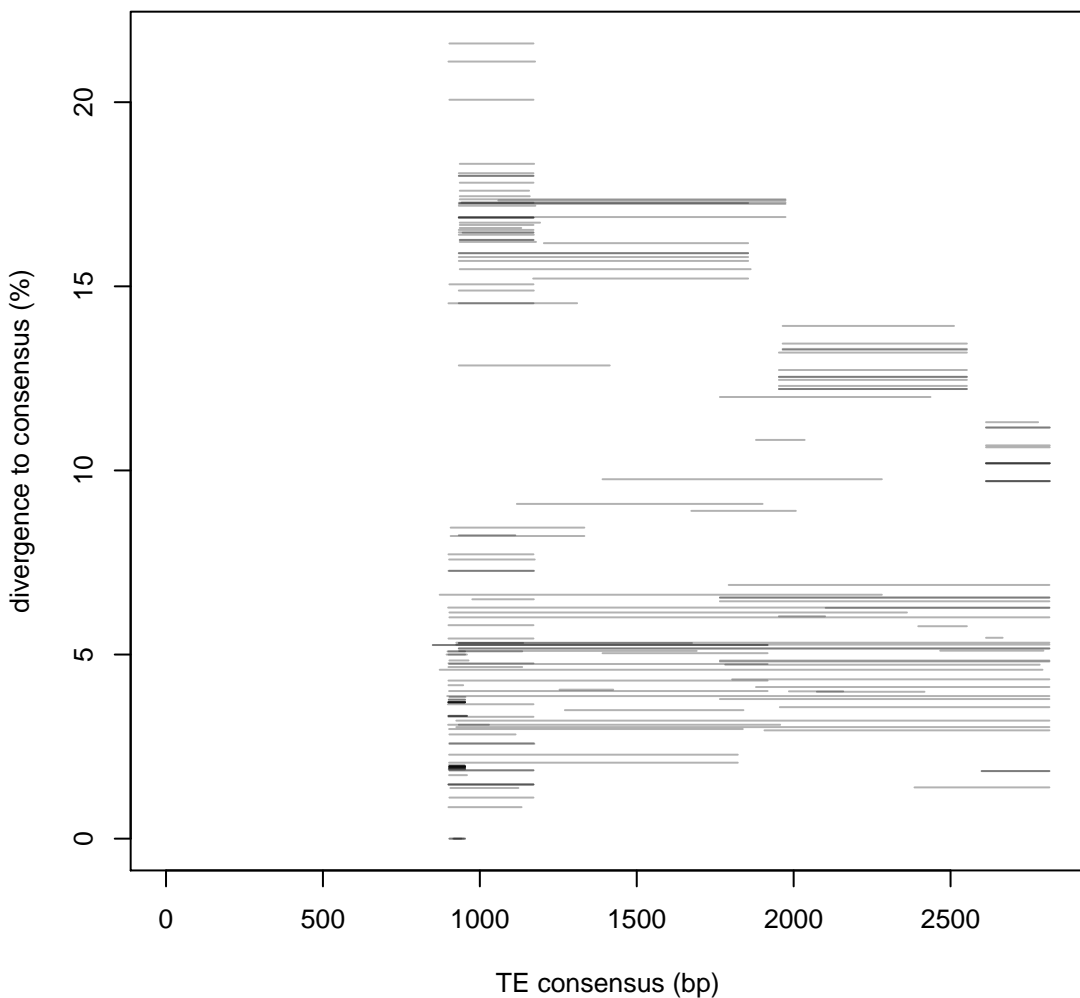
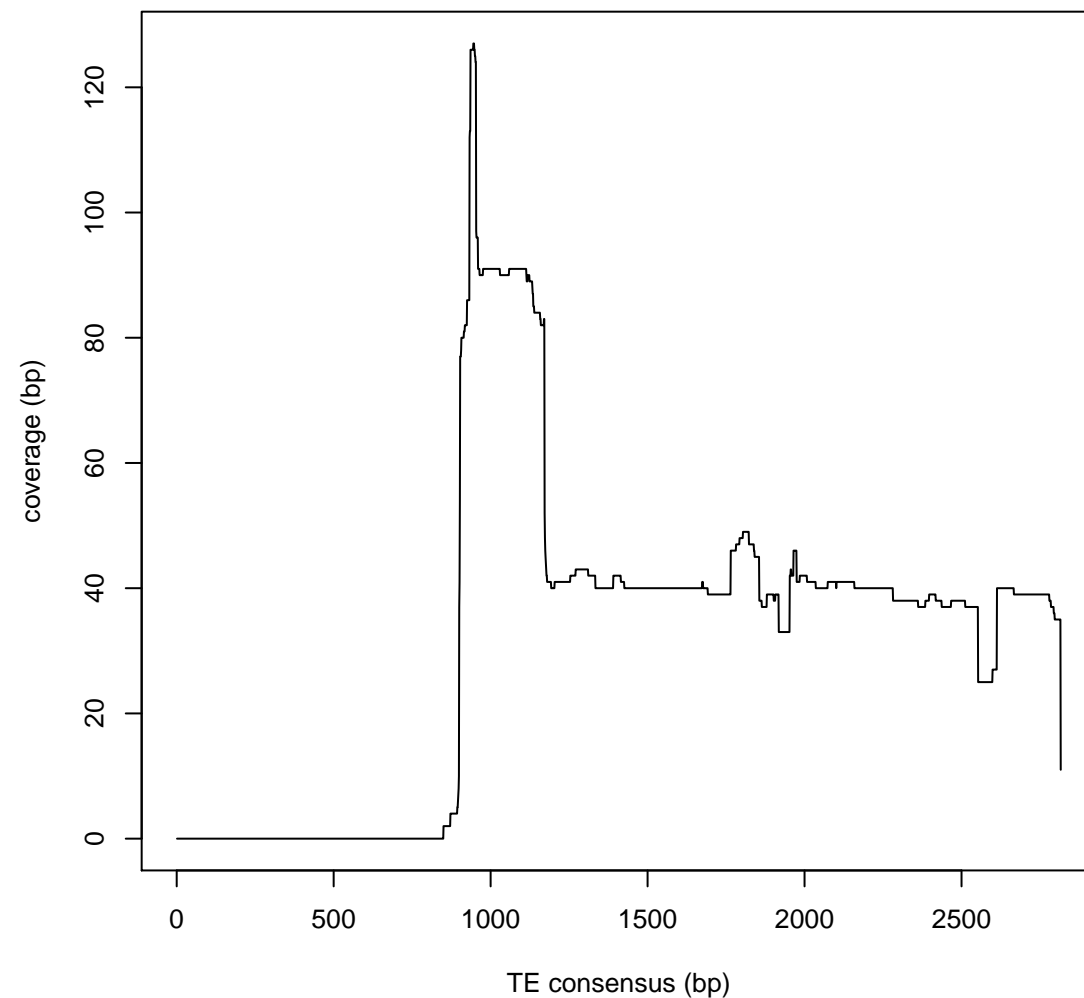


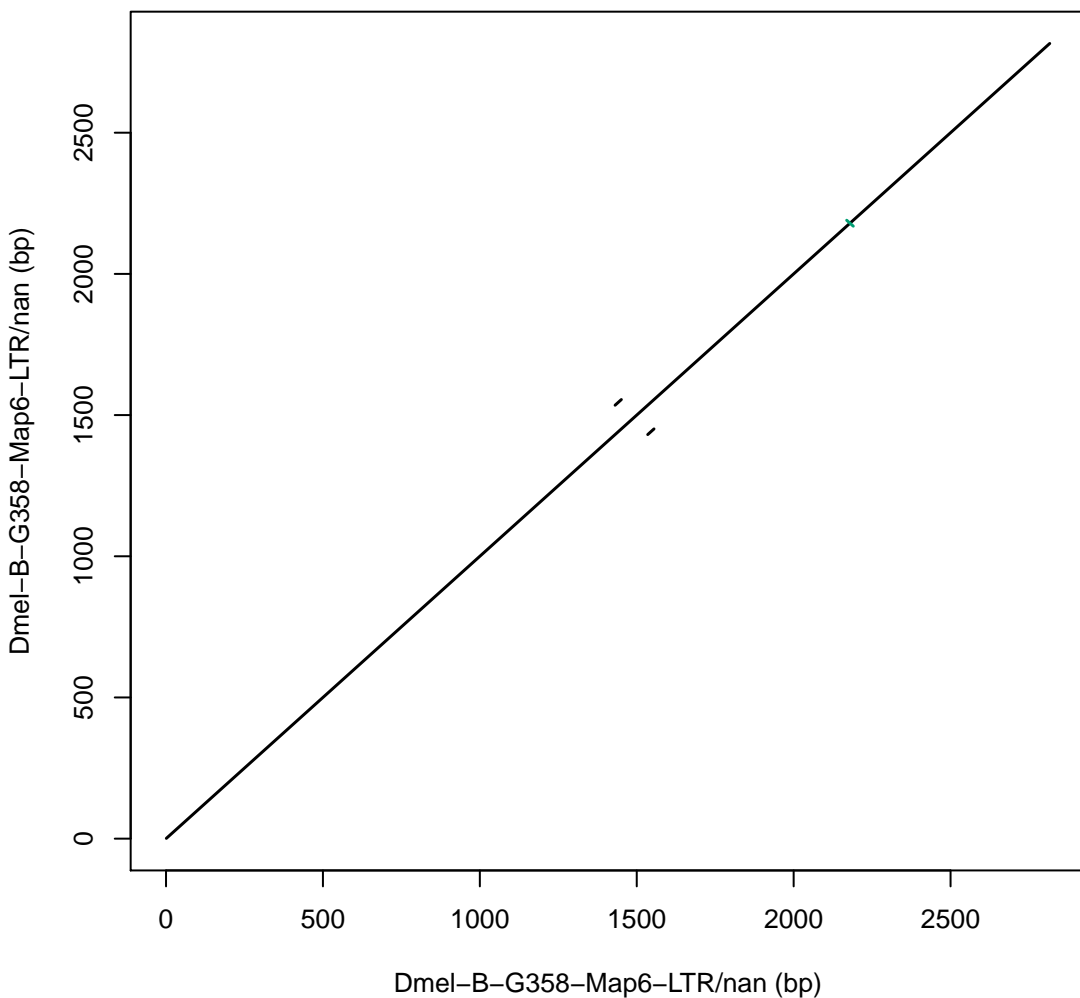
TE: Dmel-B-G358-Map6-LTR/nan
consensus size: 2816bp; fragments: 186; full length: 0 (≥ 2534.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

