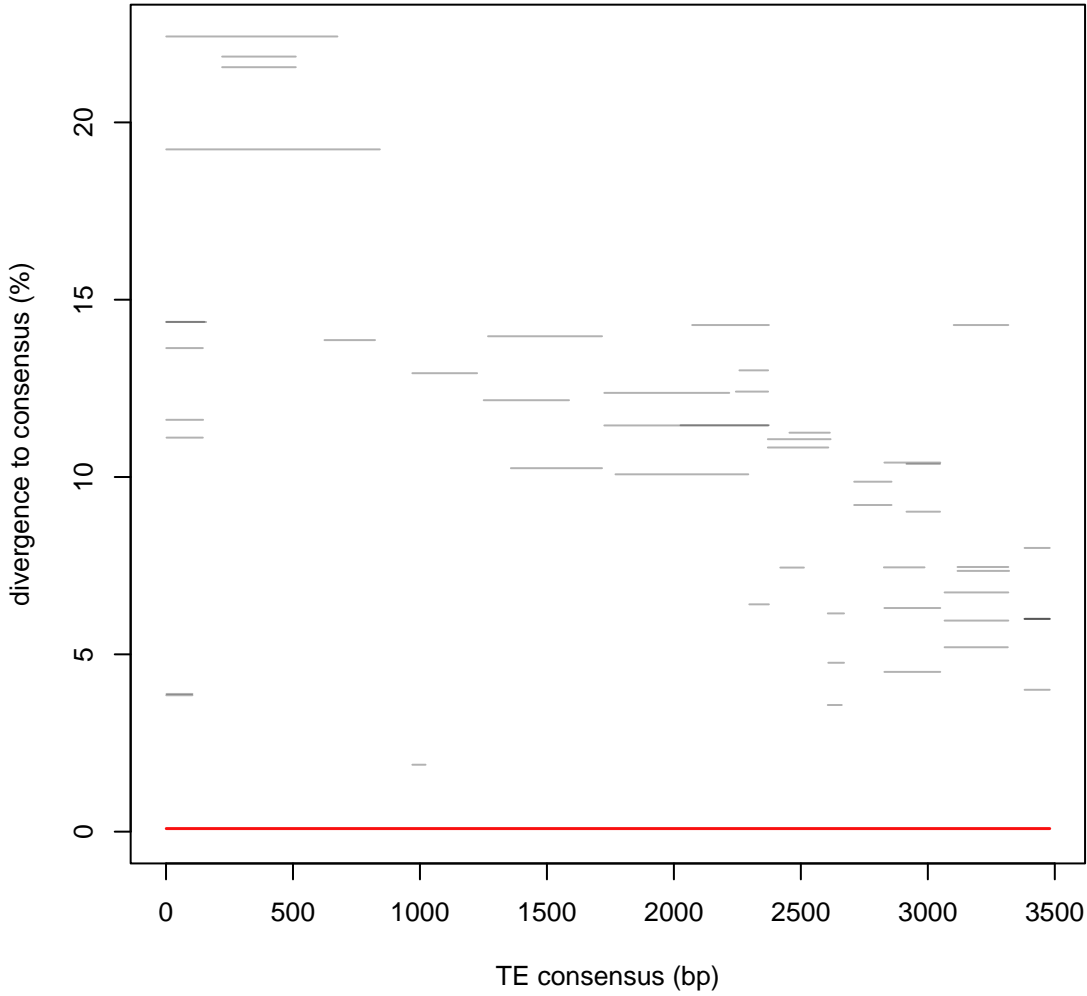
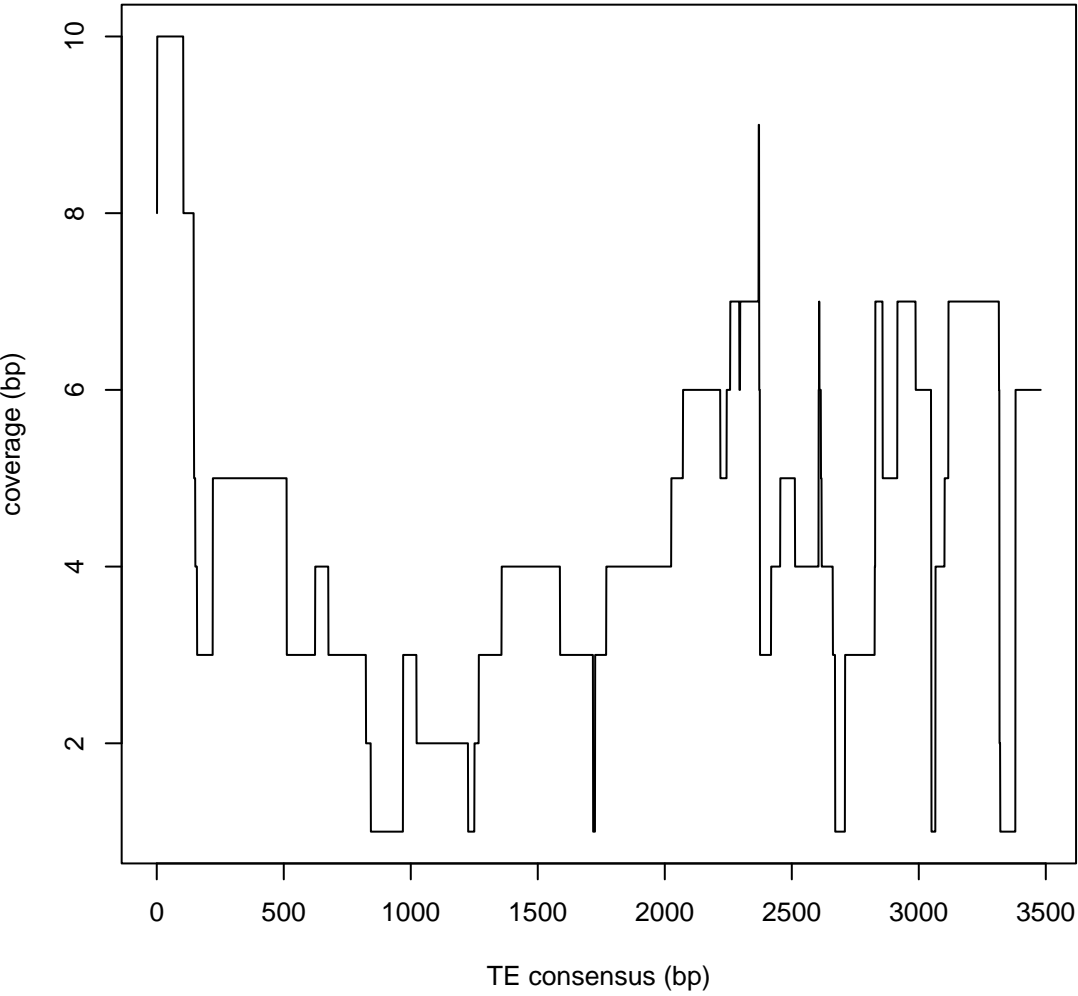


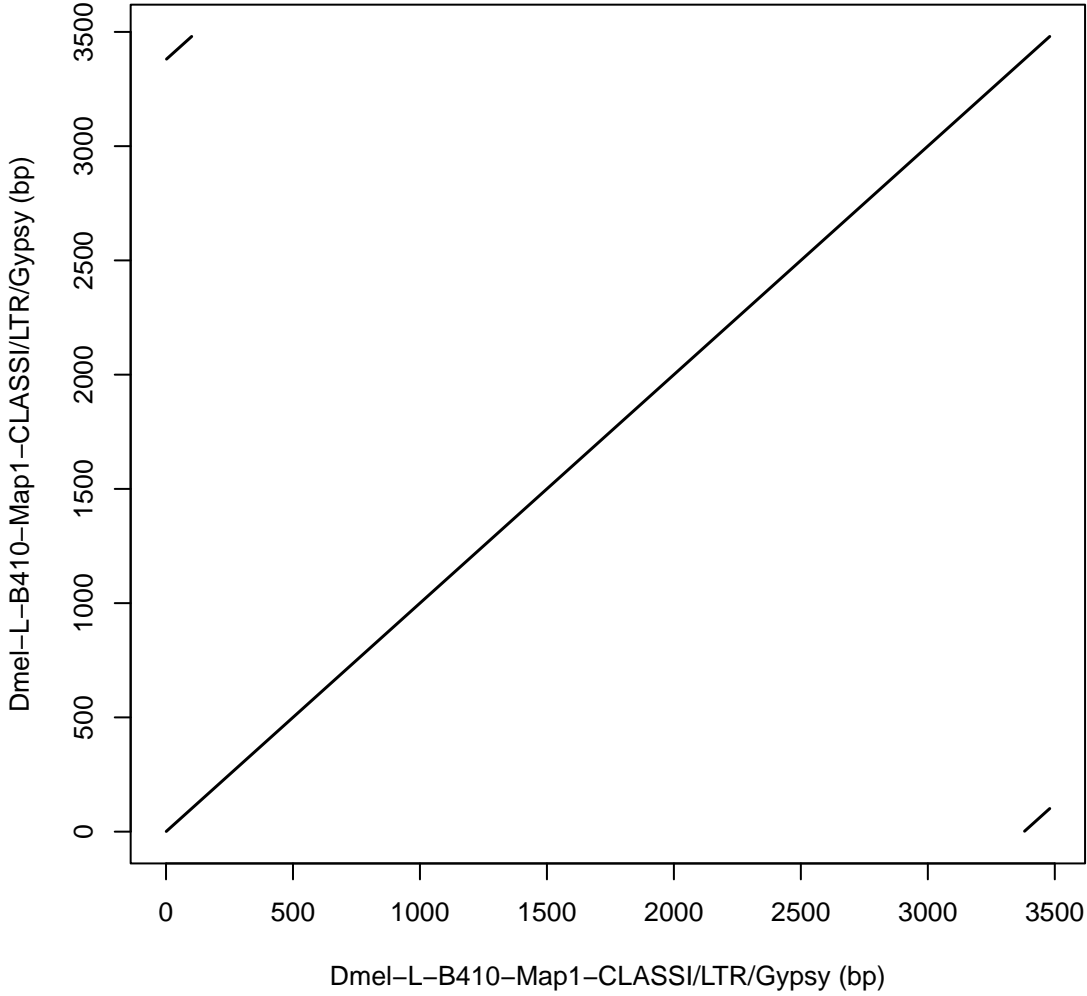
TE: Dmel-L-B410-Map1-CLASSI/LTR/Gypsy
consensus size: 3480bp; fragments: 52; full length: 1 (>=3132bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

