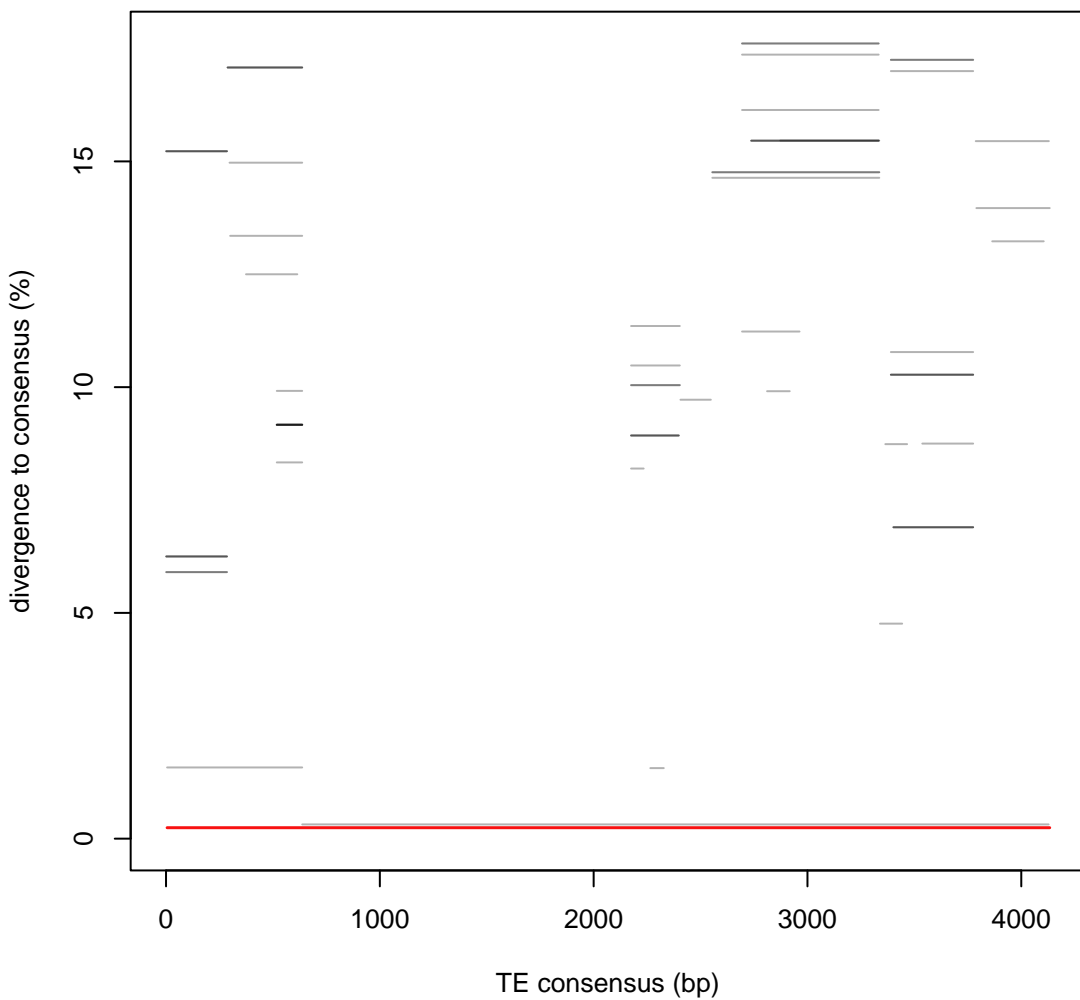
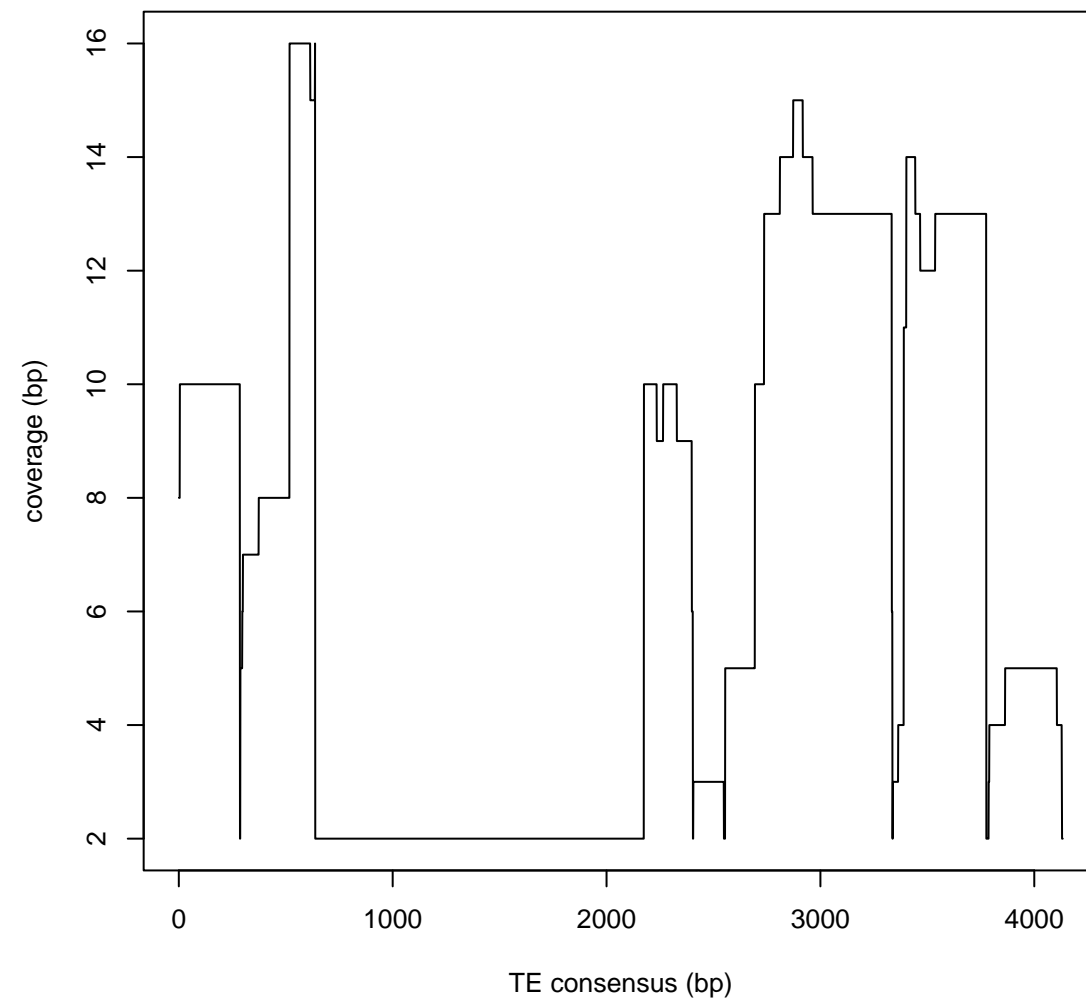


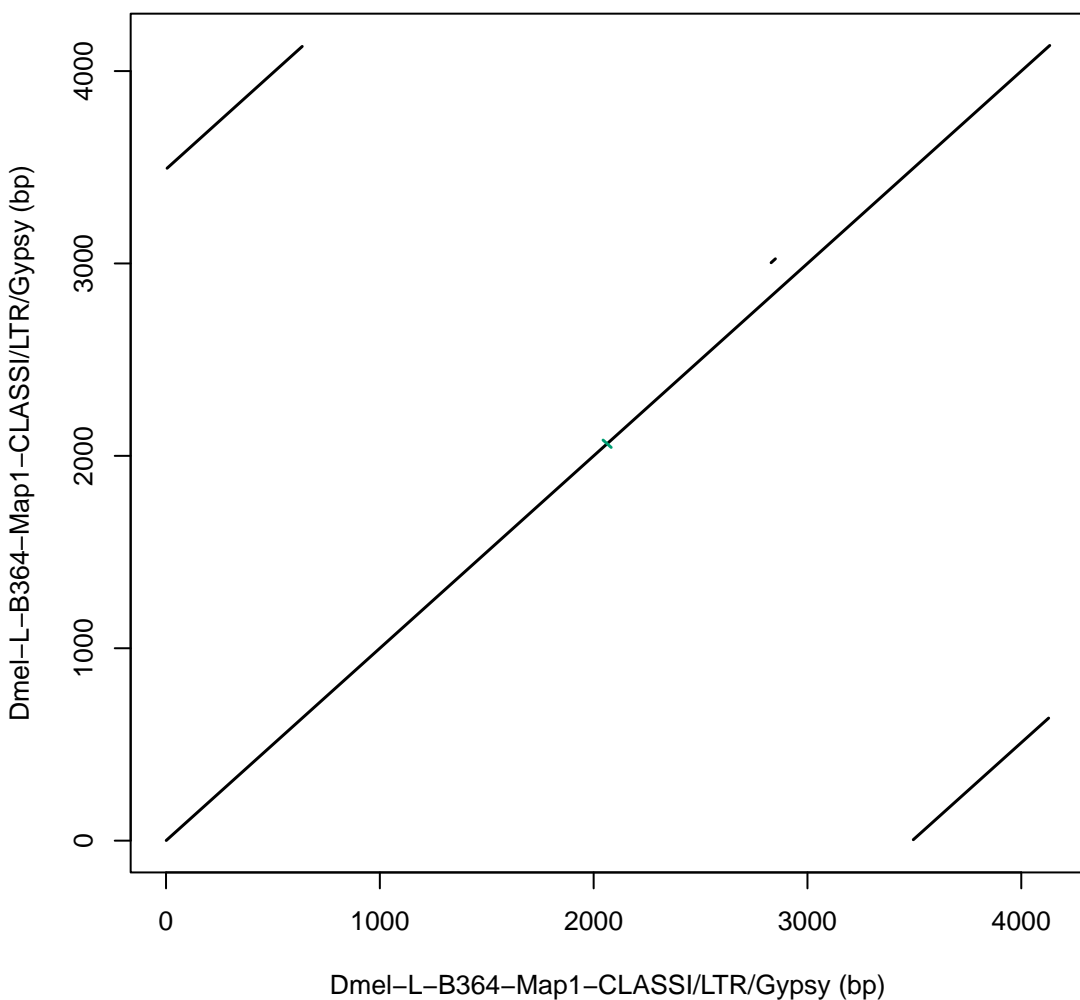
TE: Dmel-L-B364-Map1-CLASSI/LTR/Gypsy
consensus size: 4133bp; fragments: 64; full length: 1 (>=3719.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

