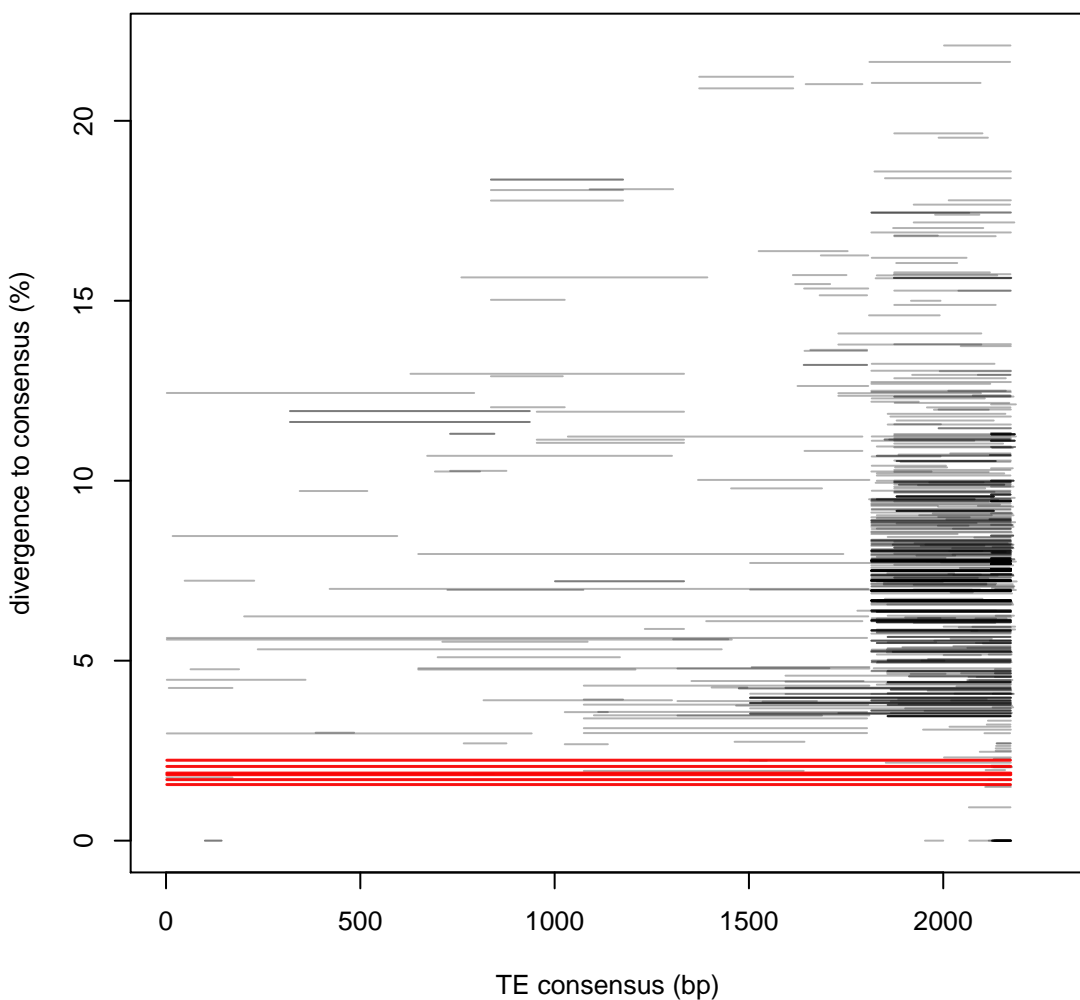
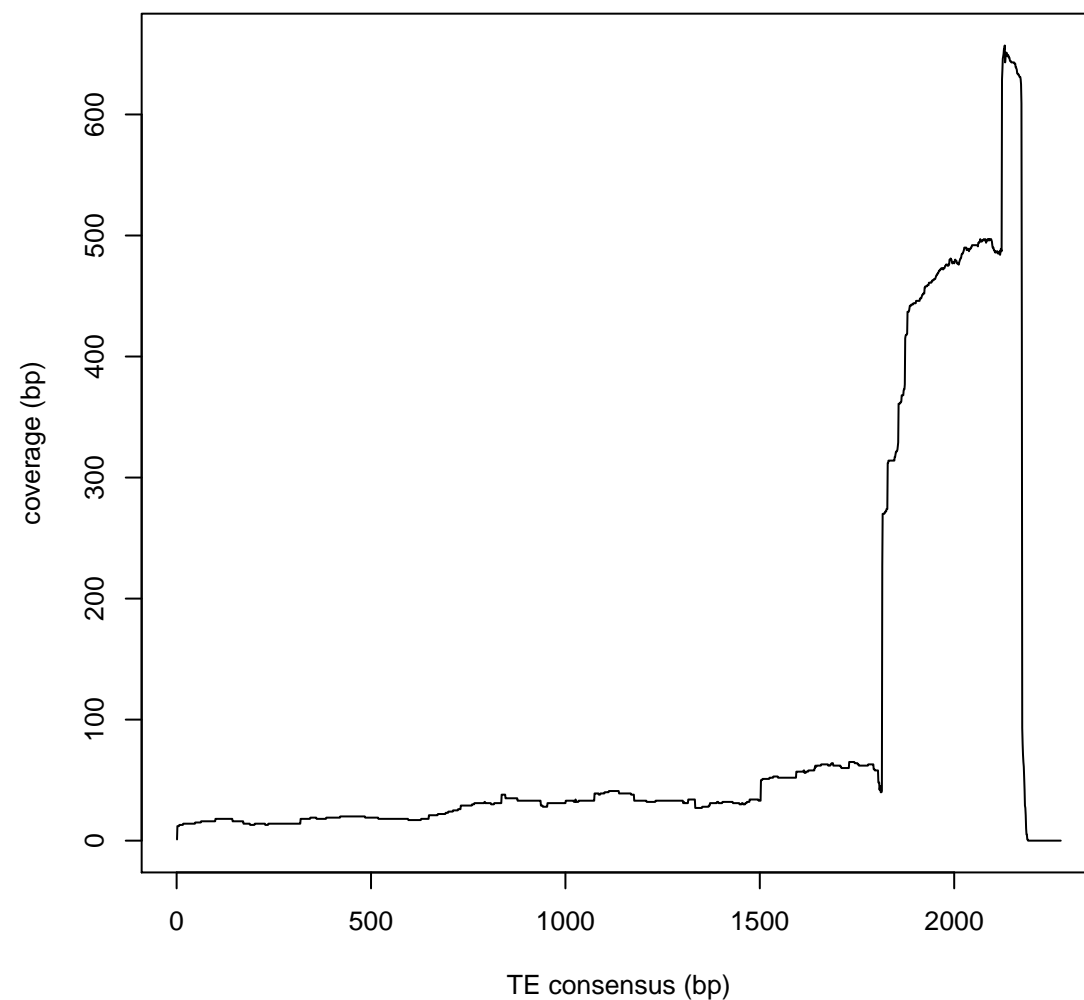


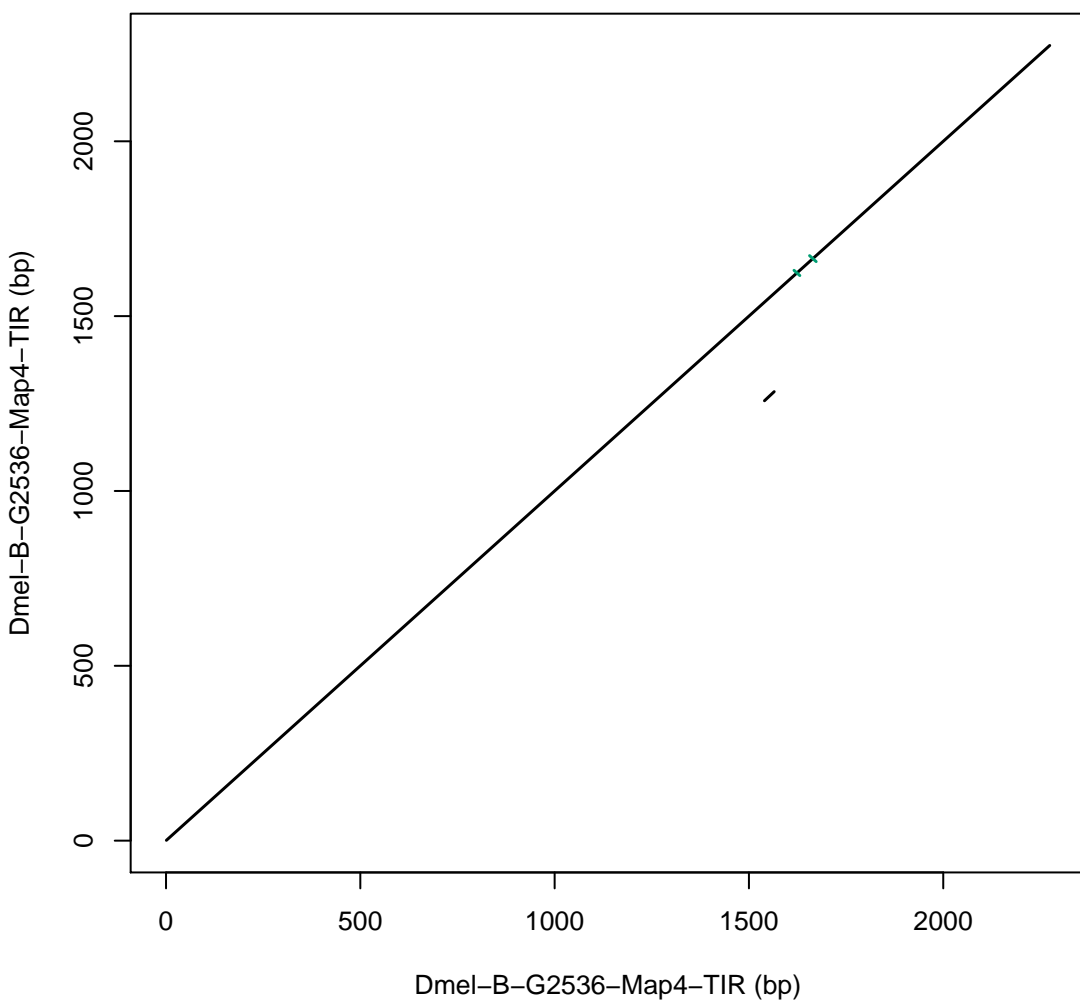
TE: Dmel-B-G2536-Map4-TIR
consensus size: 2274bp; fragments: 829; full length: 6 (≥ 2046.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

