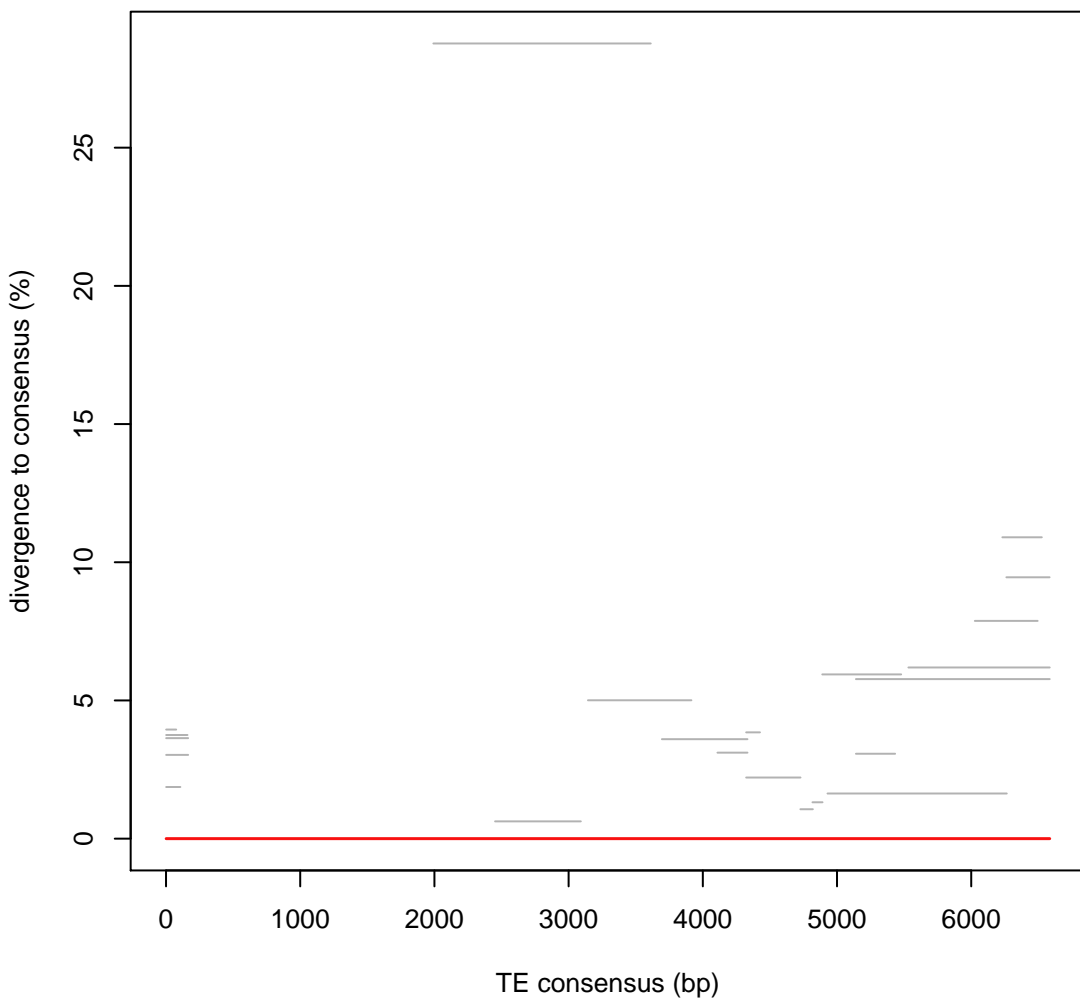
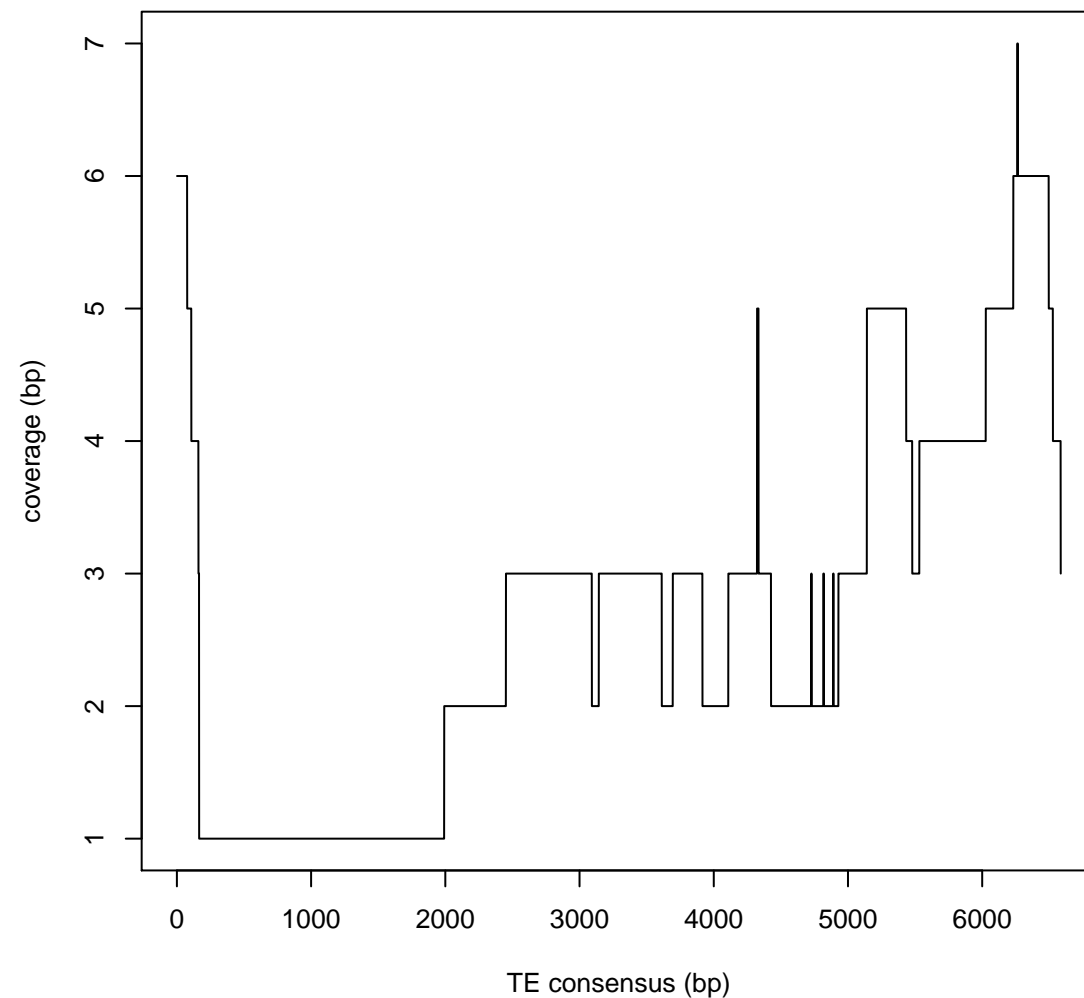


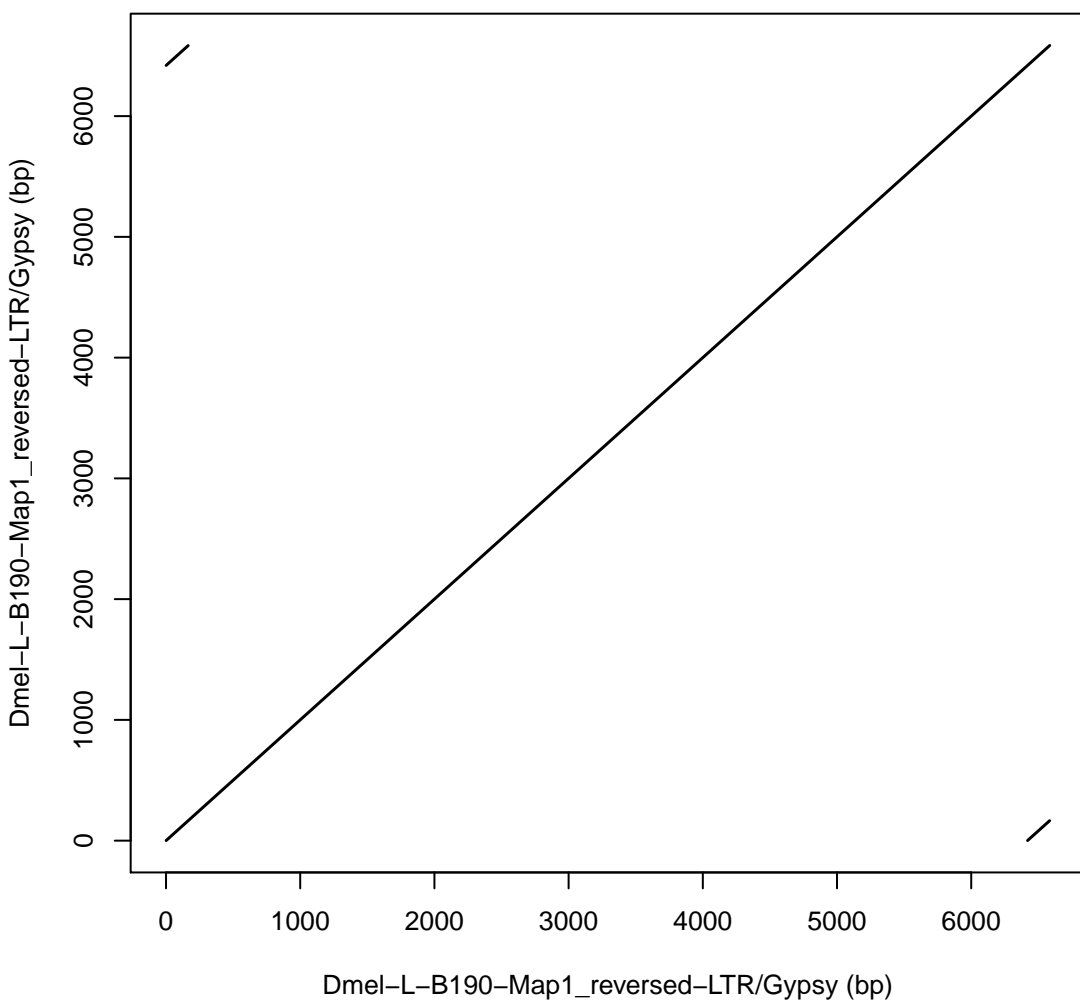
TE: Dmel-L-B190-Map1_reversed-LTR/Gypsy
consensus size: 6585bp; fragments: 23; full length: 1 (>=5926.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

