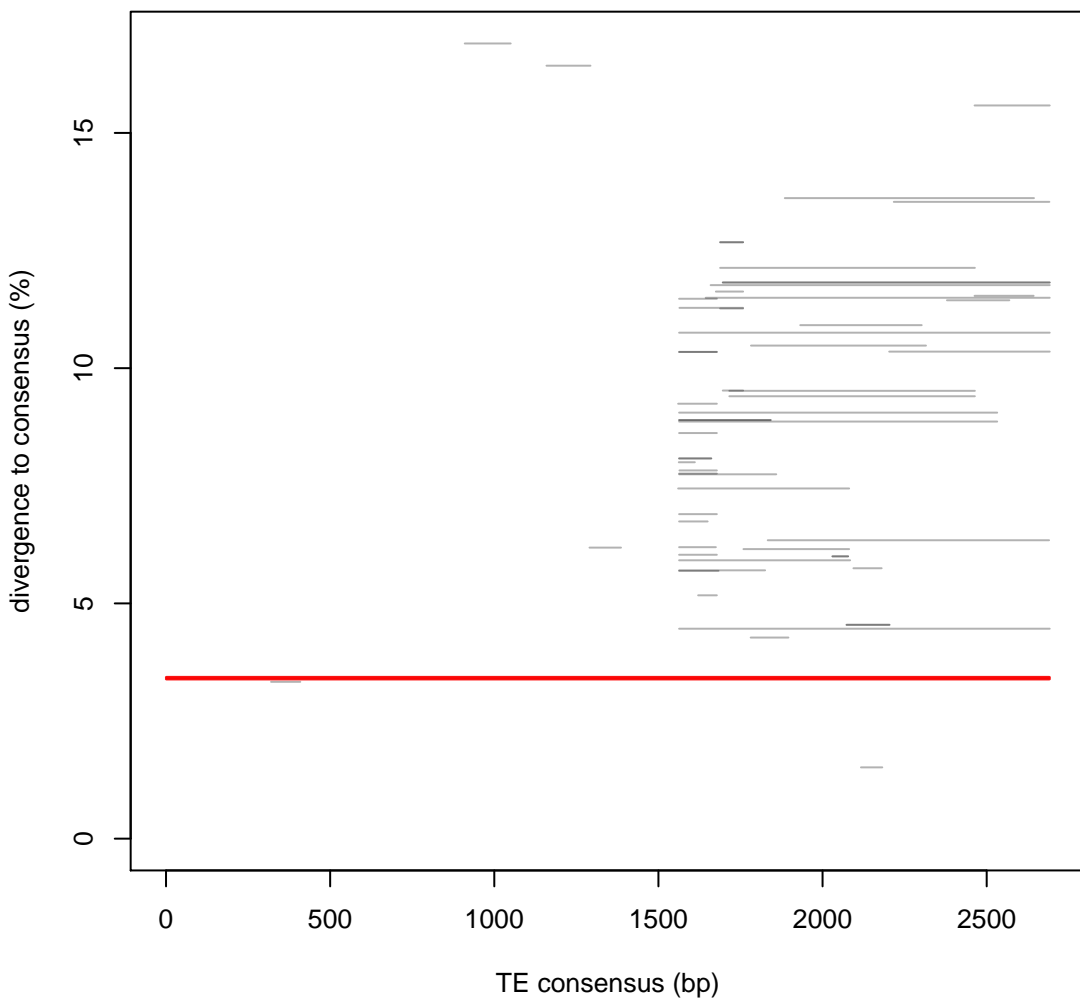
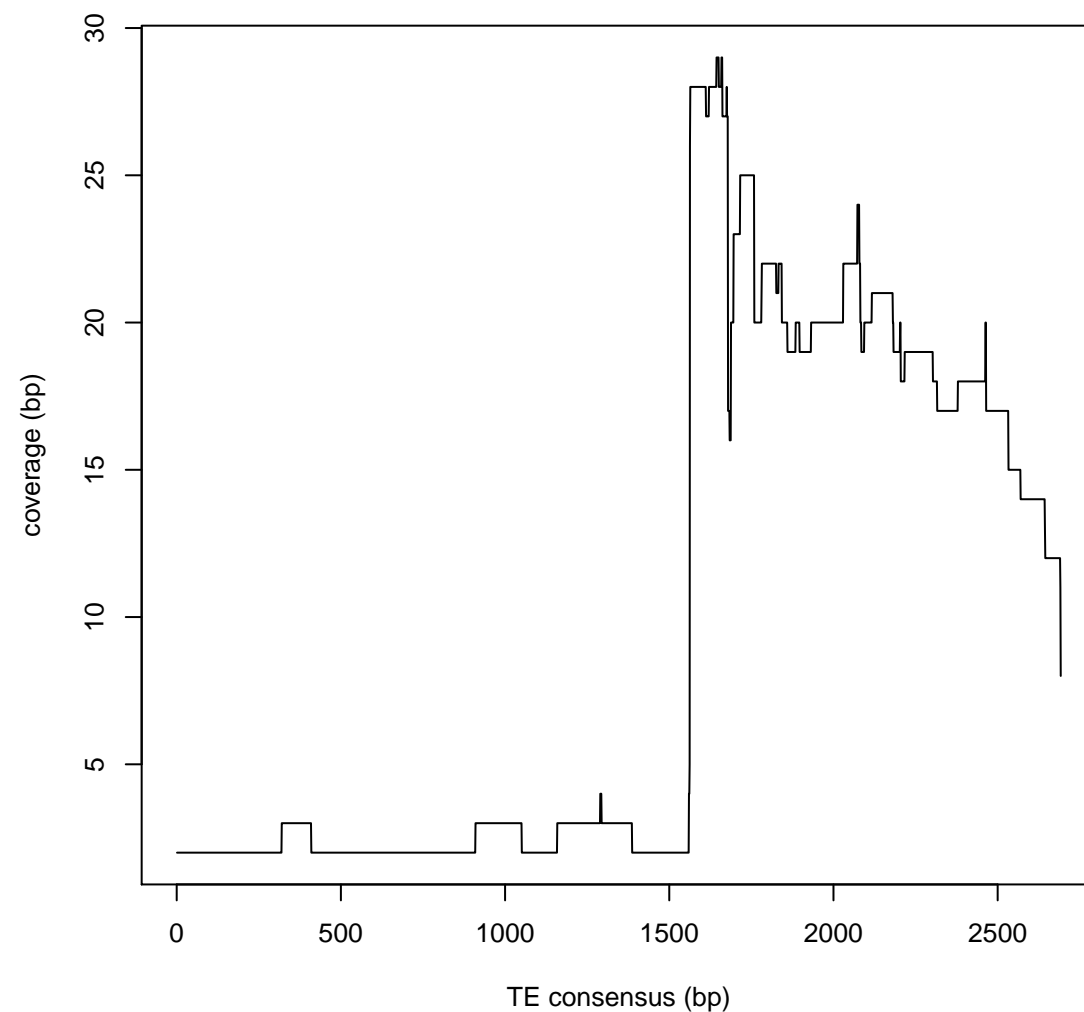


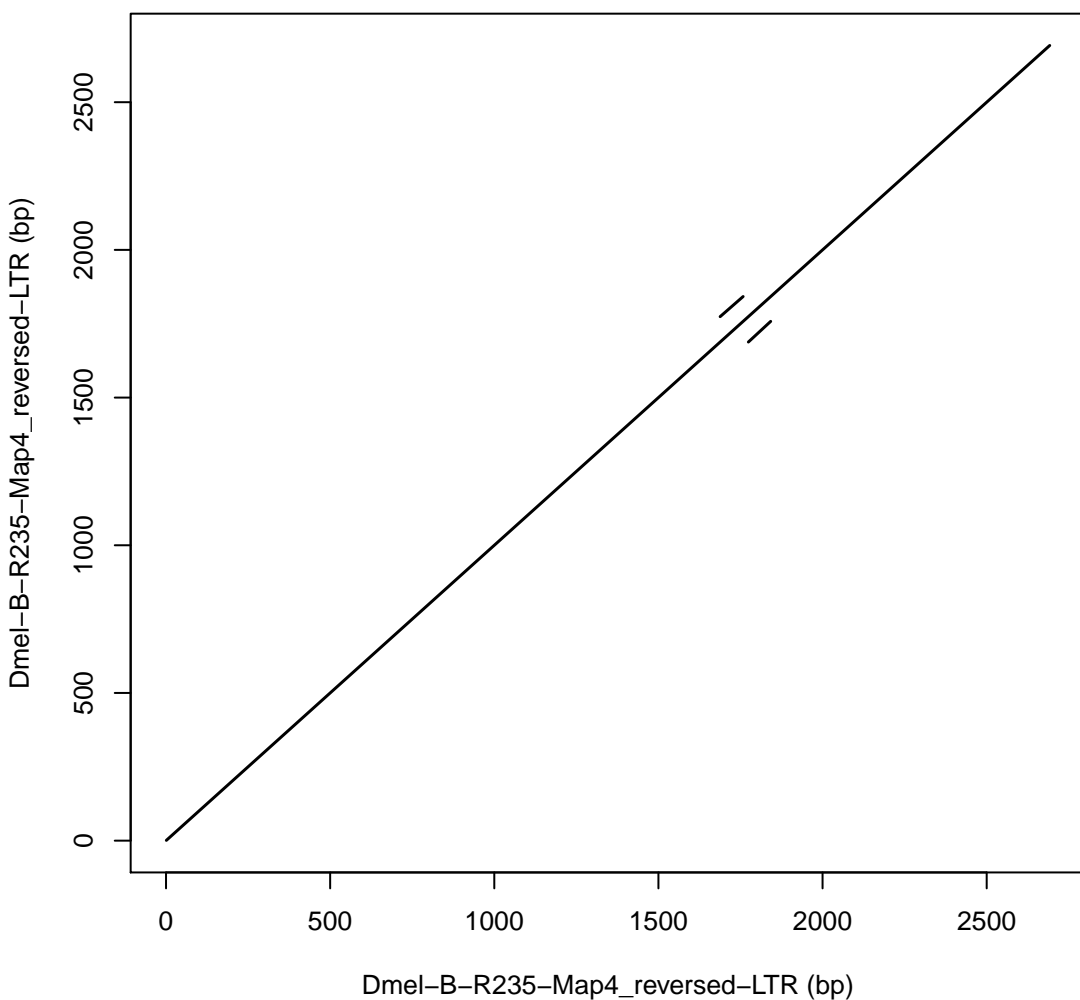
TE: Dmel-B-R235-Map4_reversed-LTR
consensus size: 2692bp; fragments: 62; full length: 2 (≥ 2422.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

