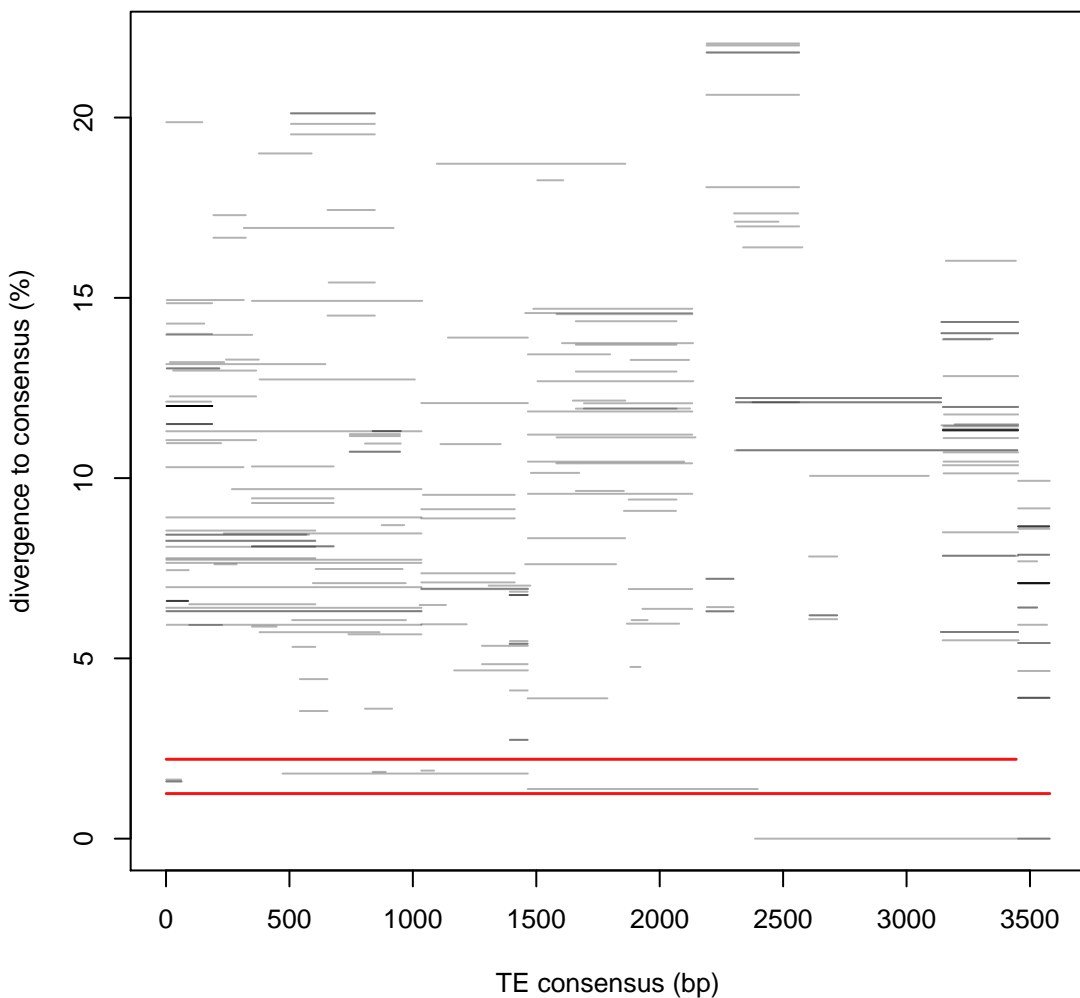
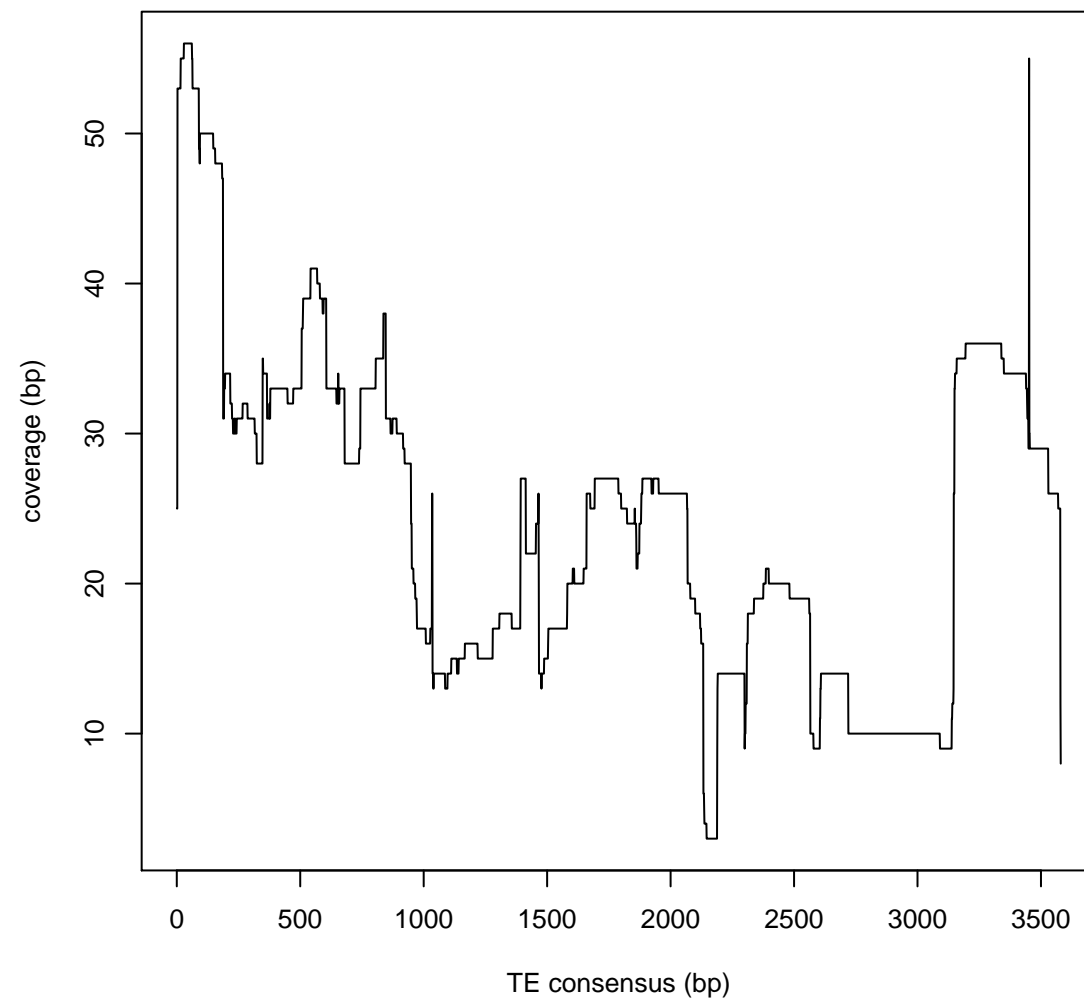


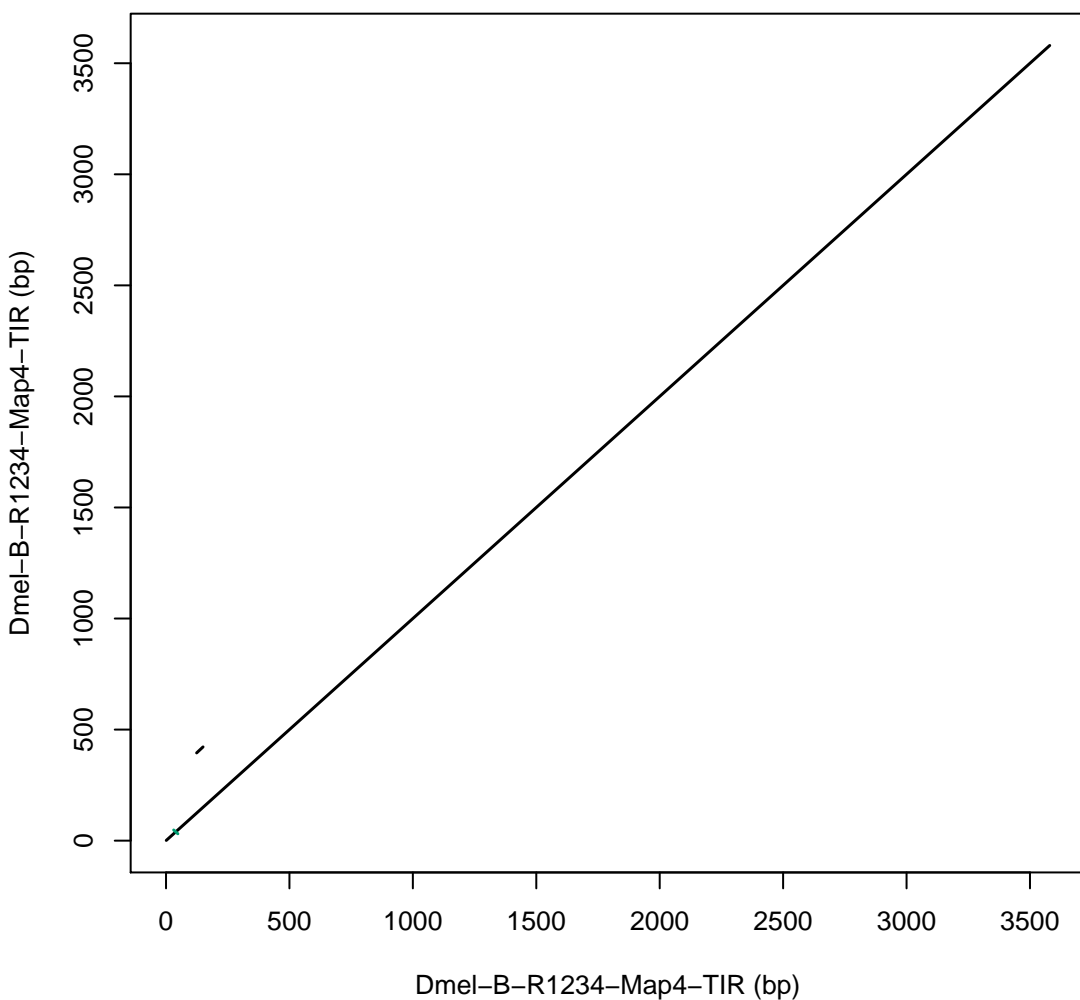
TE: Dmel-B-R1234-Map4-TIR
consensus size: 3580bp; fragments: 248; full length: 2 (≥ 3222 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

