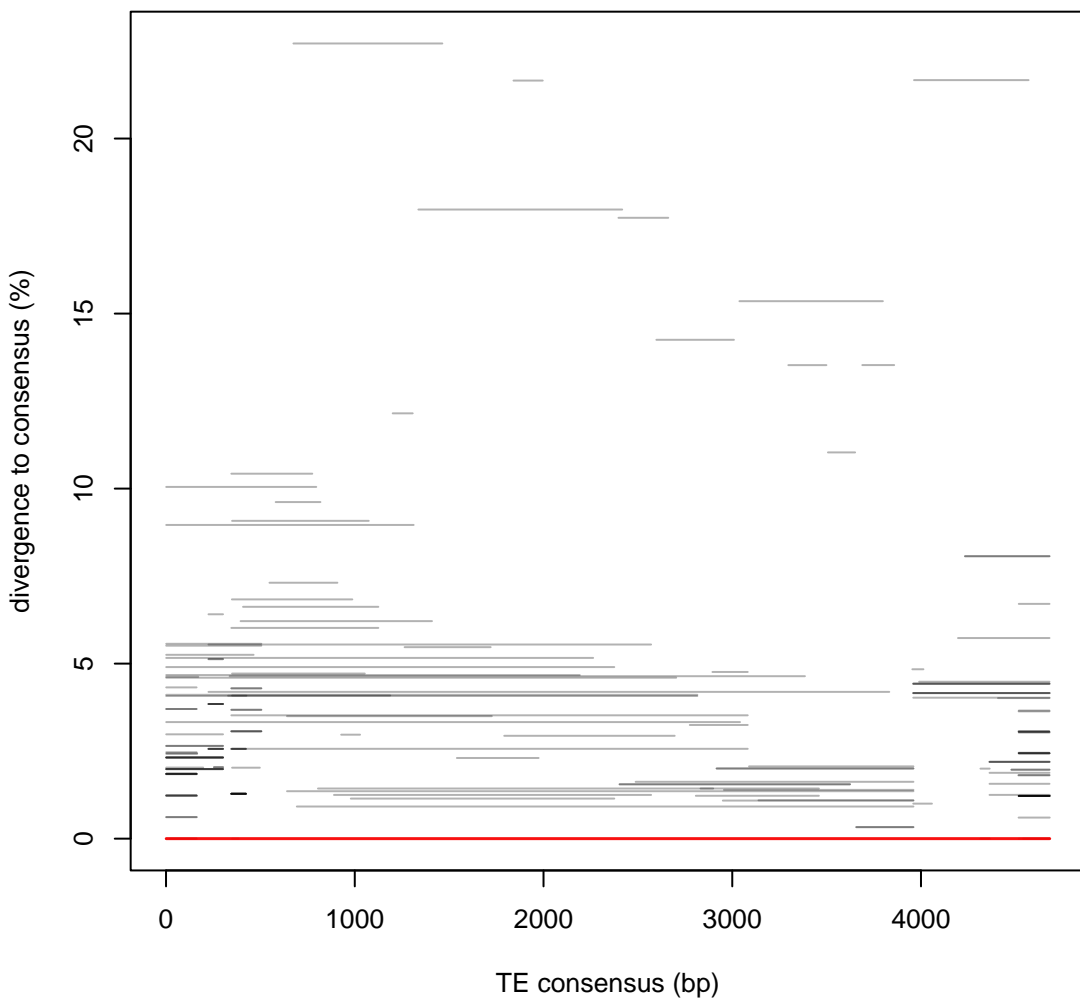
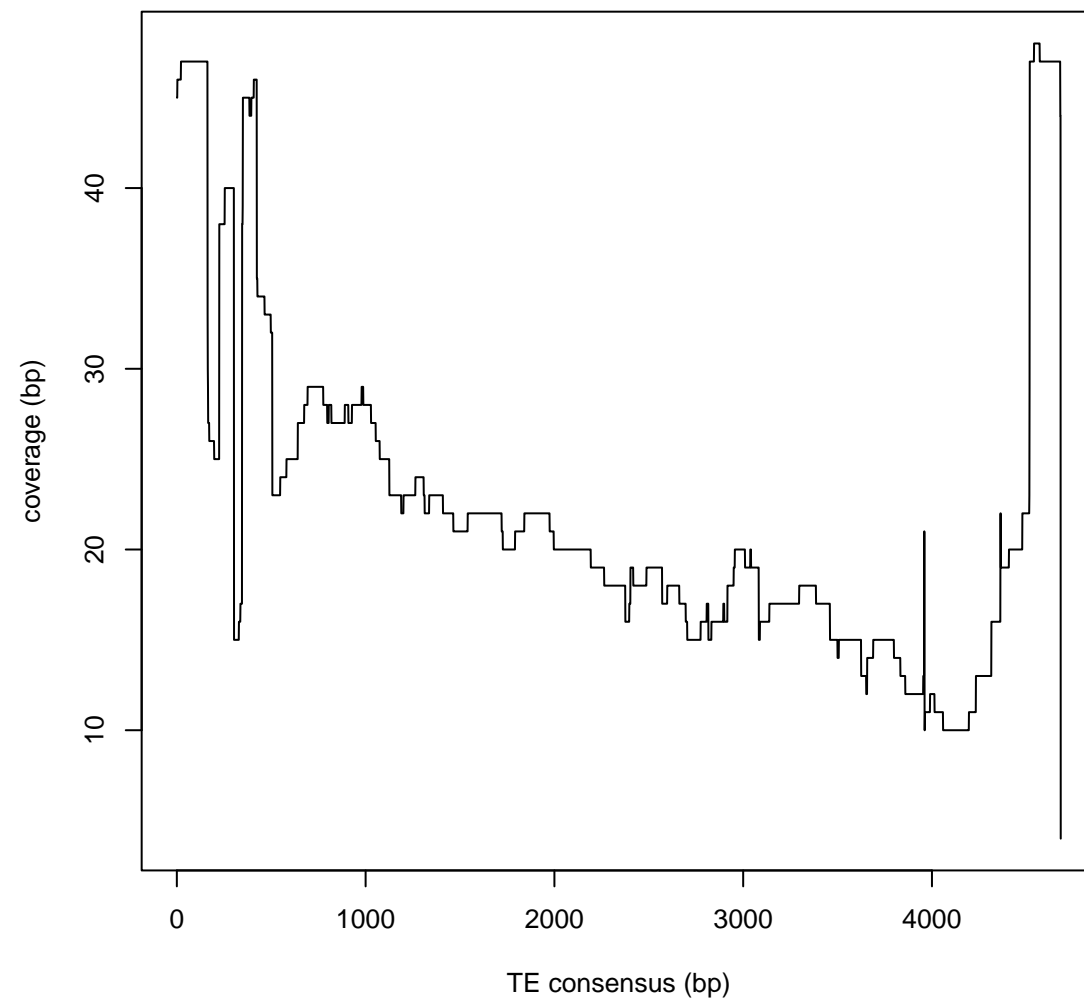


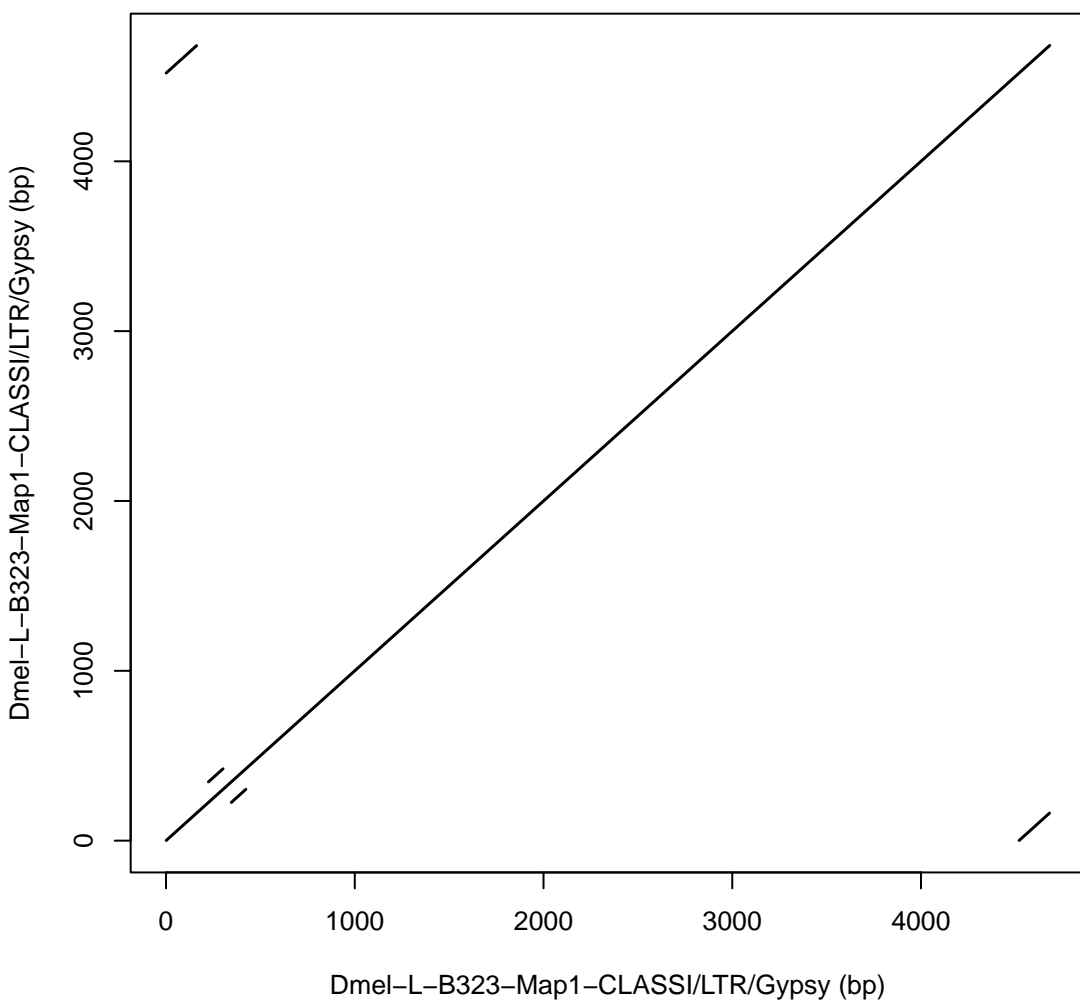
TE: Dmel-L-B323-Map1-CLASSI/LTR/Gypsy  
consensus size: 4682bp; fragments: 183; full length: 1 (>=4213.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

