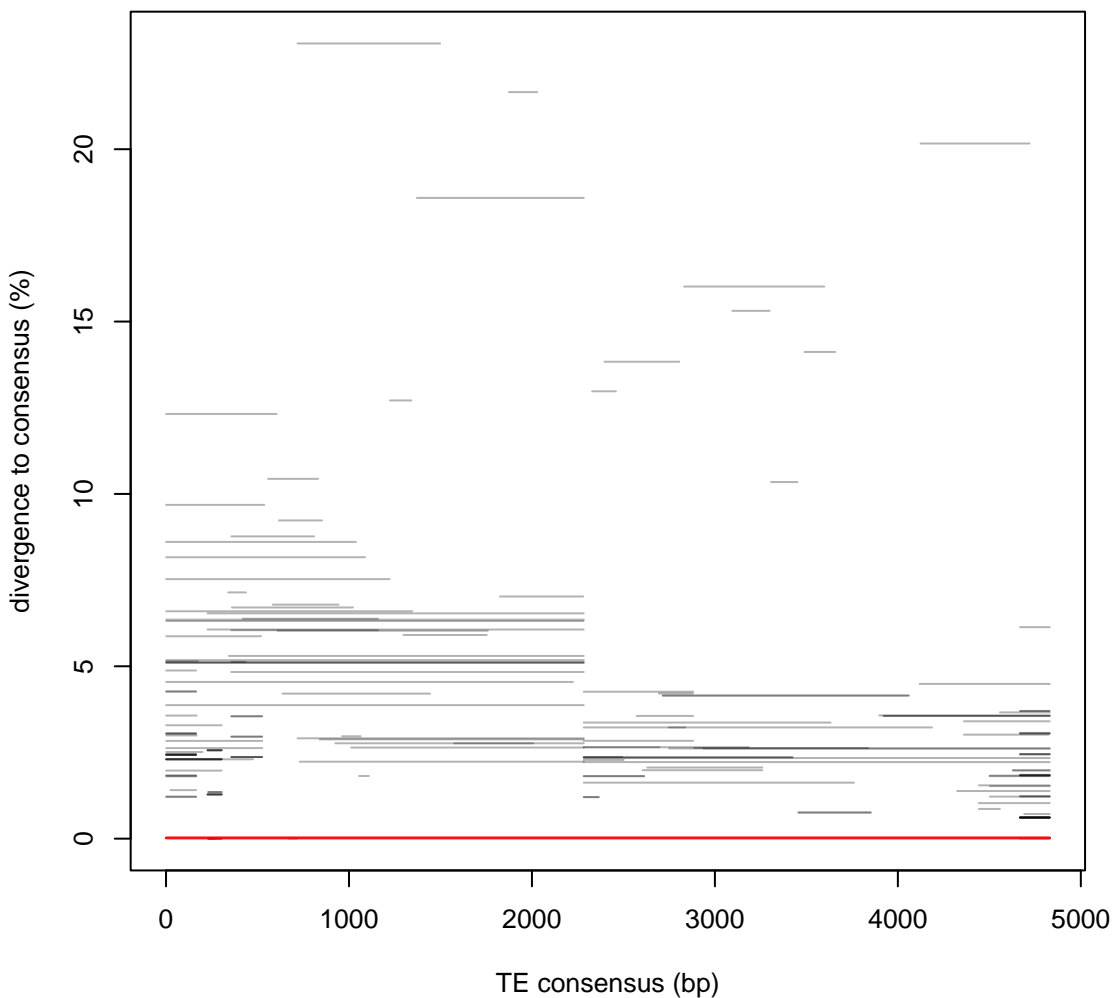
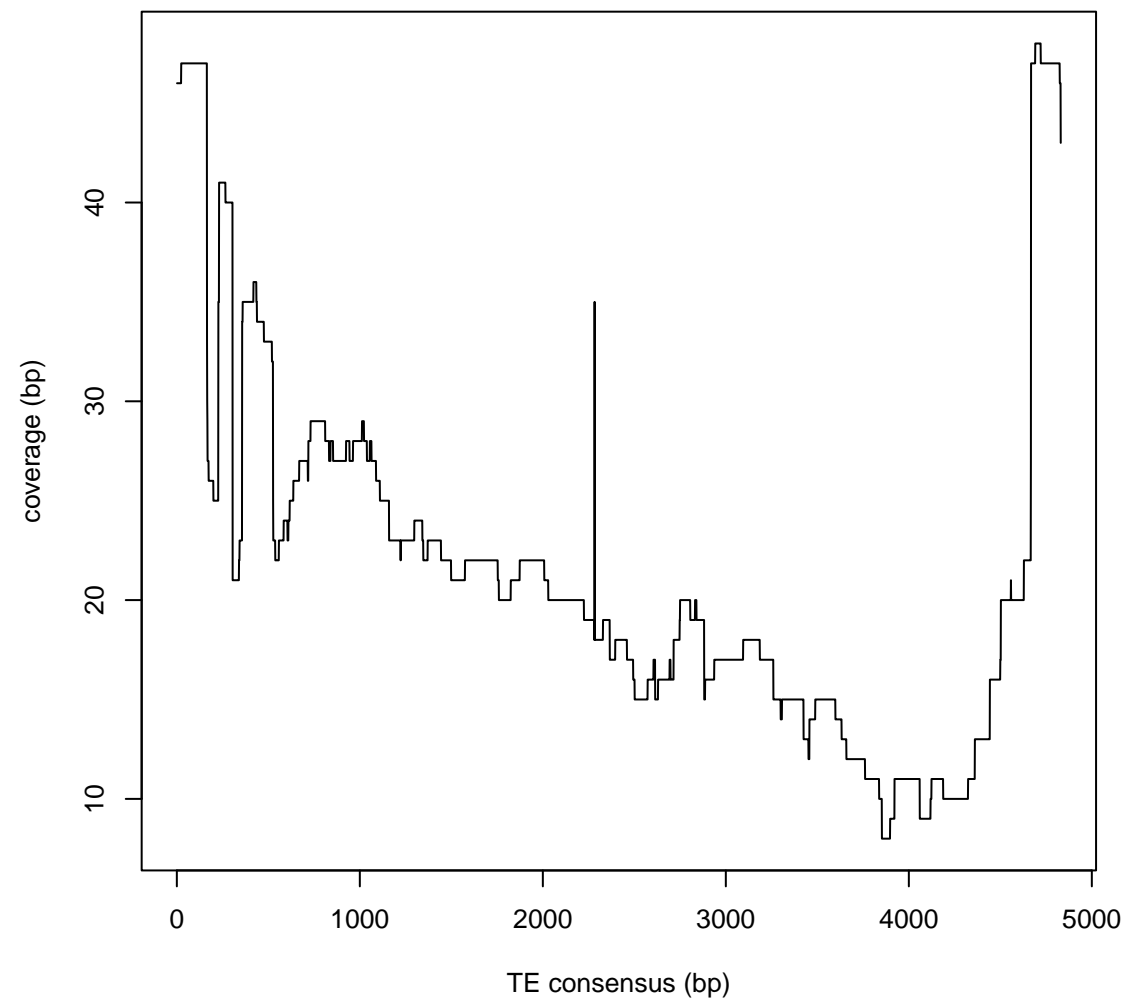


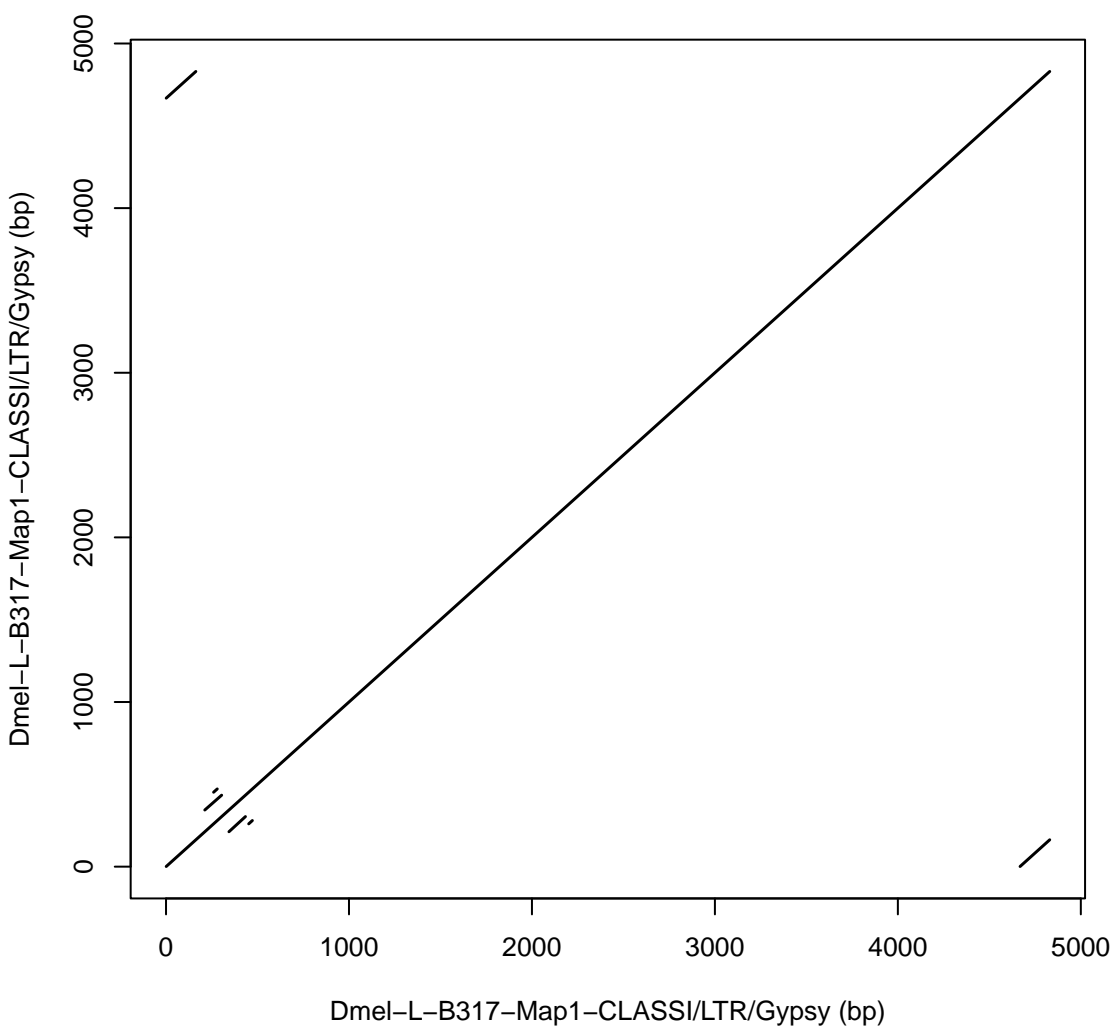
TE: Dmel-L-B317-Map1-CLASSI/LTR/Gypsy  
consensus size: 4830bp; fragments: 177; full length: 1 (>=4347bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

