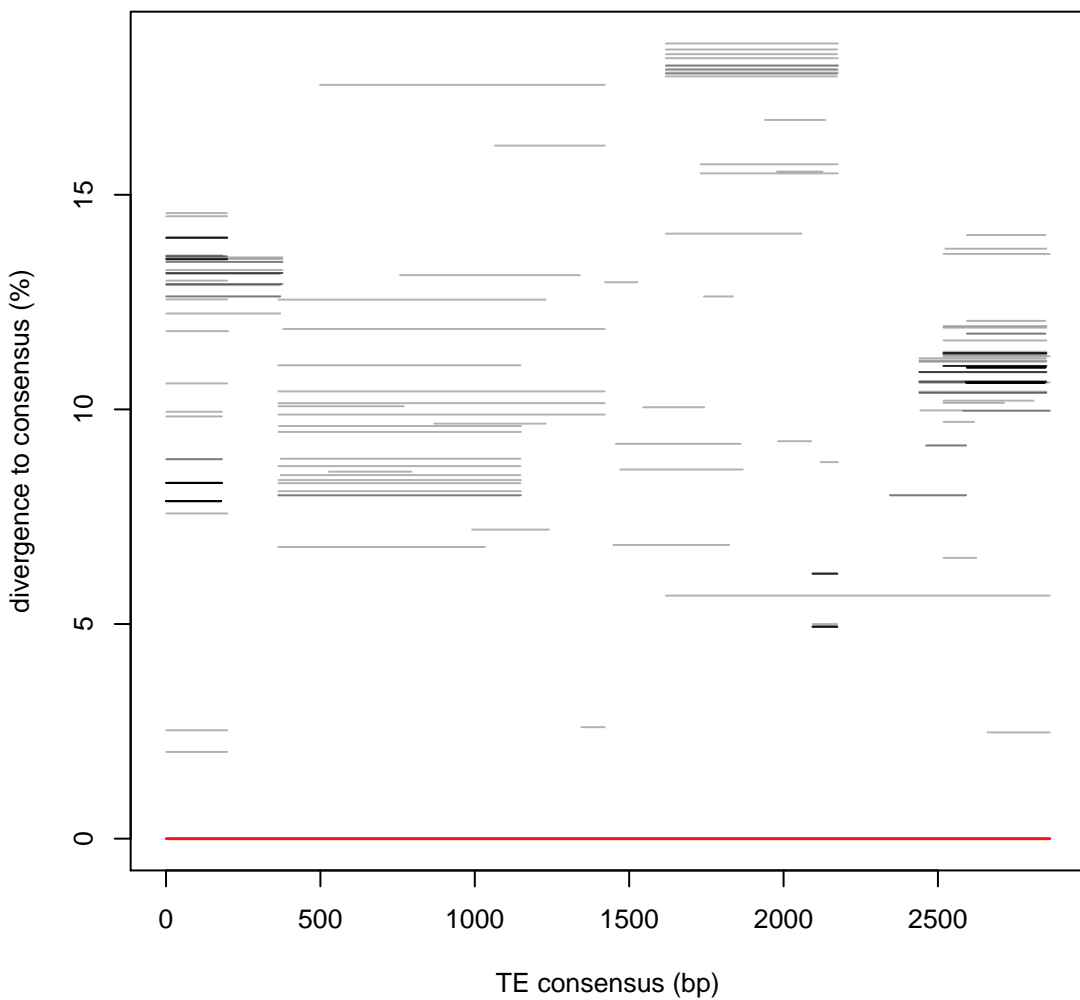
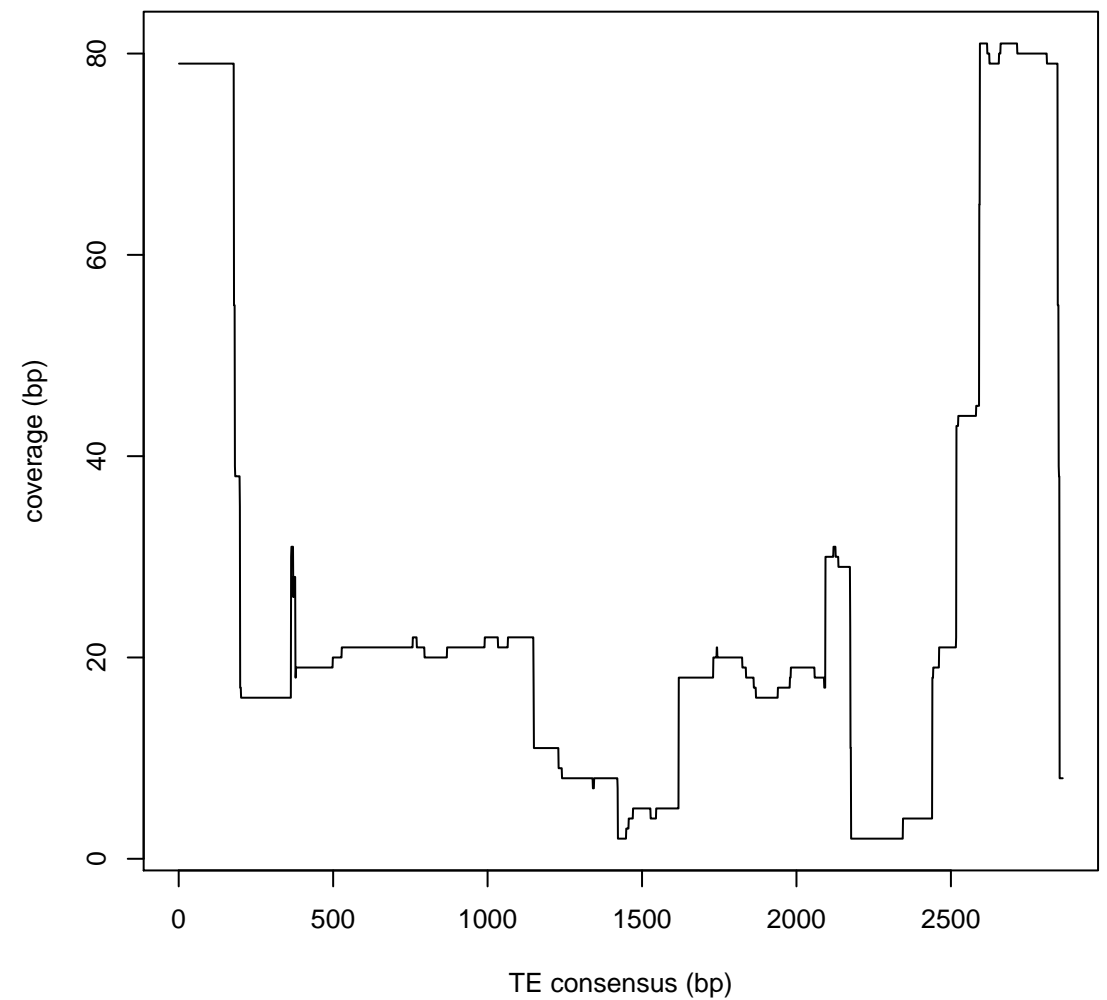


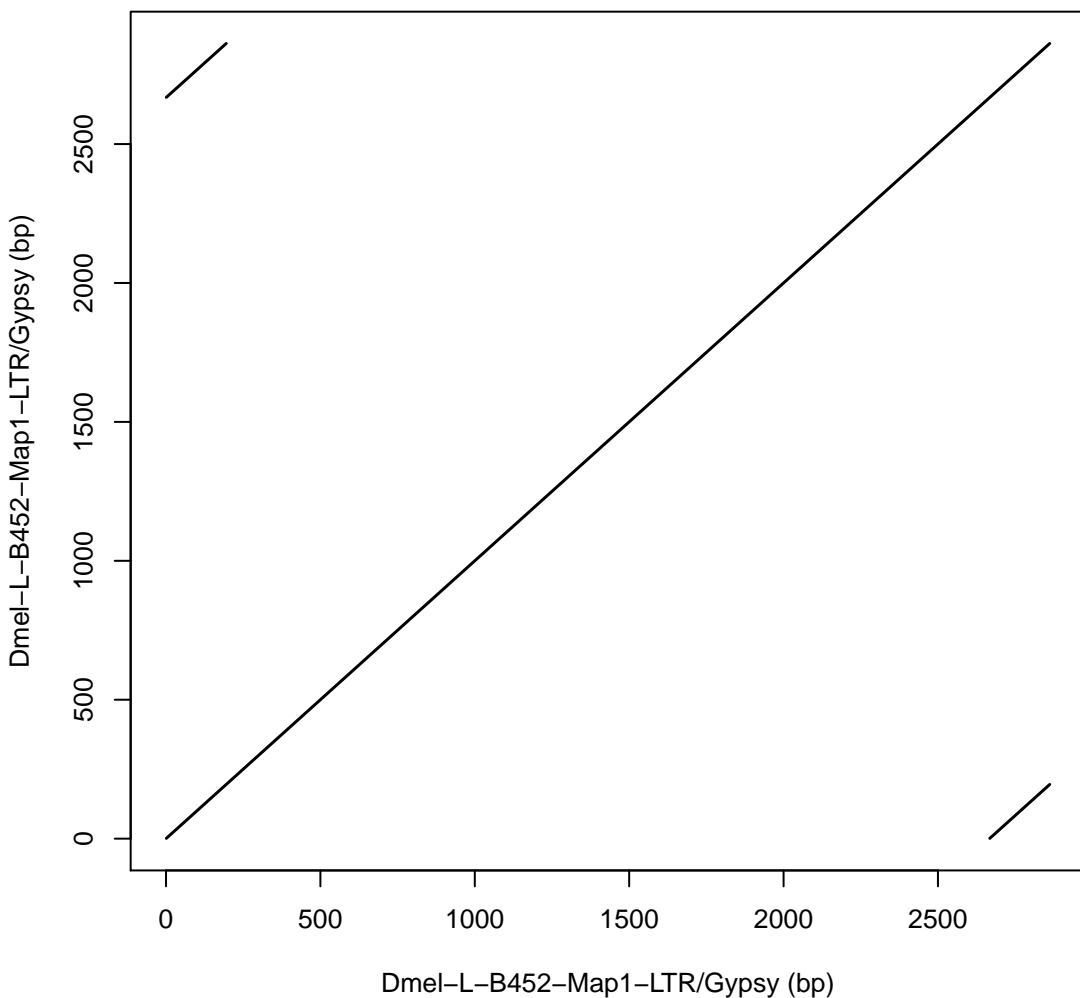
TE: Dmel-L-B452-Map1-LTR/Gypsy
consensus size: 2862bp; fragments: 227; full length: 1 (>=2575.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

