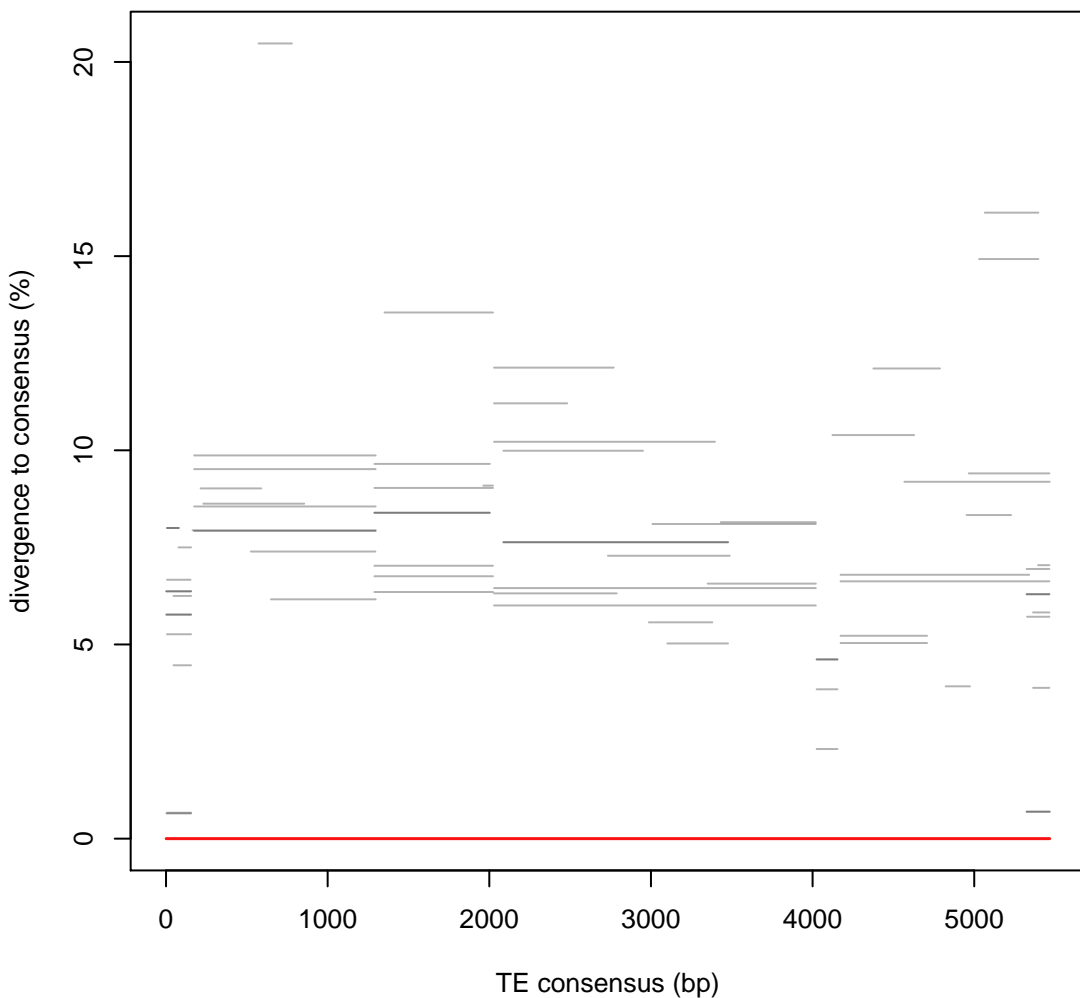
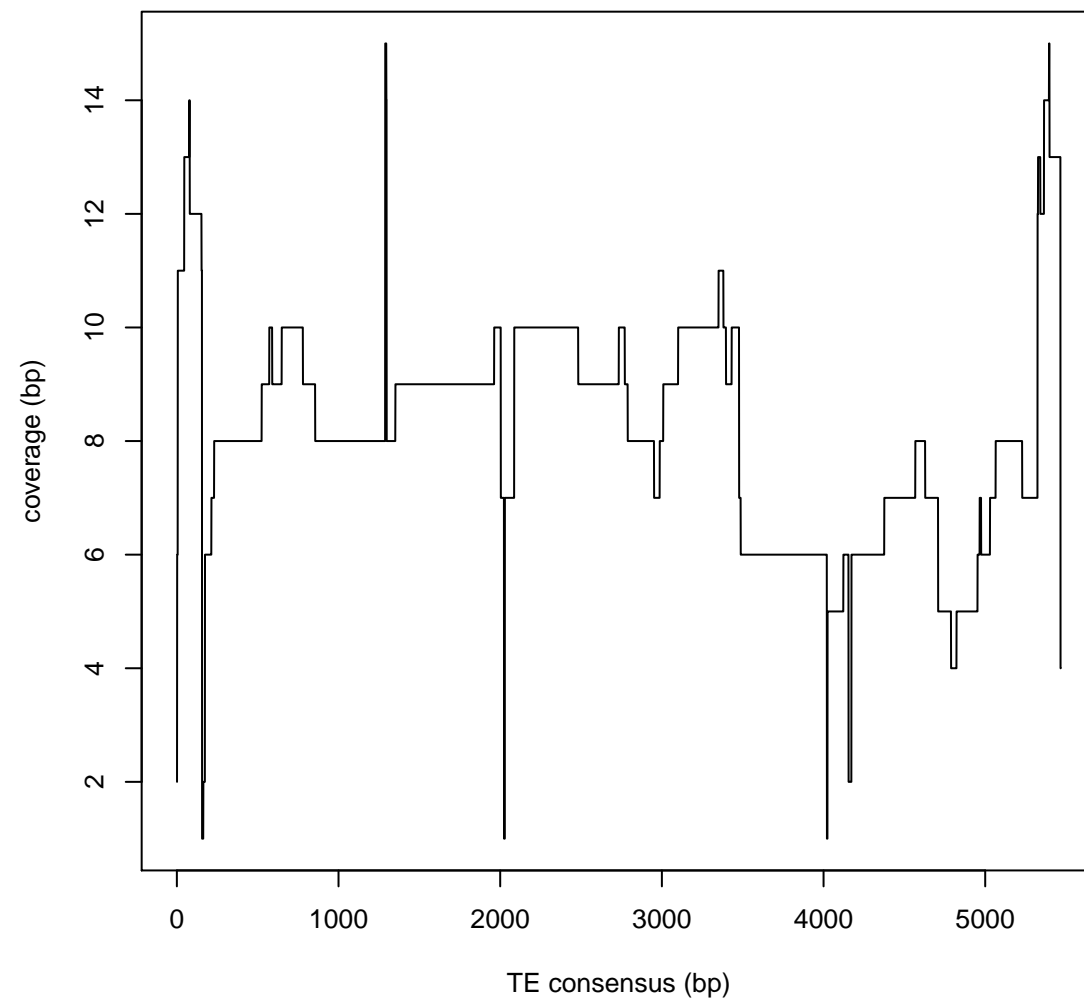


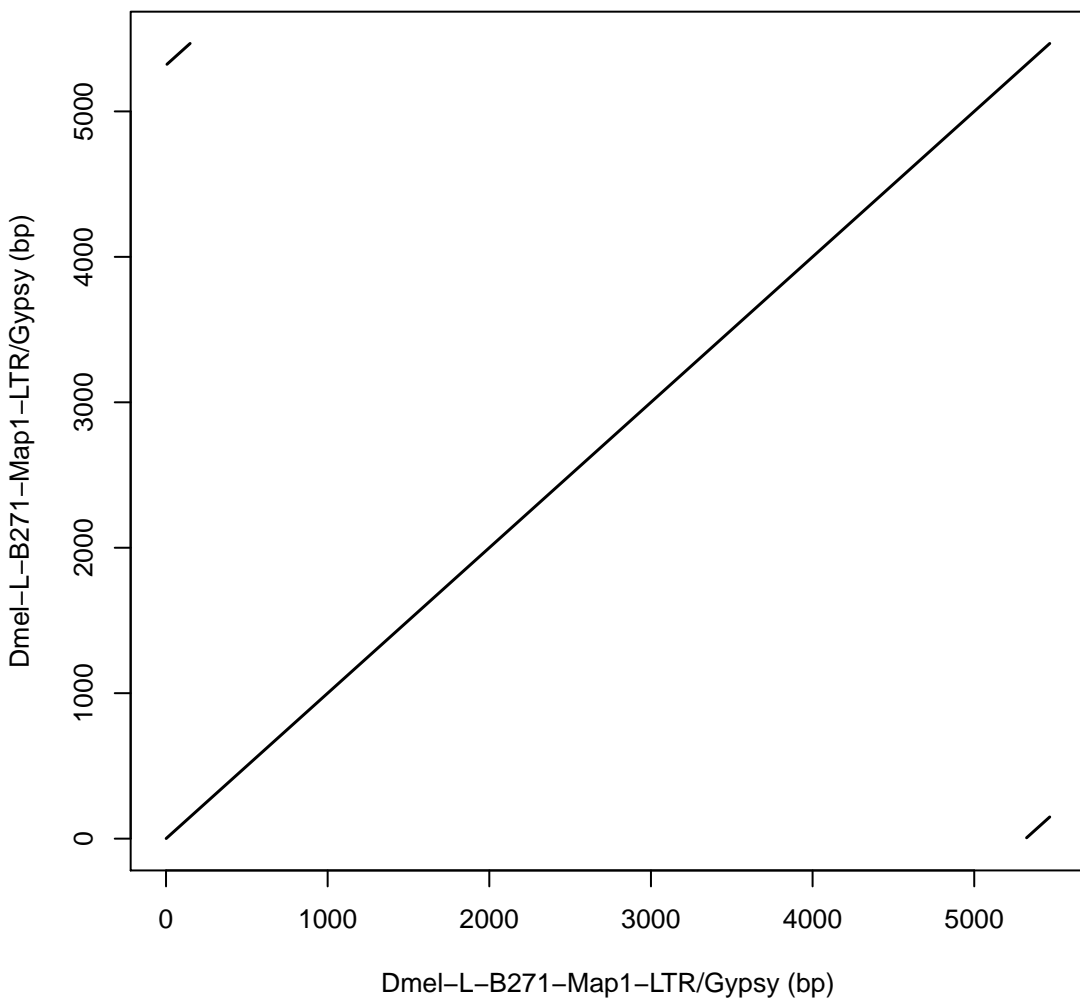
TE: Dmel-L-B271-Map1-LTR/Gypsy  
consensus size: 5467bp; fragments: 73; full length: 1 (>=4920.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

