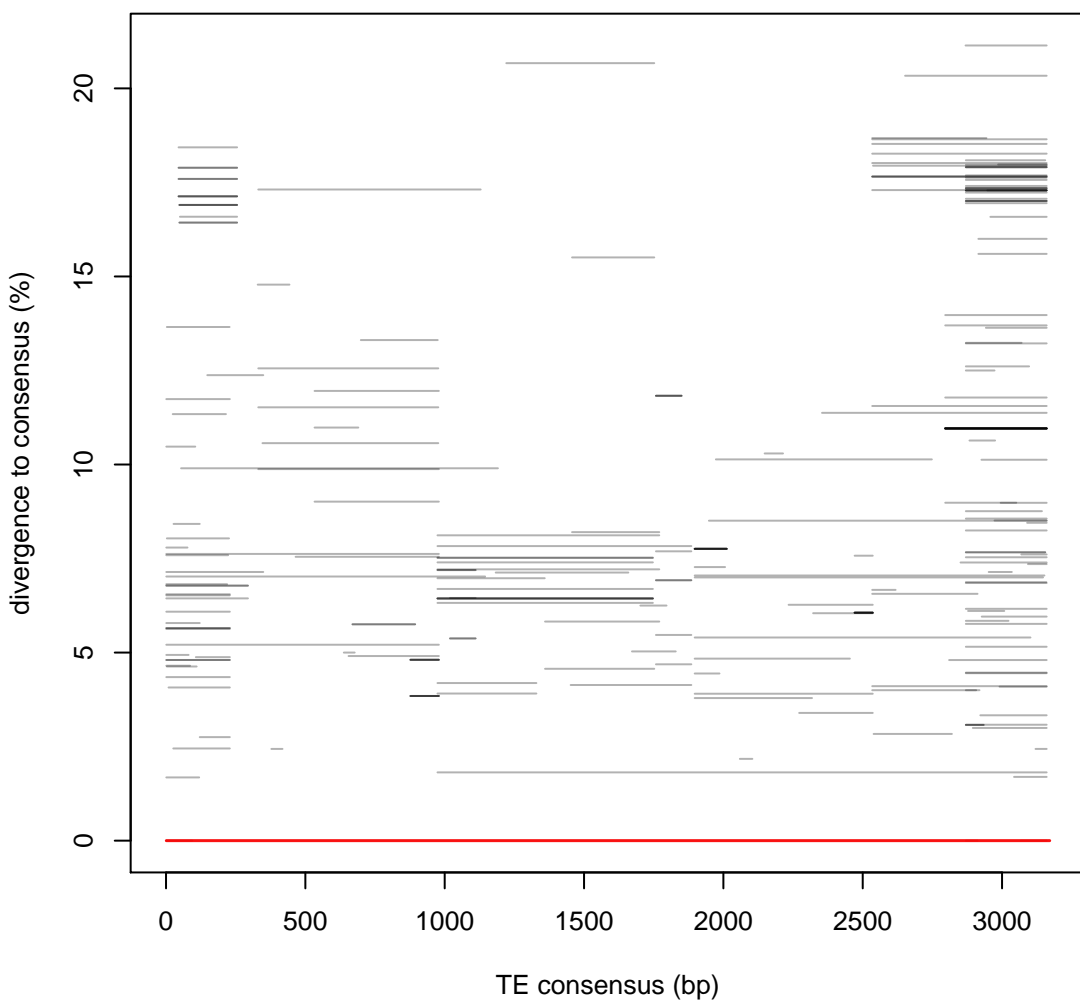
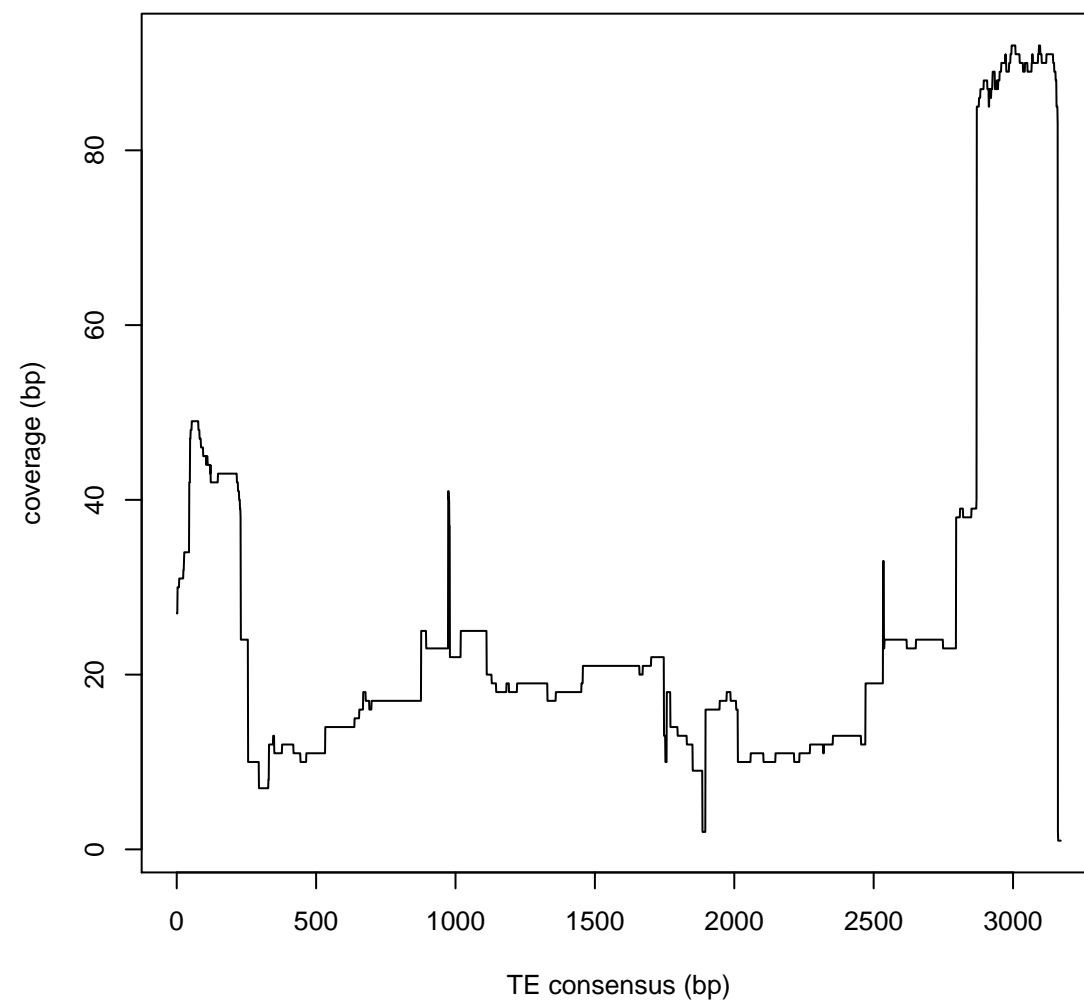


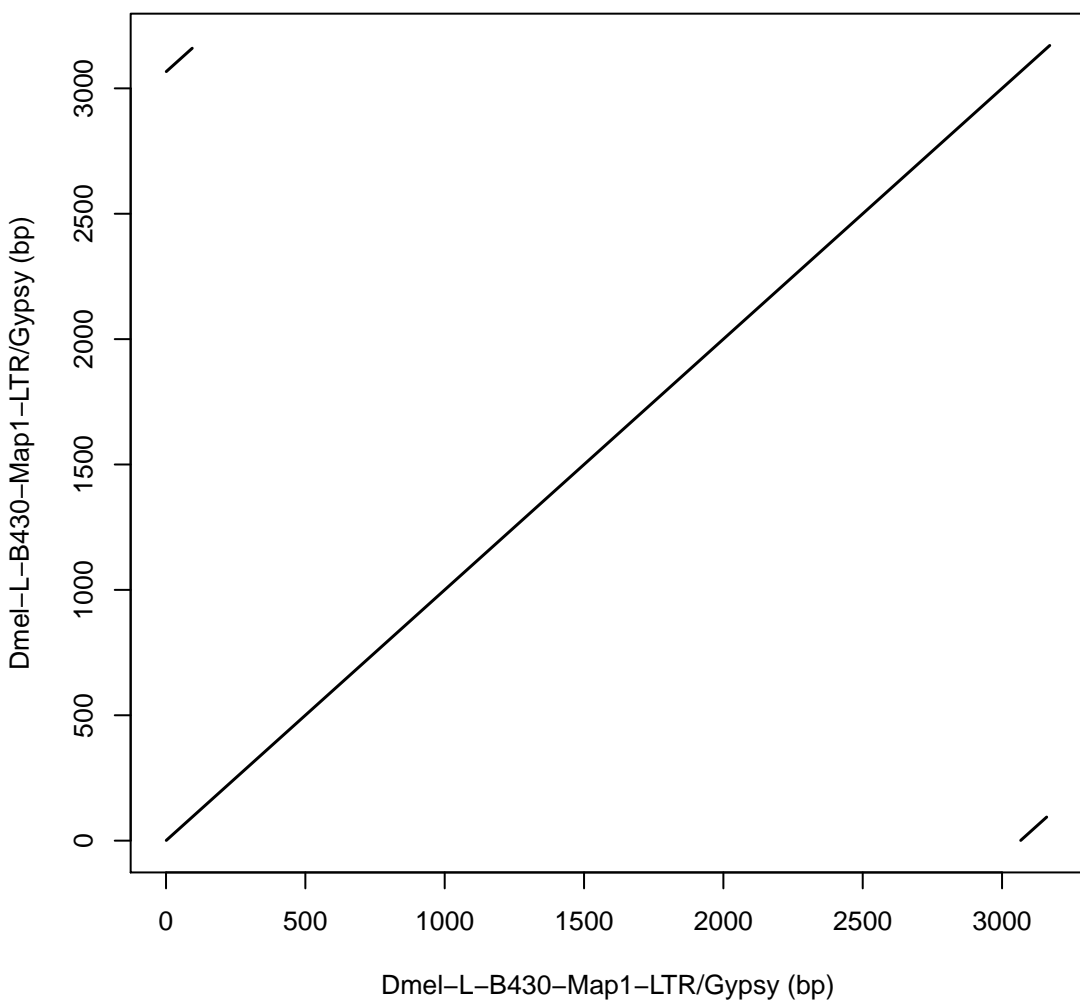
TE: Dmel-L-B430-Map1-LTR/Gypsy  
consensus size: 3171bp; fragments: 245; full length: 1 (>=2853.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

