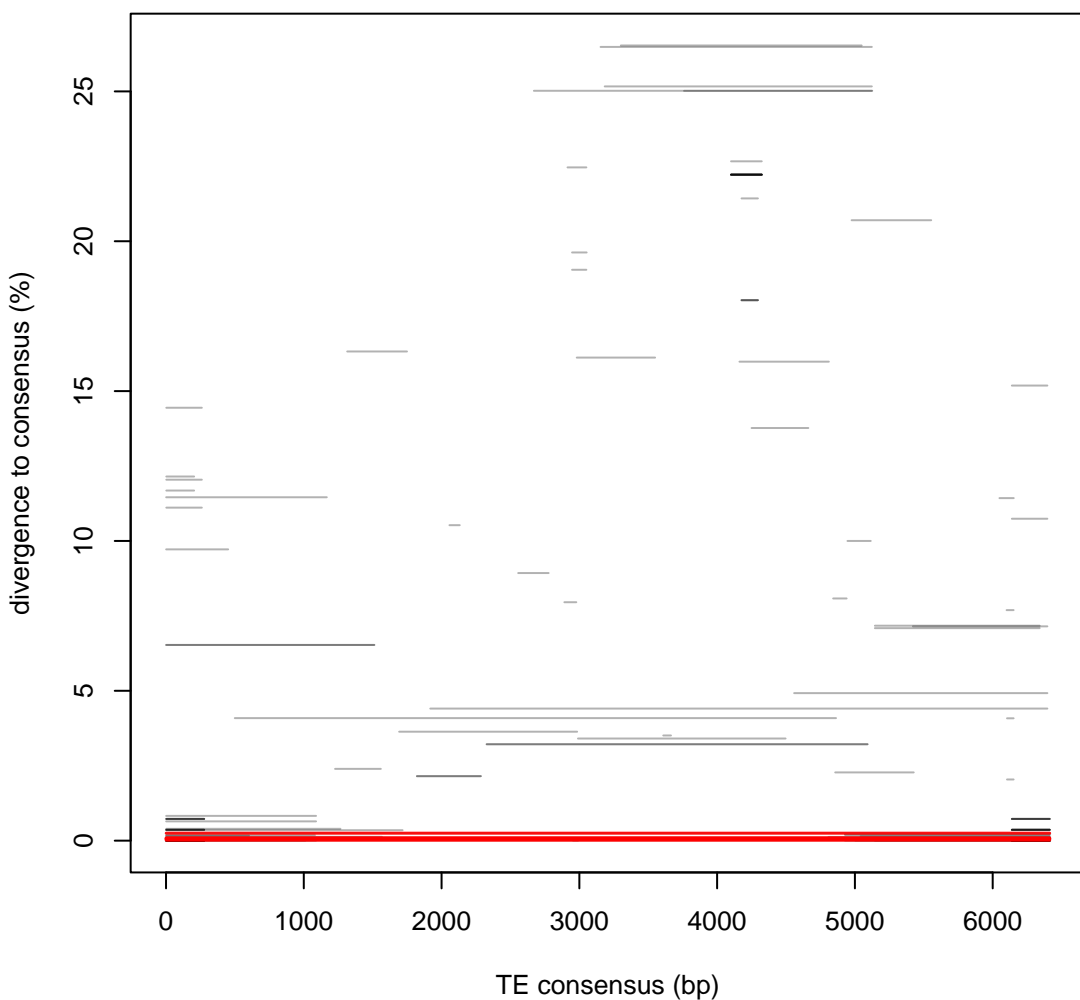
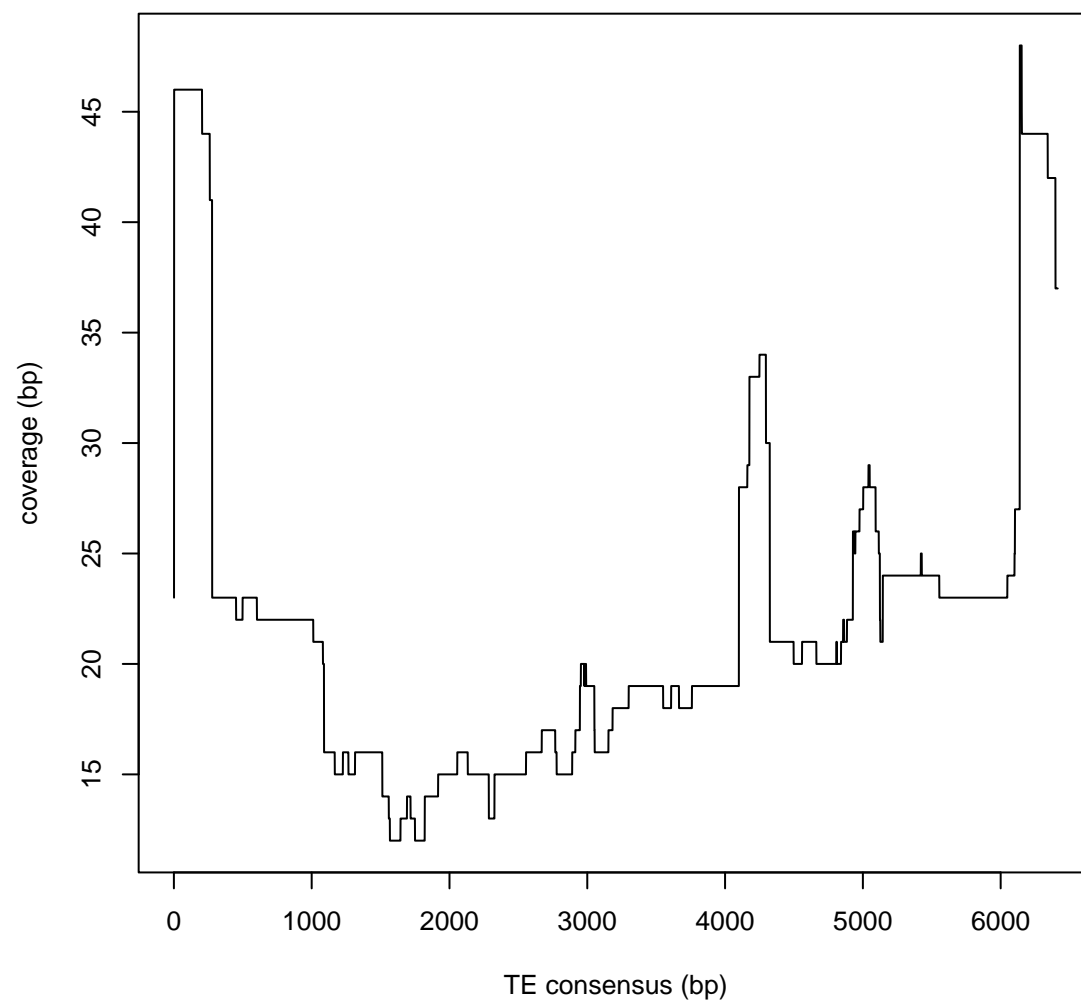


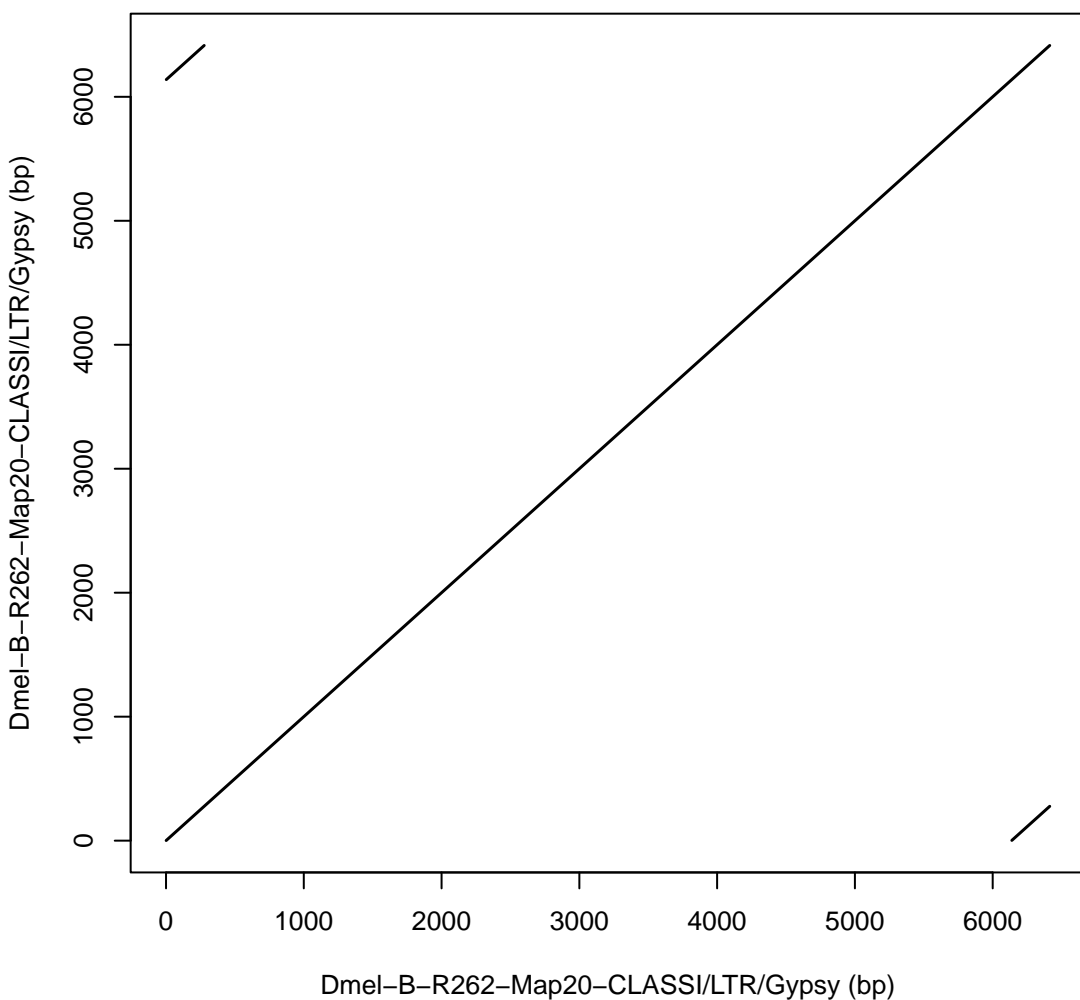
TE: Dmel-B-R262-Map20-CLASSI/LTR/Gypsy
consensus size: 6414bp; fragments: 128; full length: 8 (>=5772.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

