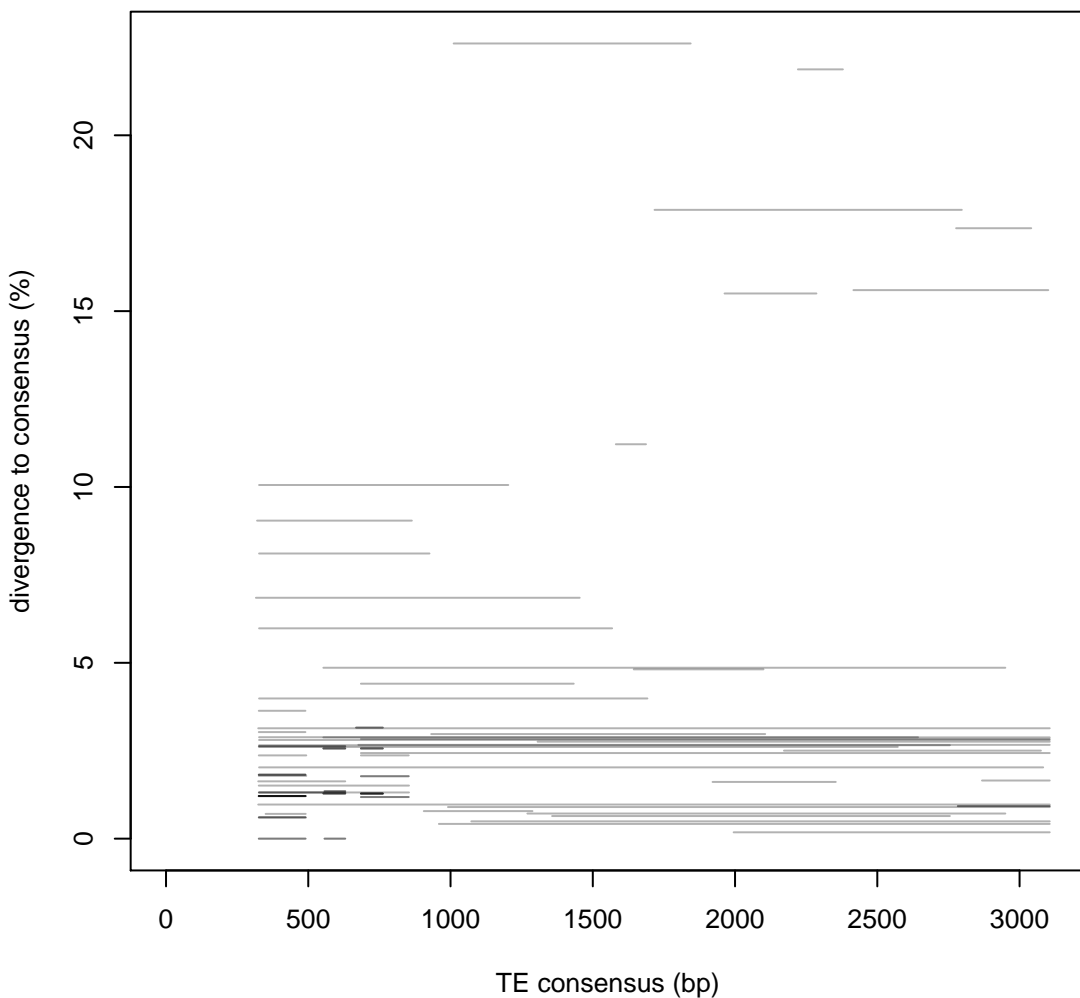
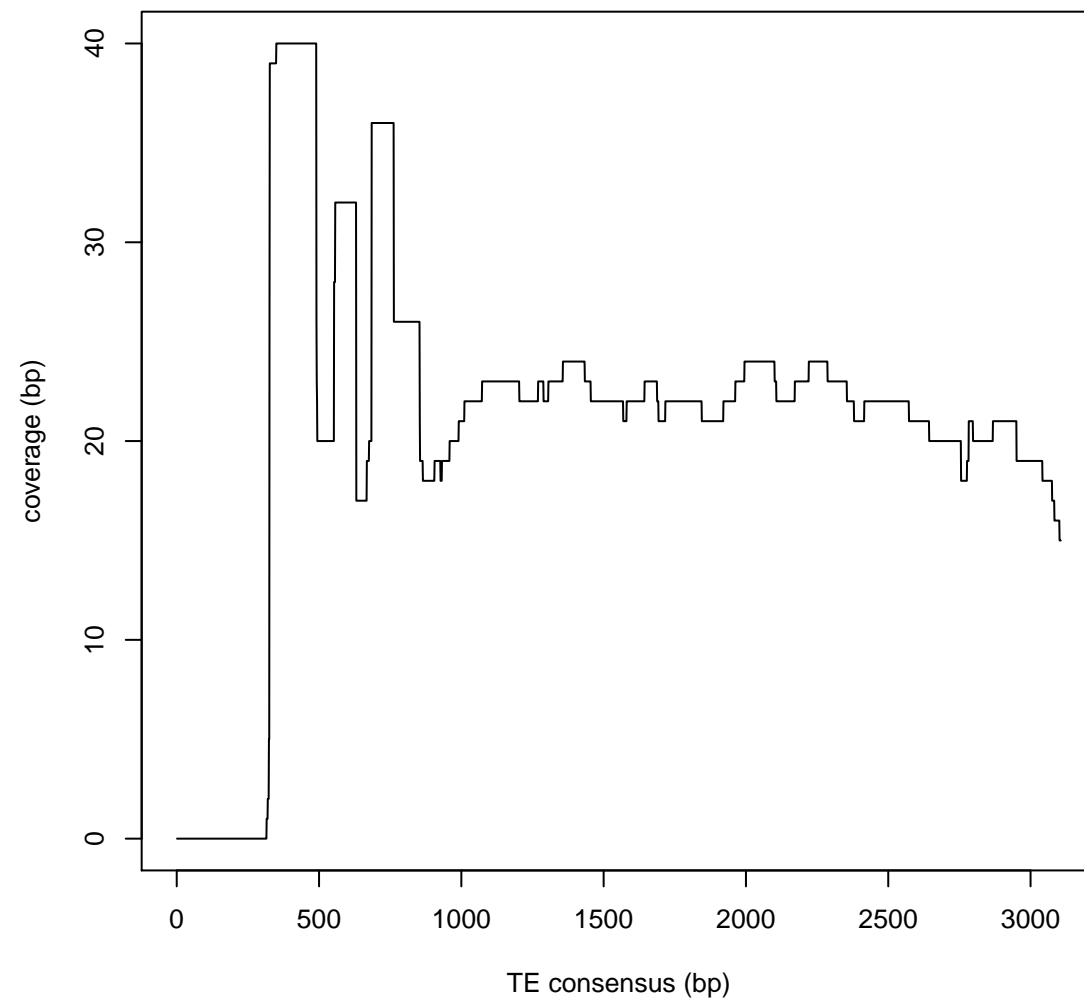


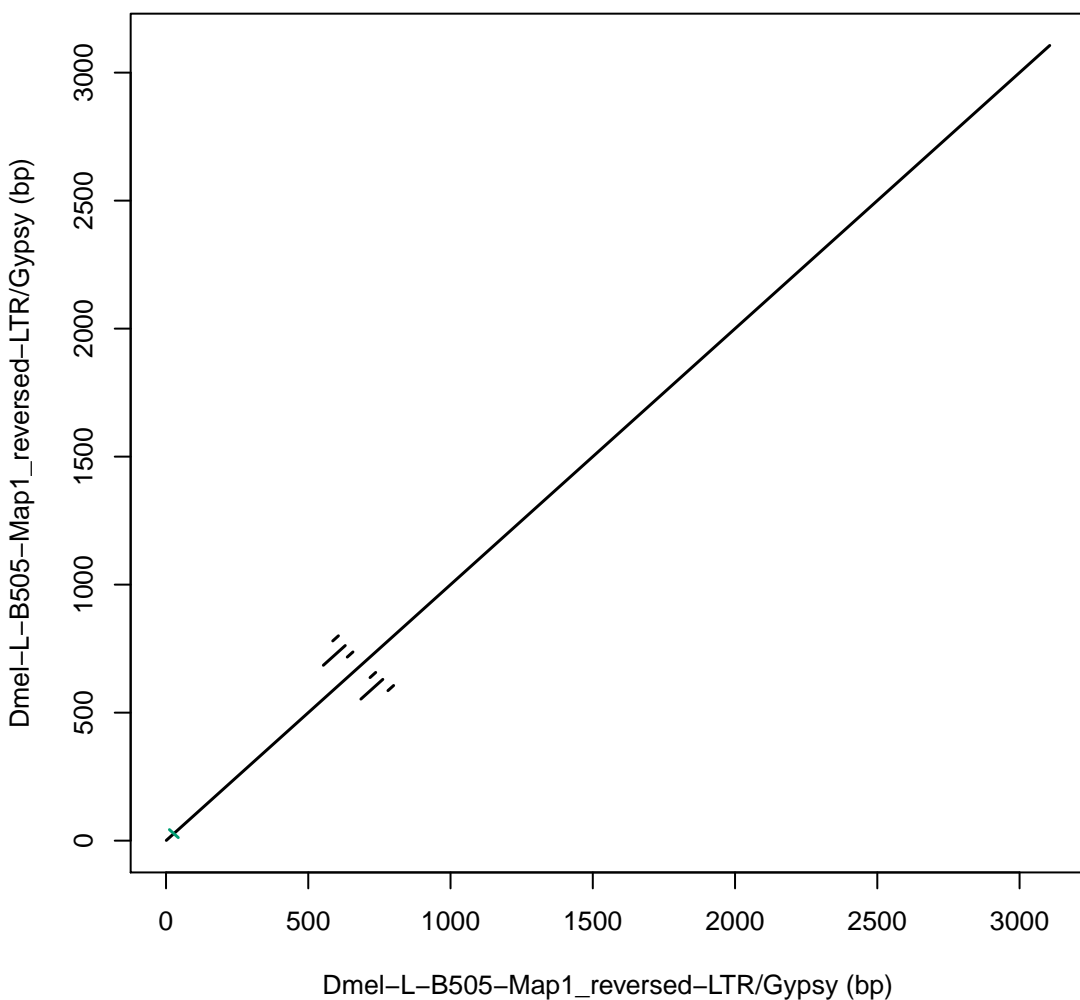
TE: Dmel-L-B505-Map1\_reversed-LTR/Gypsy  
consensus size: 3106bp; fragments: 93; full length: 0 (>=2795.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

