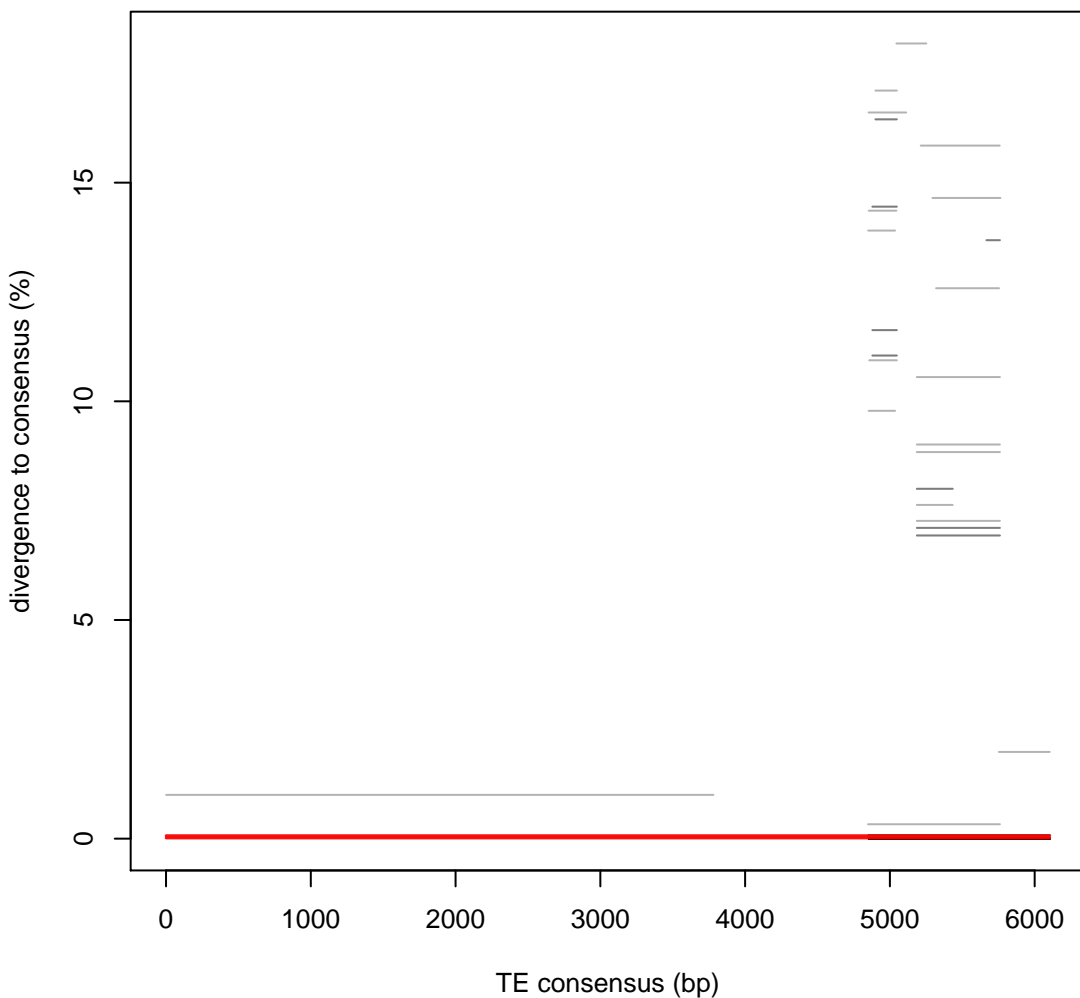
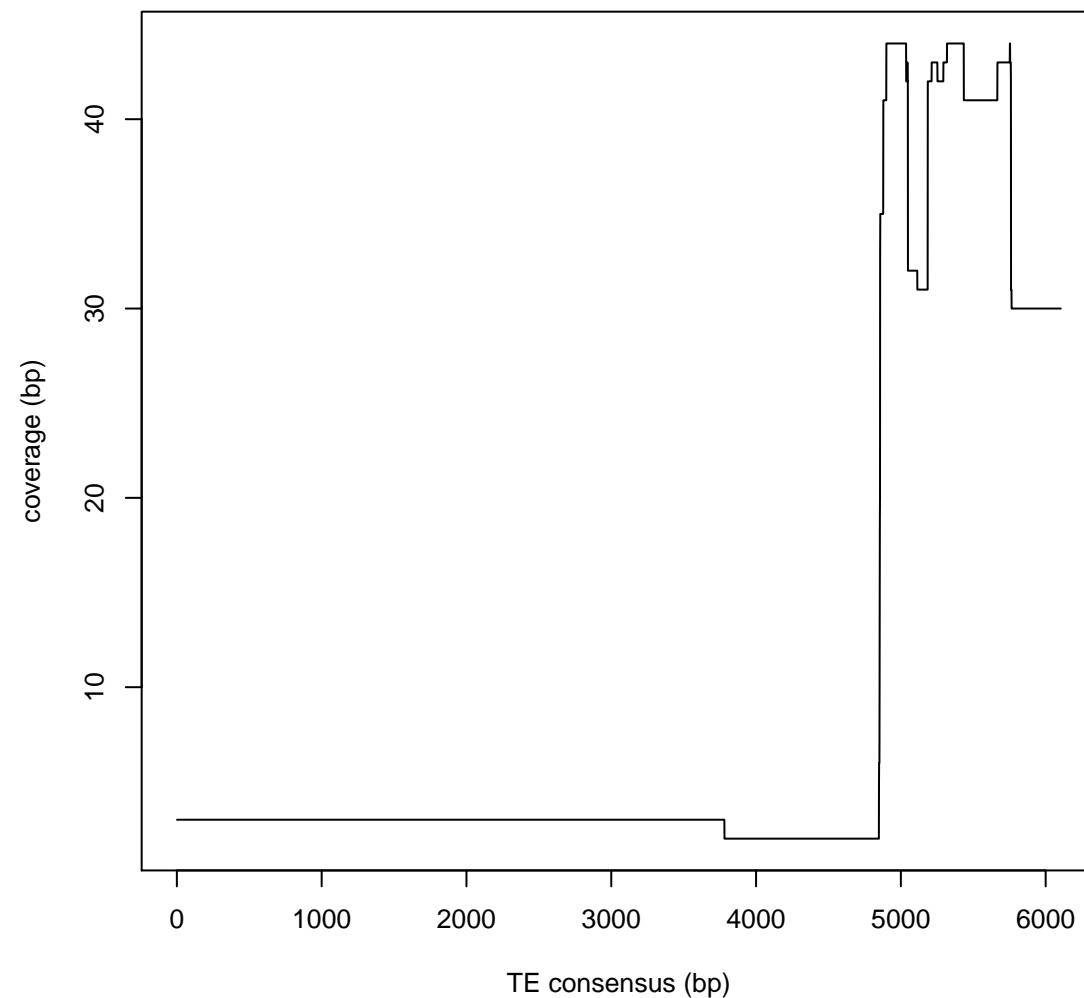


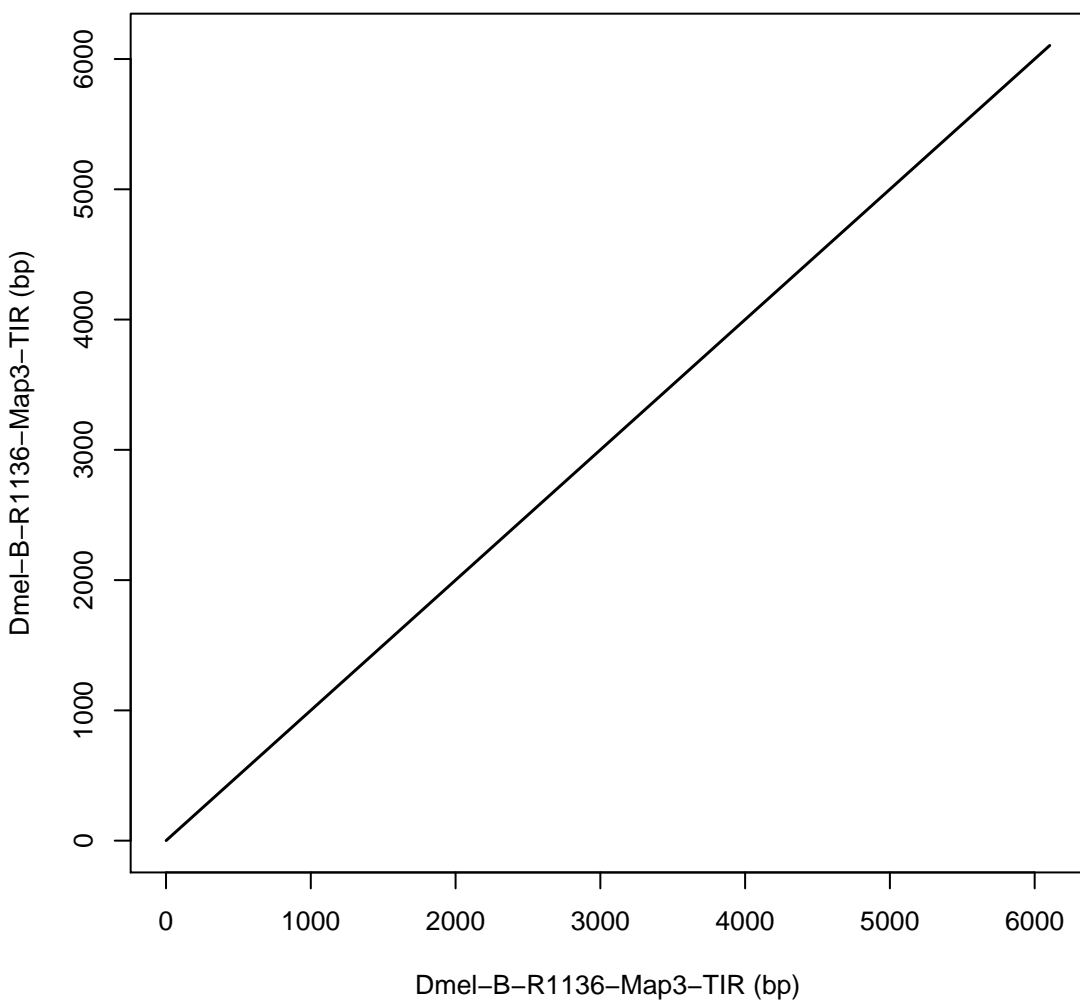
TE: Dmel-B-R1136-Map3-TIR
consensus size: 6104bp; fragments: 63; full length: 2 (≥ 5493.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

