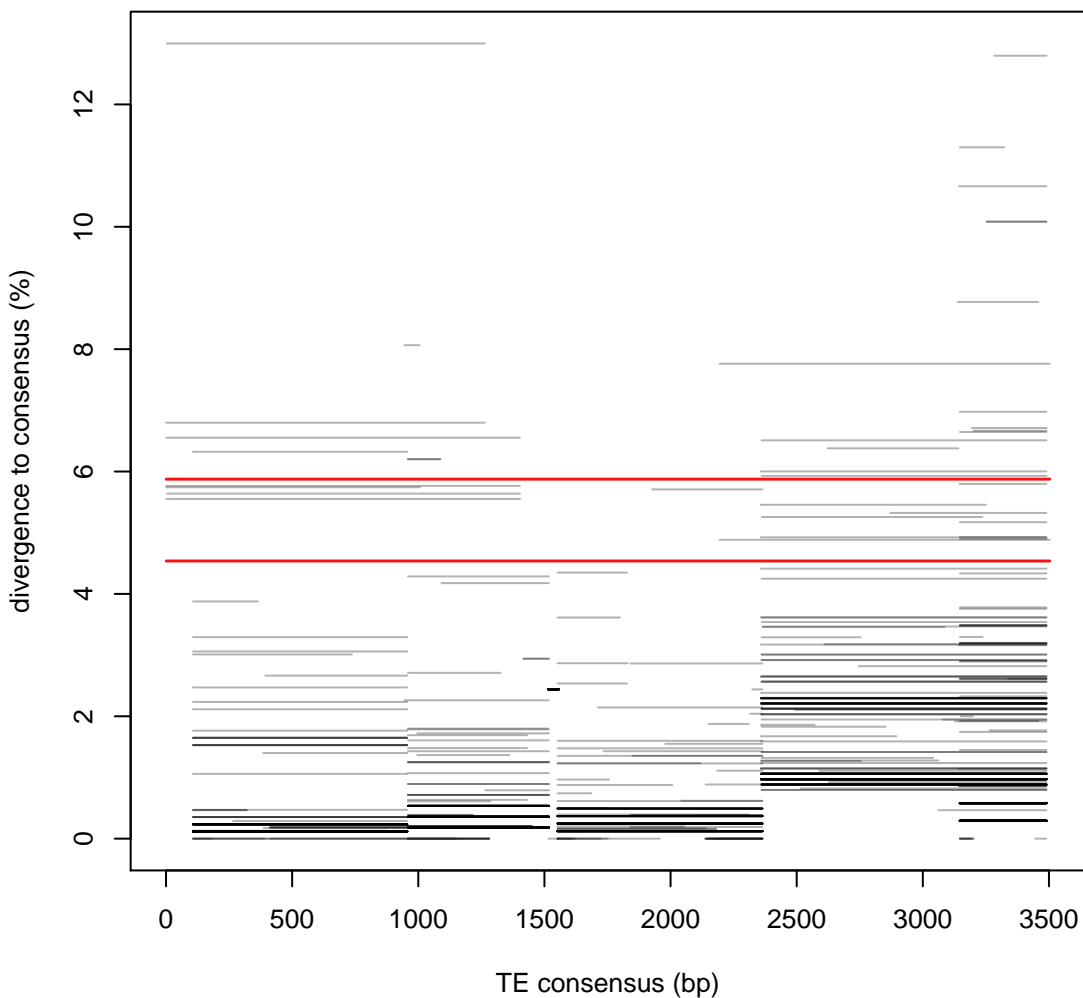
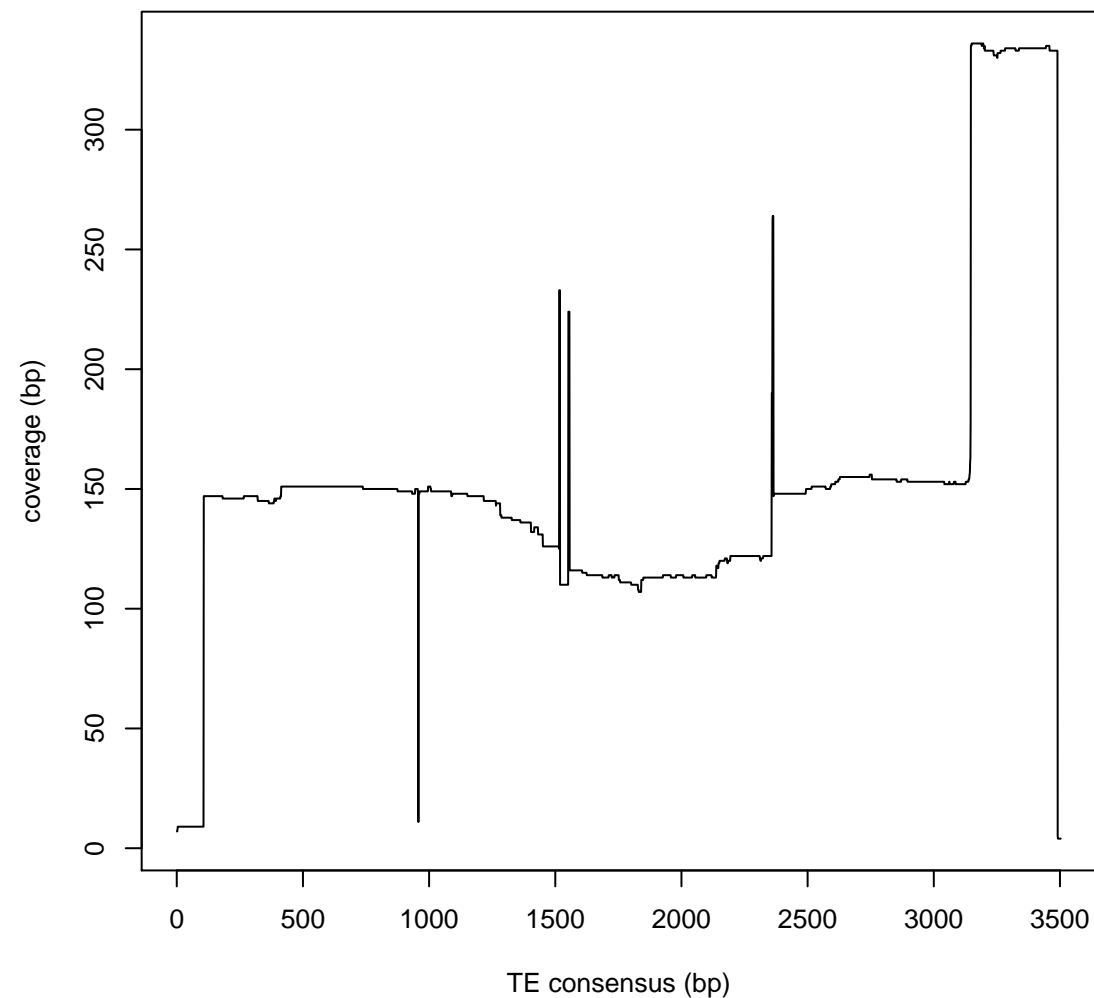


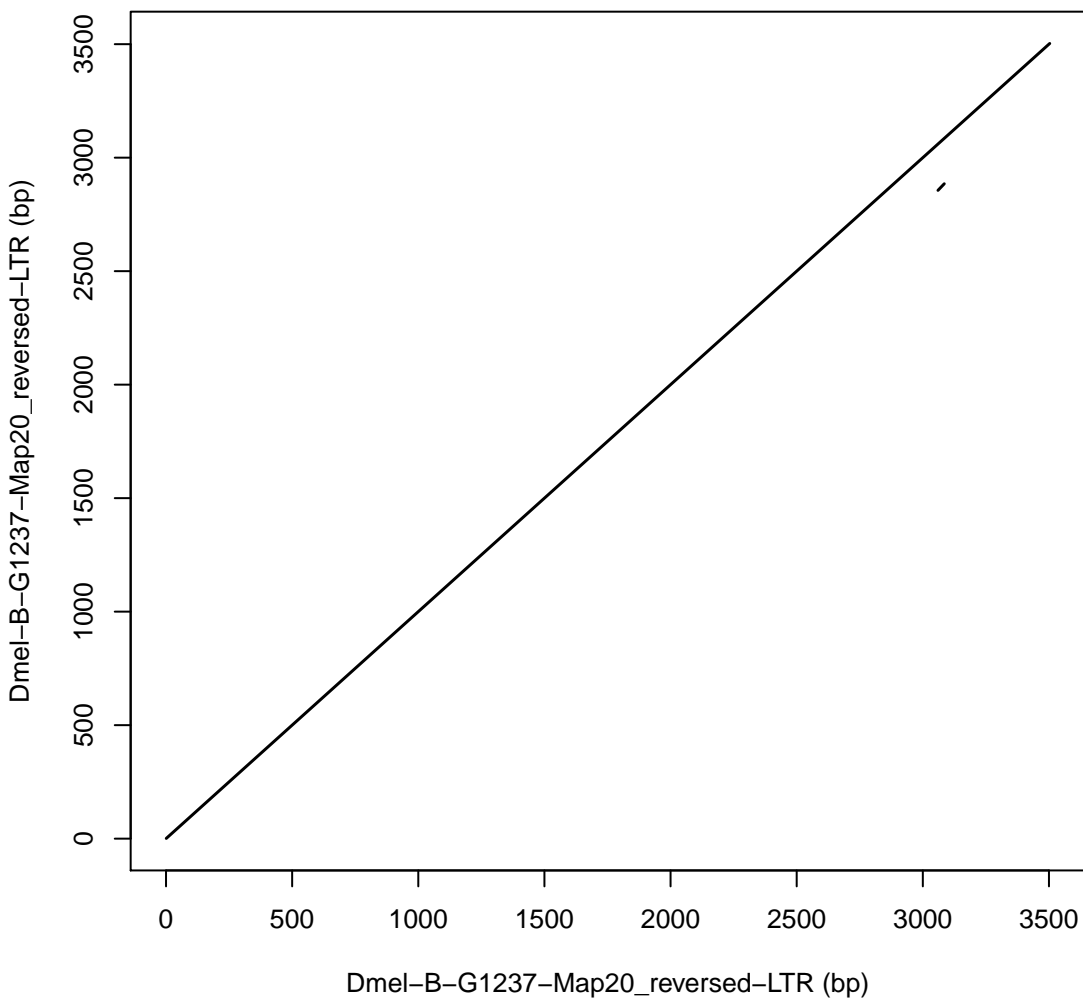
TE: Dmel-B-G1237-Map20_reversed-LTR
consensus size: 3503bp; fragments: 899; full length: 2 (≥ 3152.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

