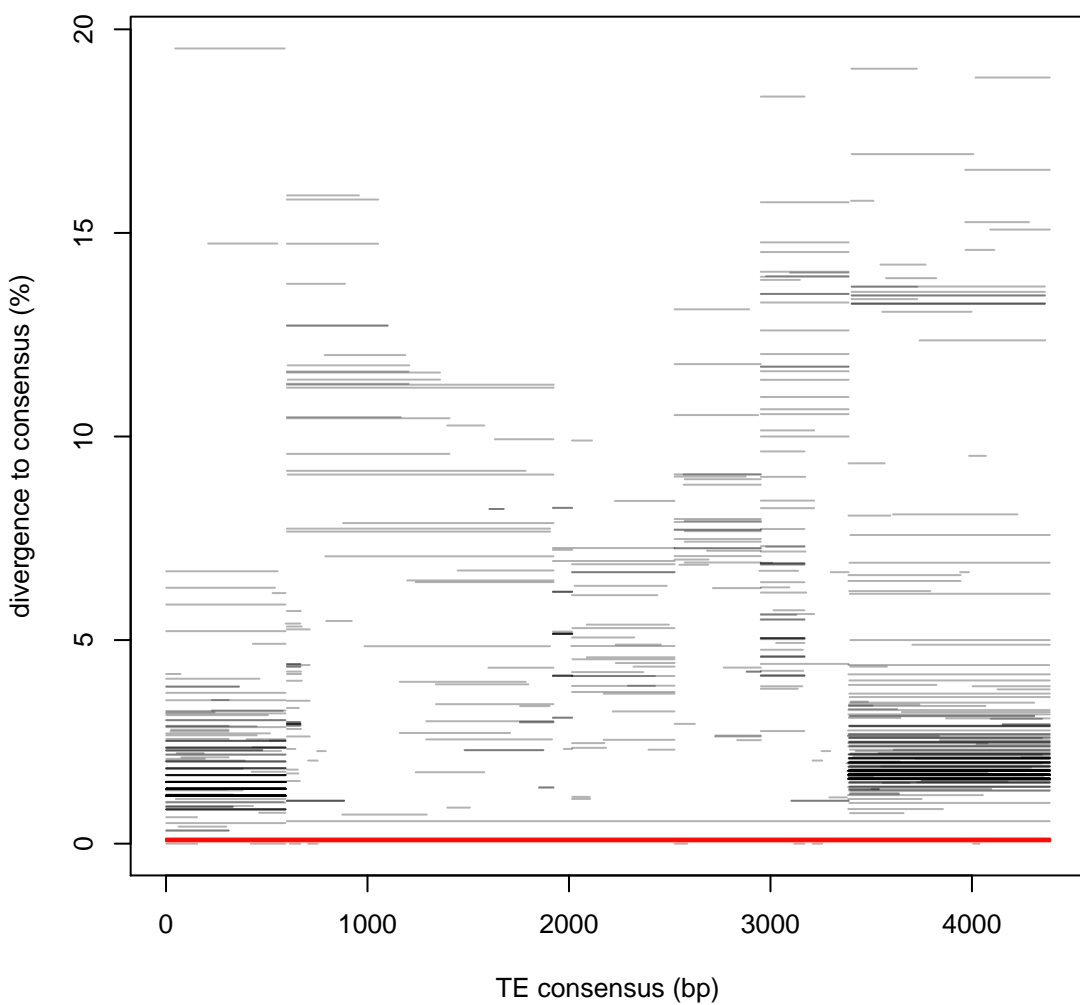
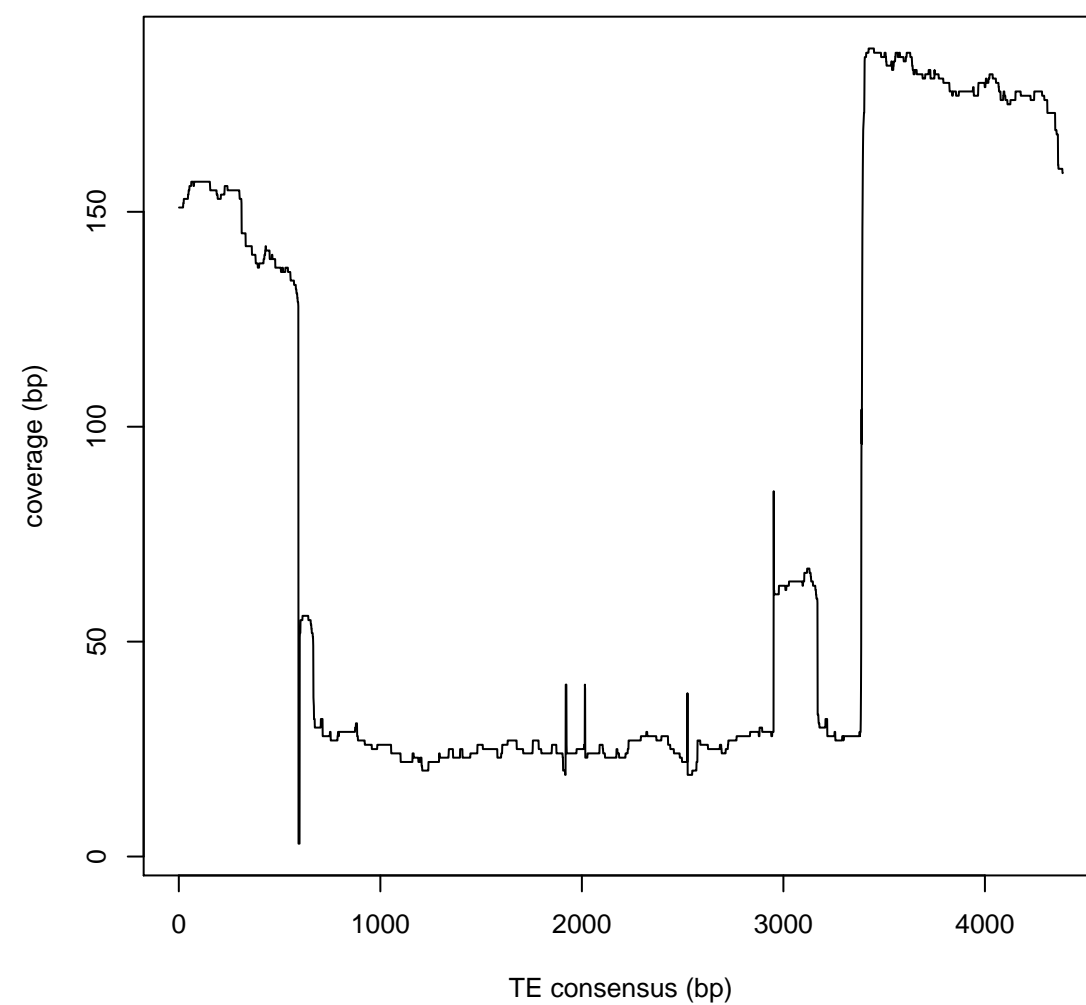


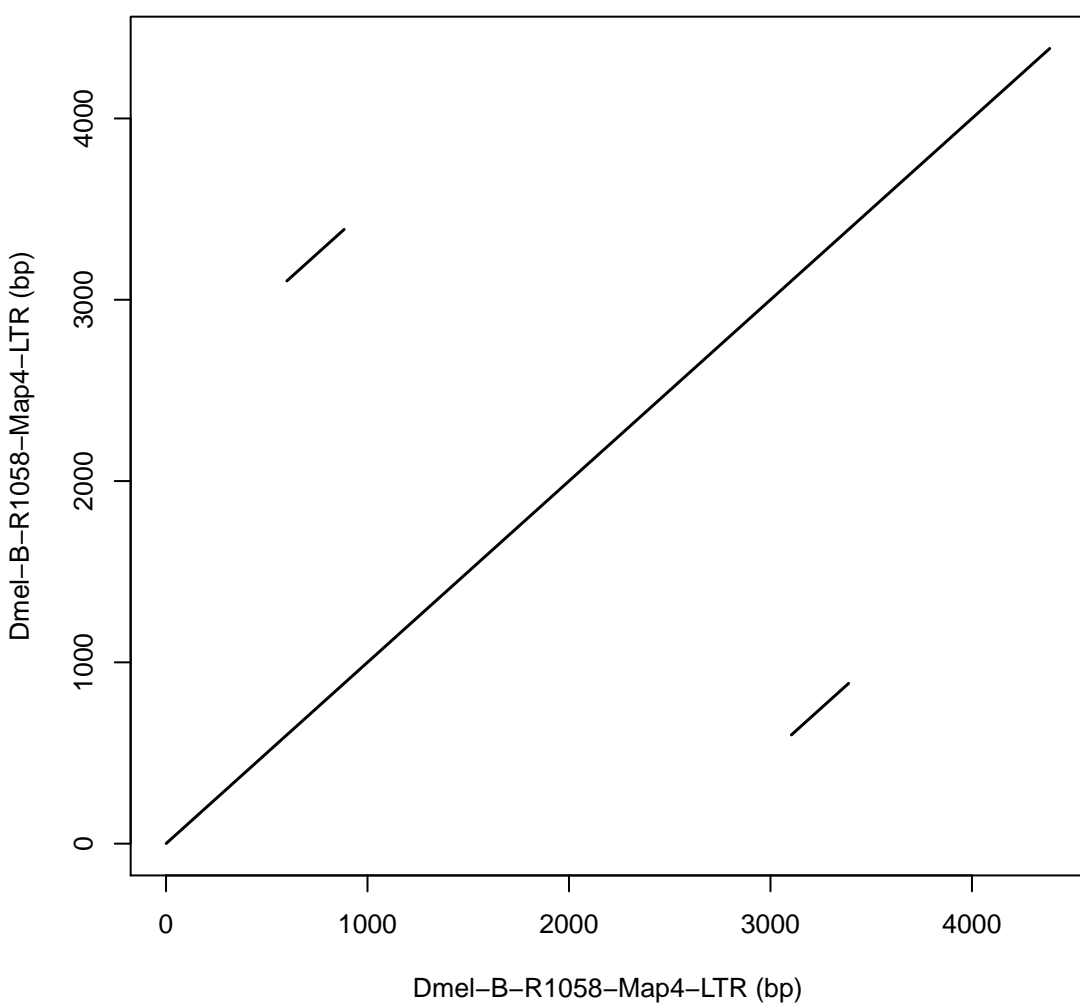
TE: Dmel-B-R1058-Map4-LTR
consensus size: 4386bp; fragments: 629; full length: 2 (≥ 3947.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

