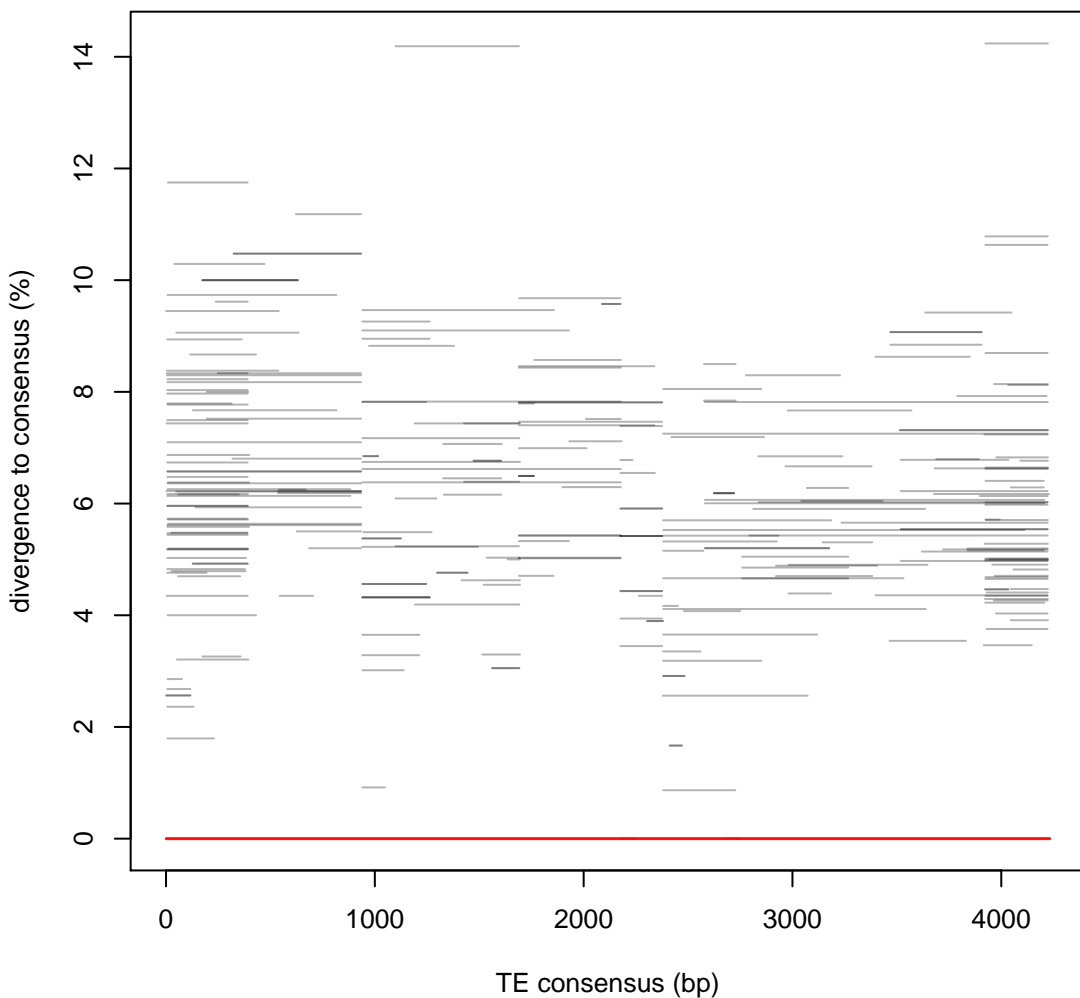
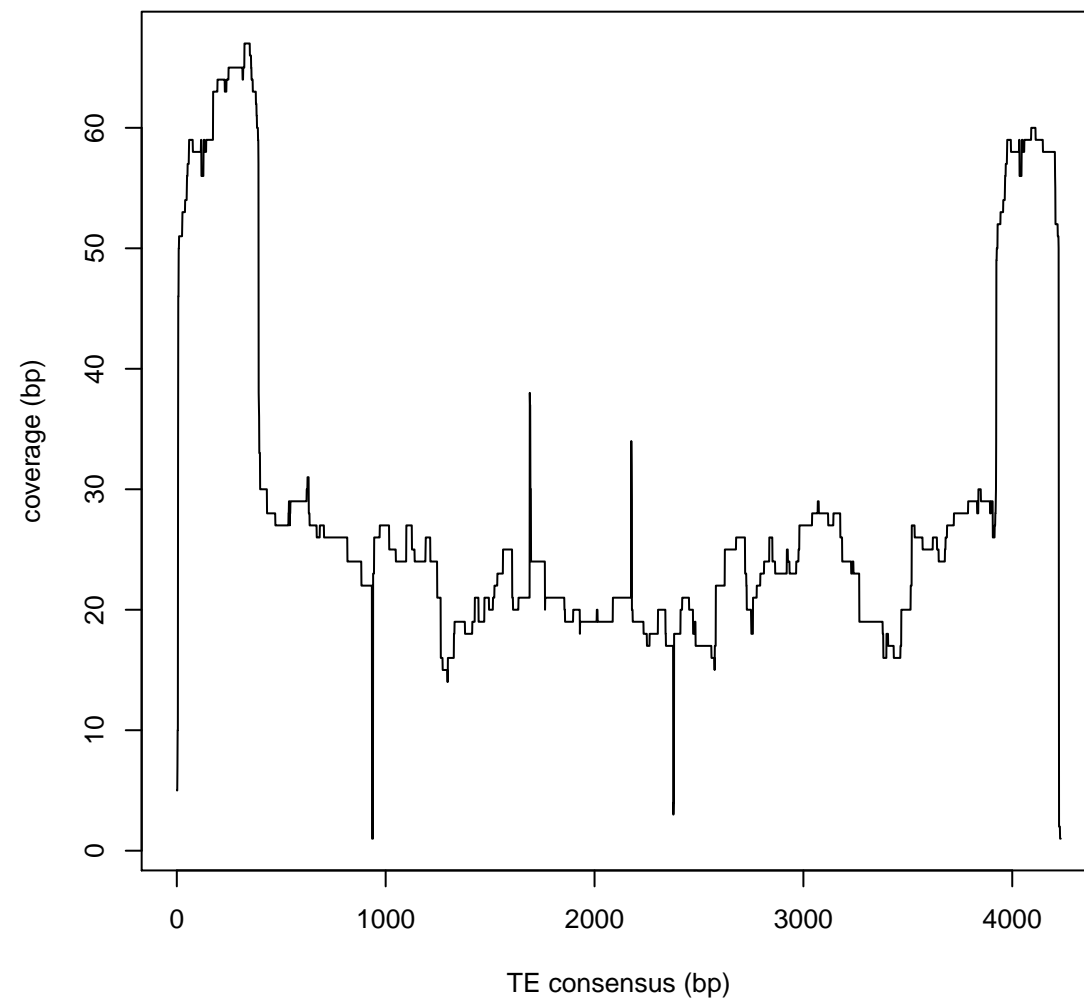


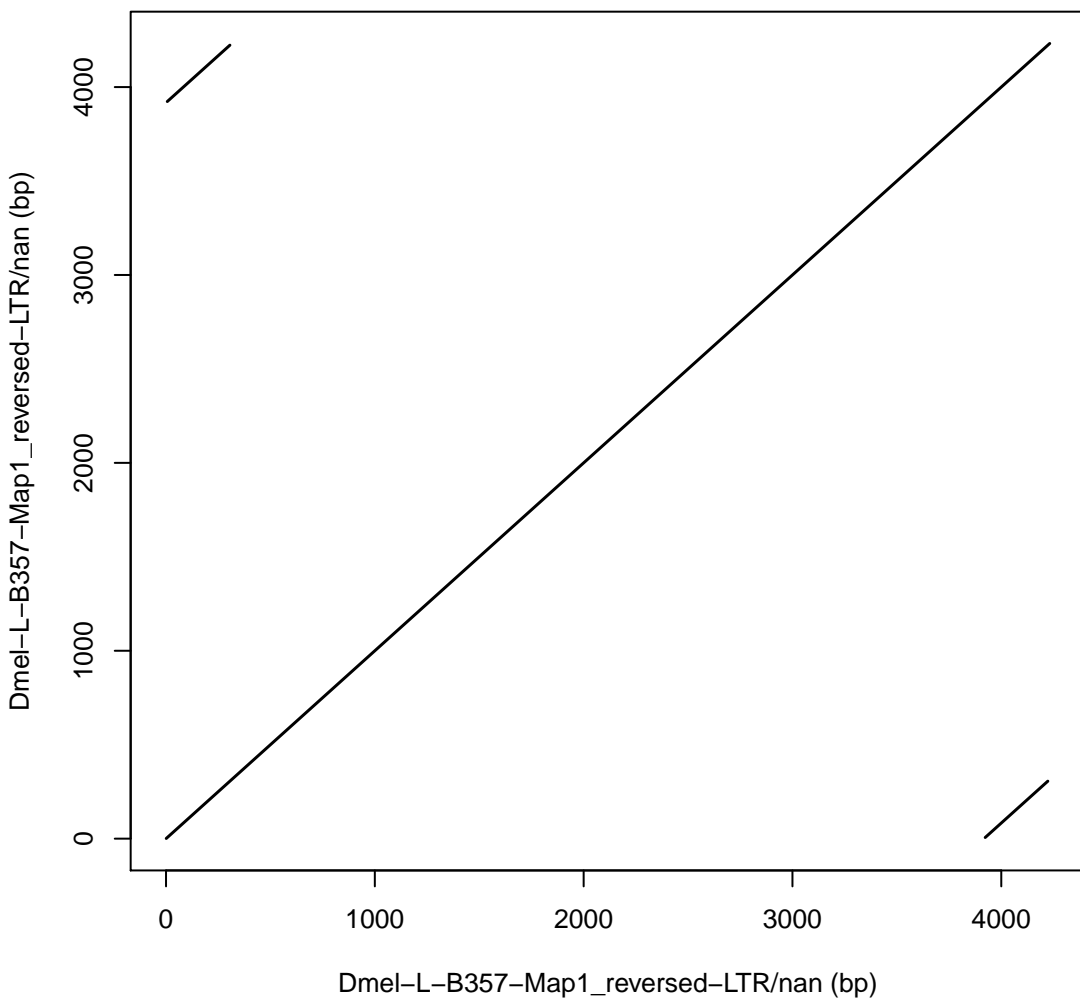
TE: Dmel-L-B357-Map1_reversed-LTR/nan
consensus size: 4232bp; fragments: 281; full length: 1 (≥ 3808.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

