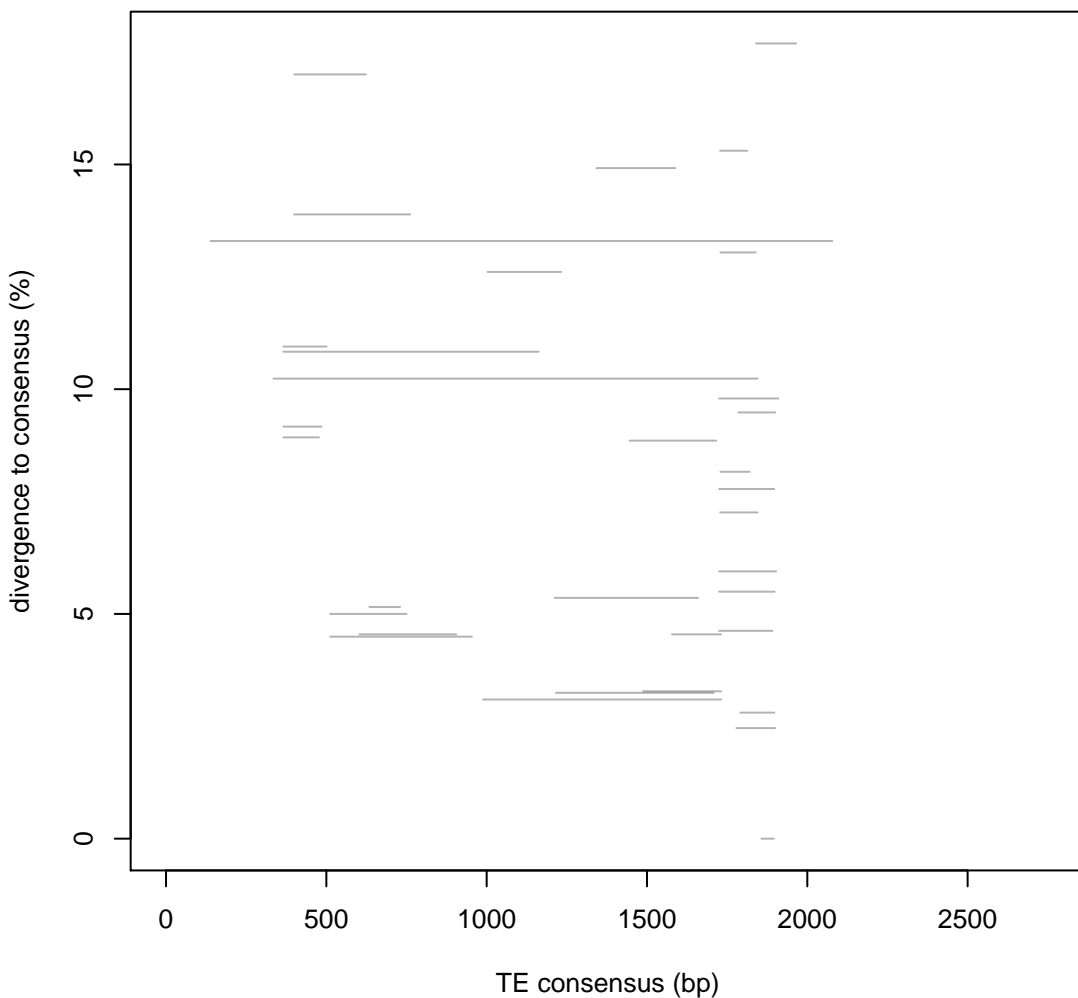
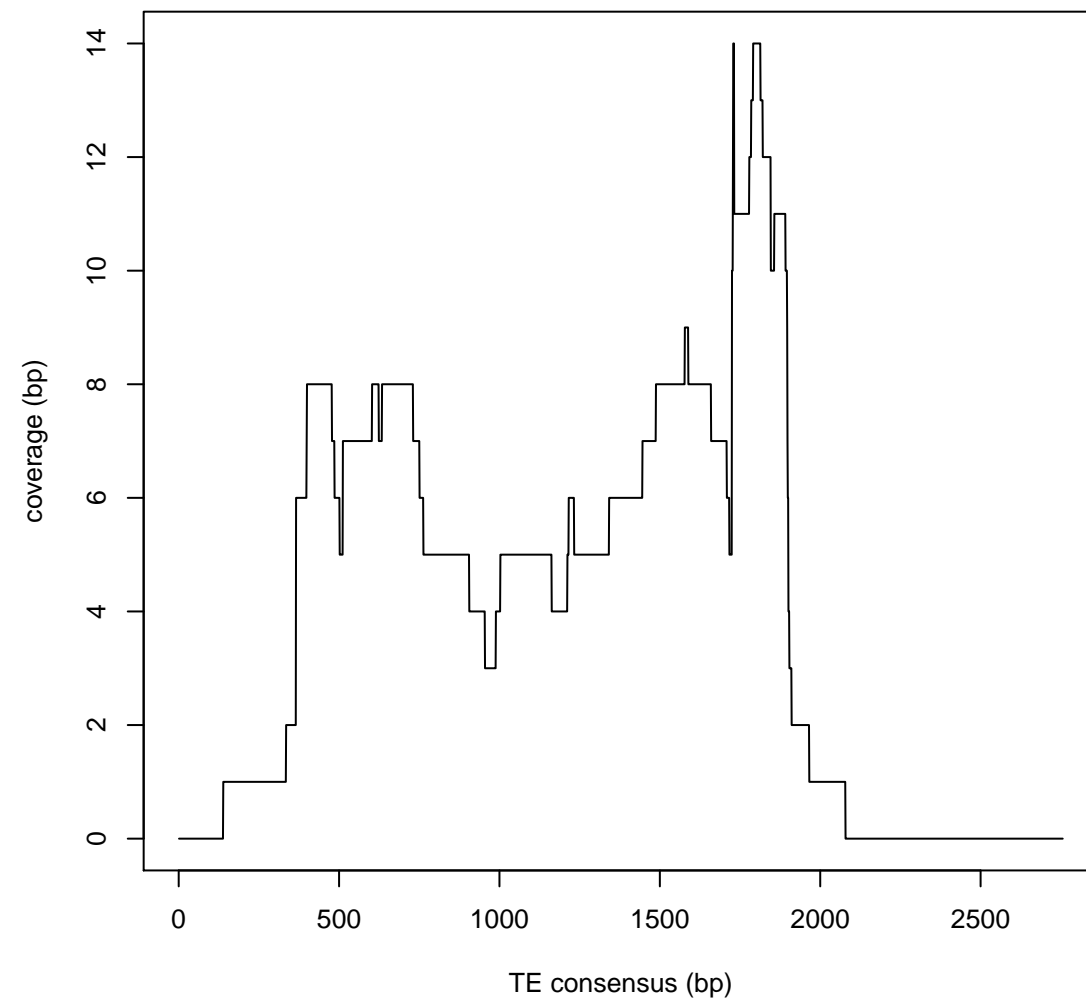


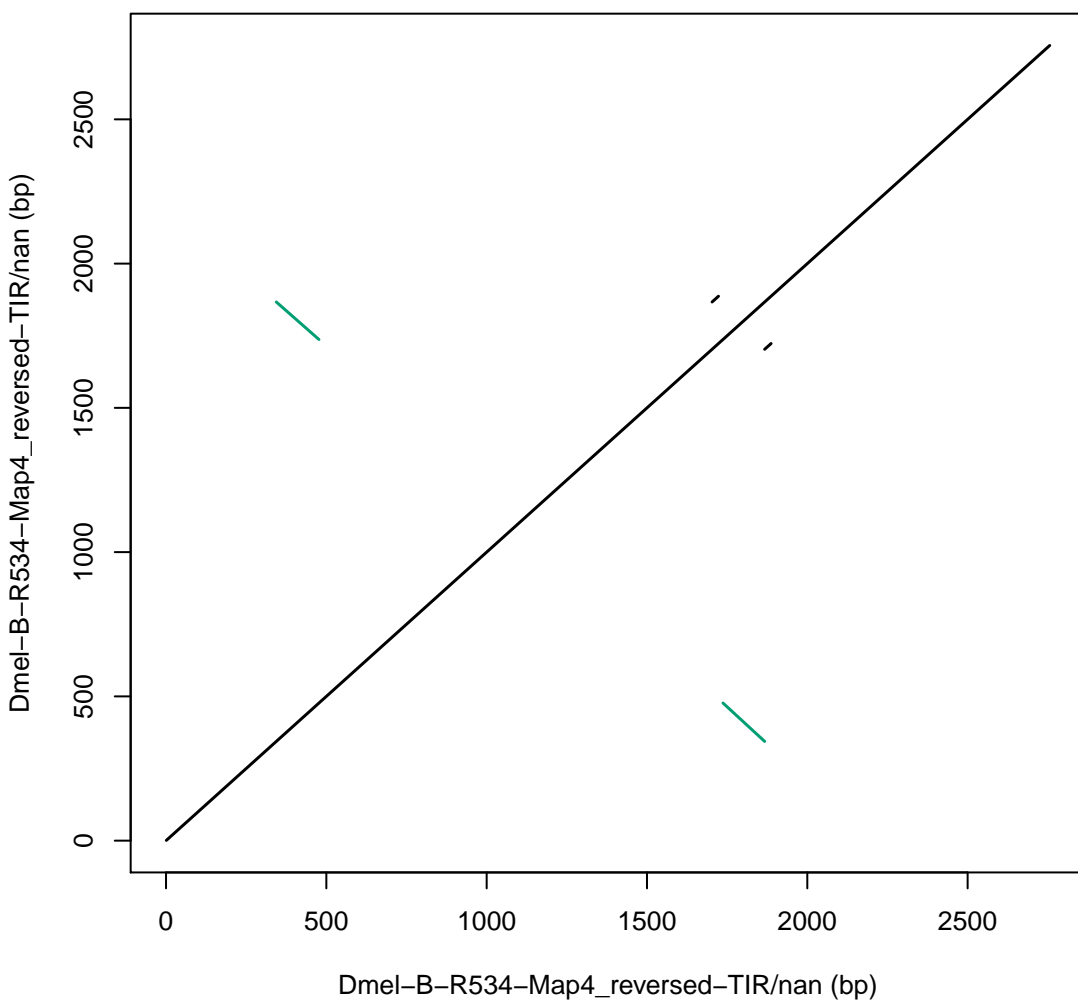
TE: Dmel-B-R534-Map4\_reversed-TIR/nan  
consensus size: 2756bp; fragments: 34; full length: 0 ( $\geq 2480.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

