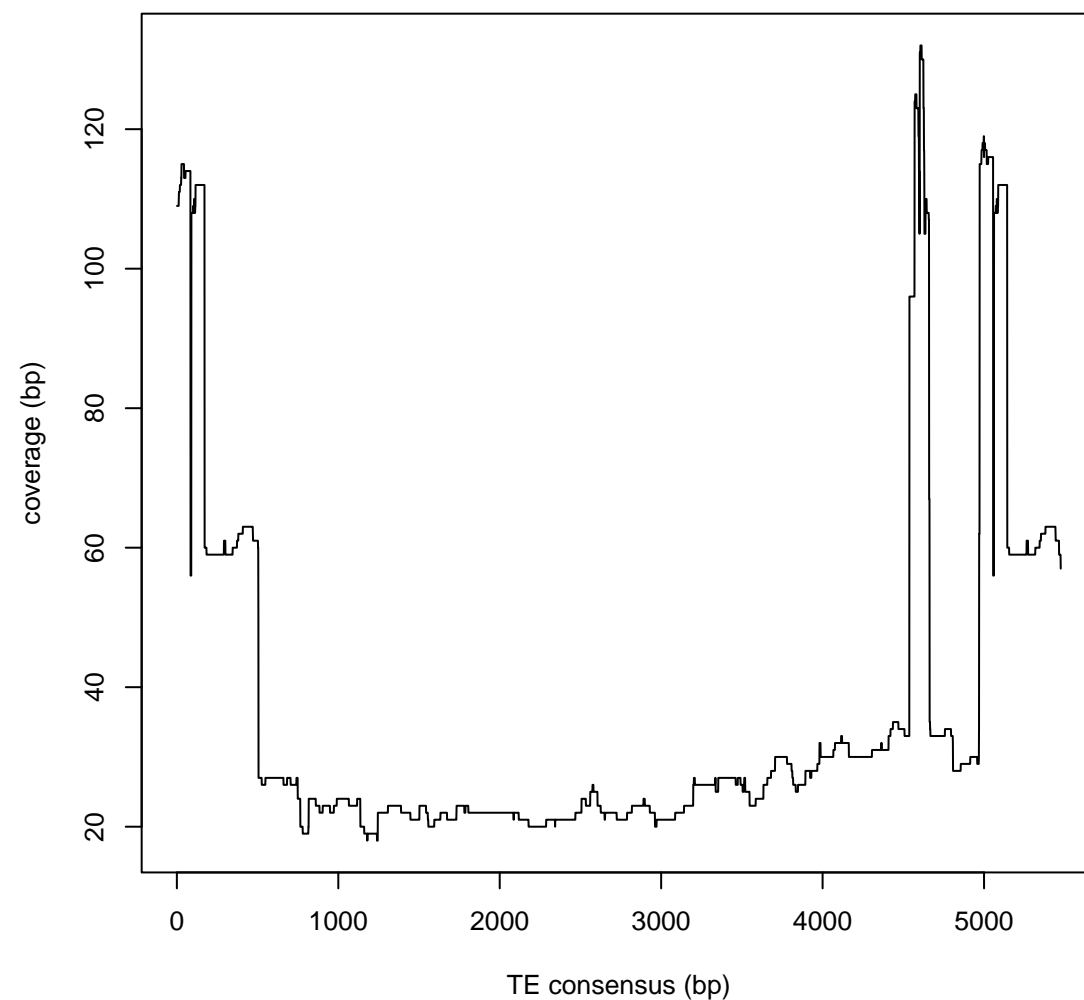


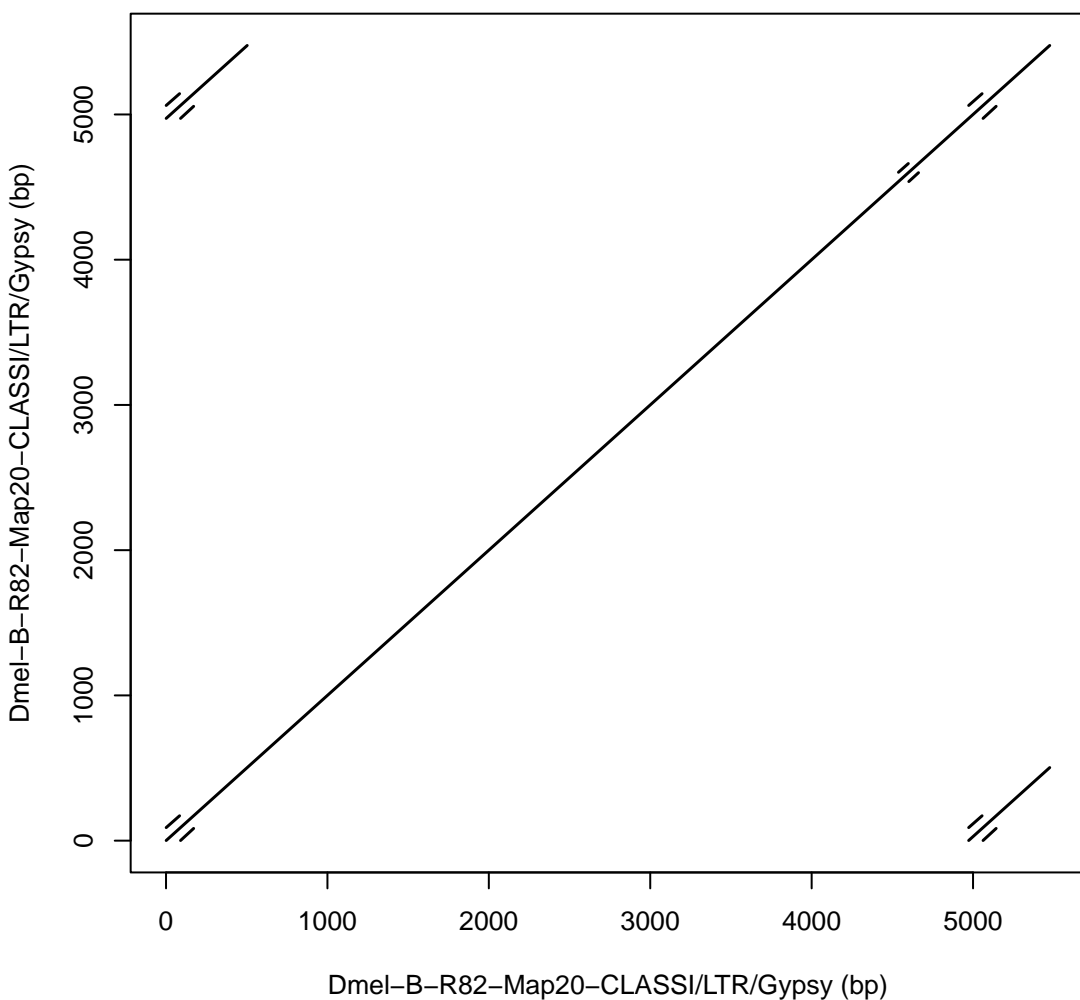
TE: Dmel-B-R82-Map20-CLASSI/LTR/Gypsy
consensus size: 5475bp; fragments: 543; full length: 6 (≥ 4927.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

