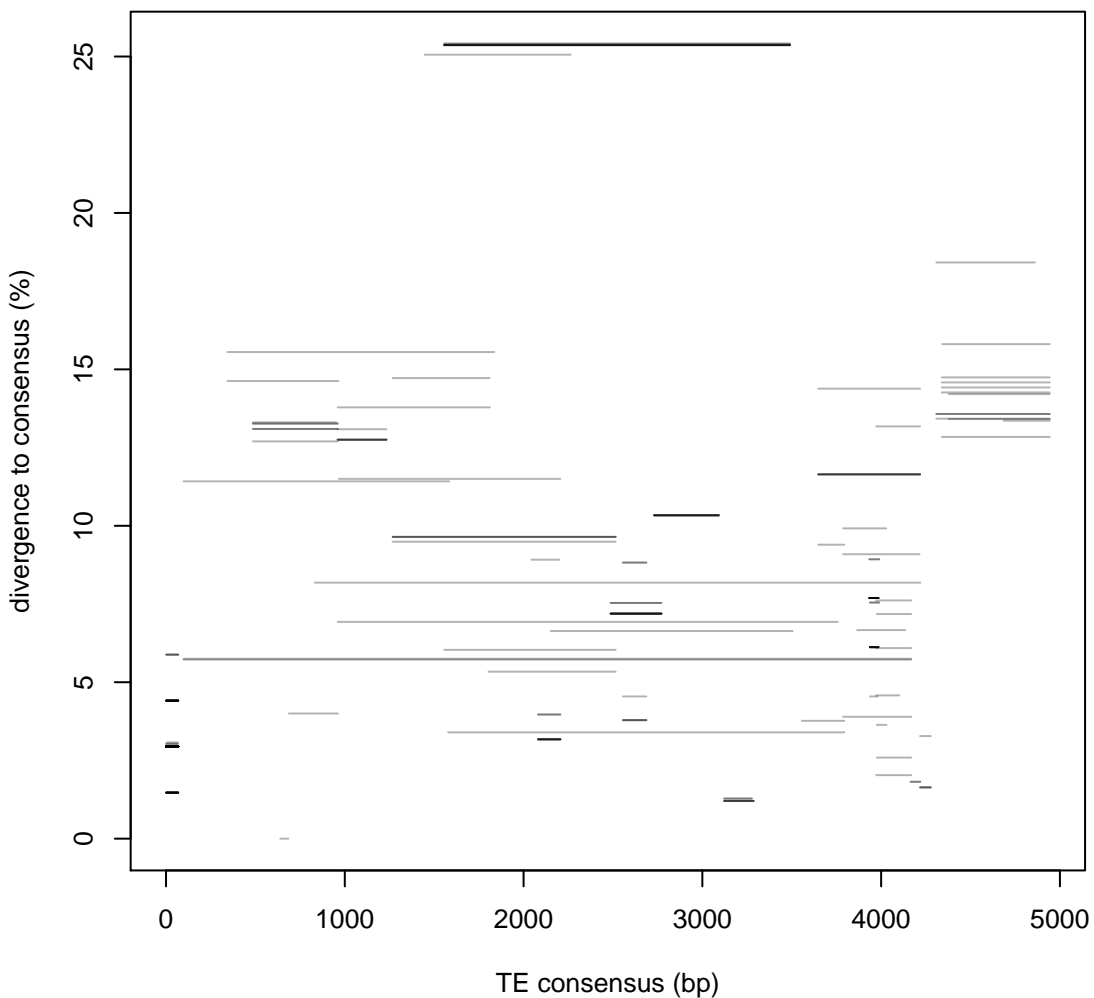
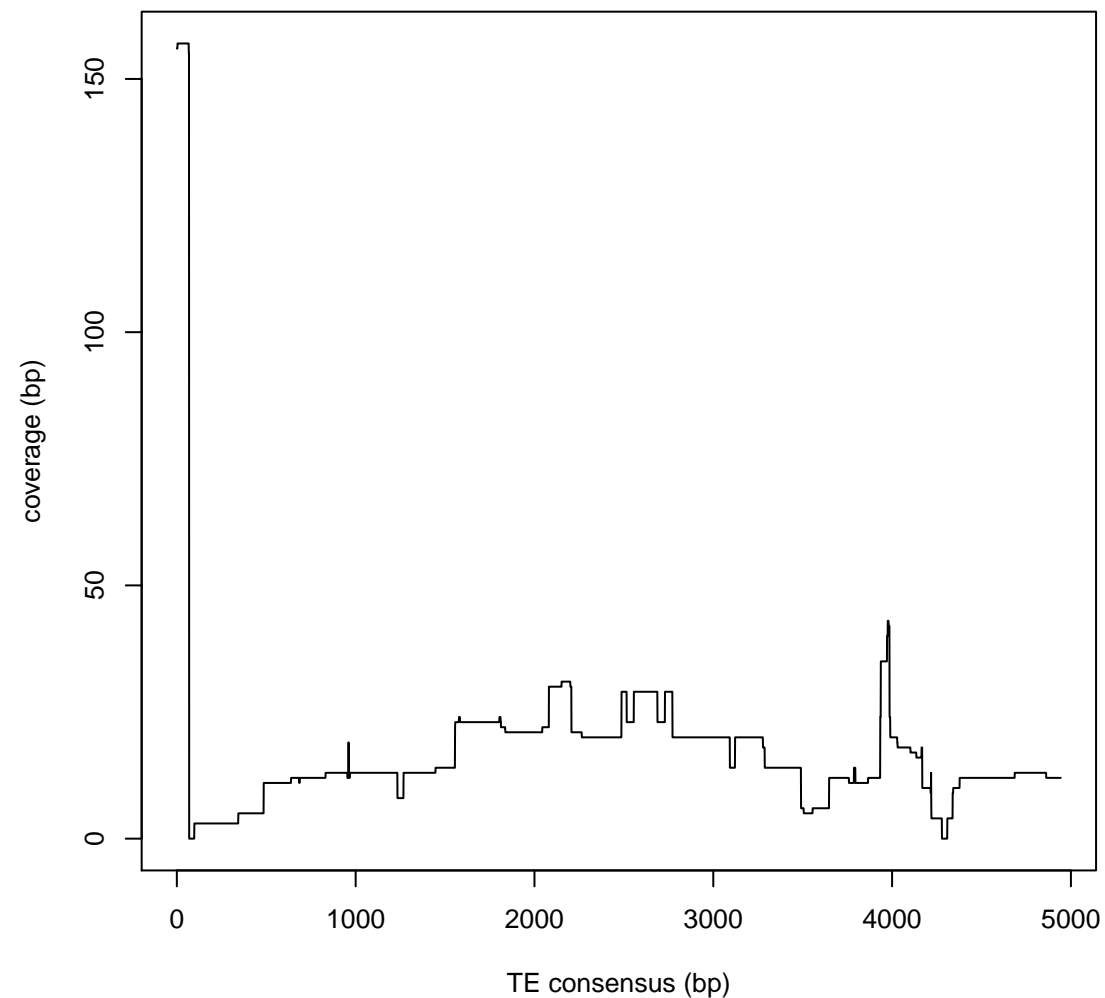


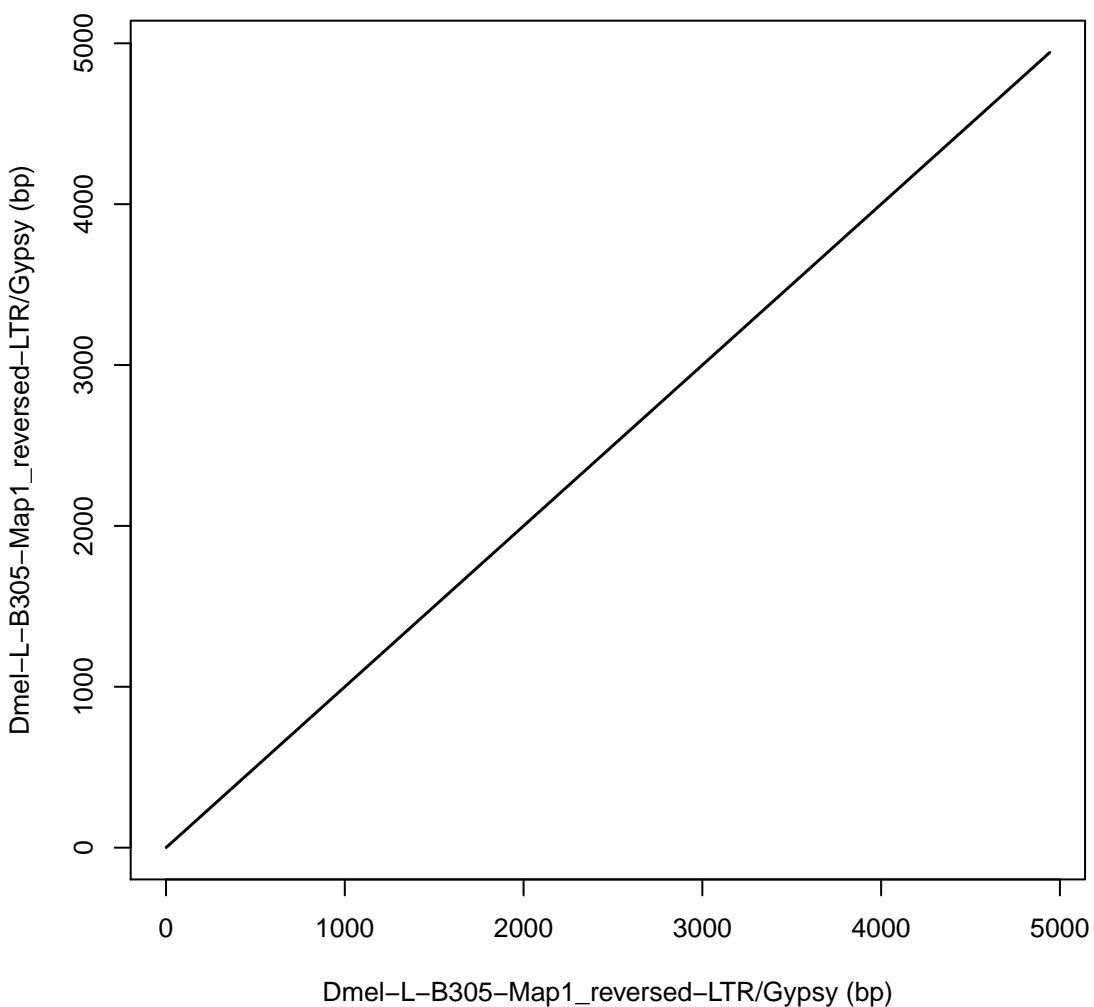
TE: Dmel-L-B305-Map1_reversed-LTR/Gypsy
consensus size: 4943bp; fragments: 294; full length: 0 (>=4448.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

