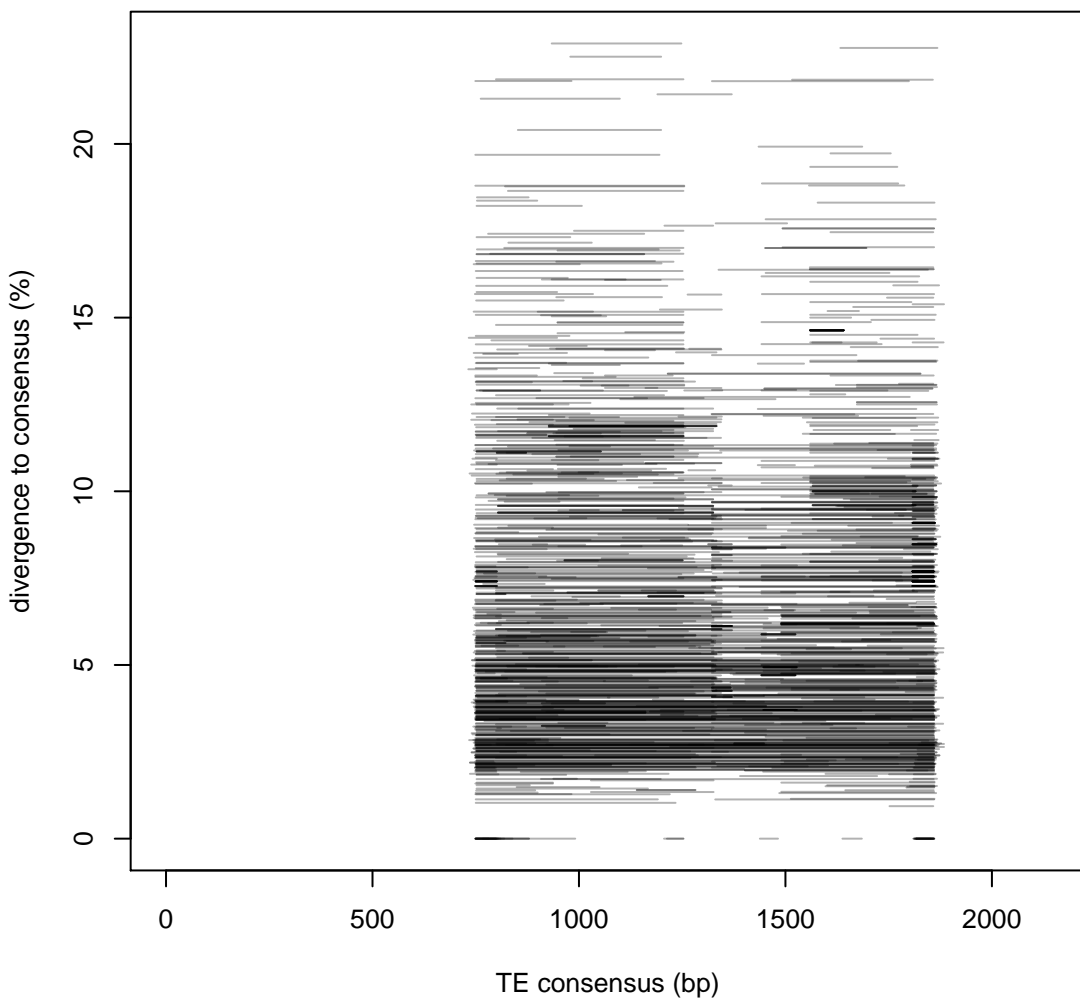
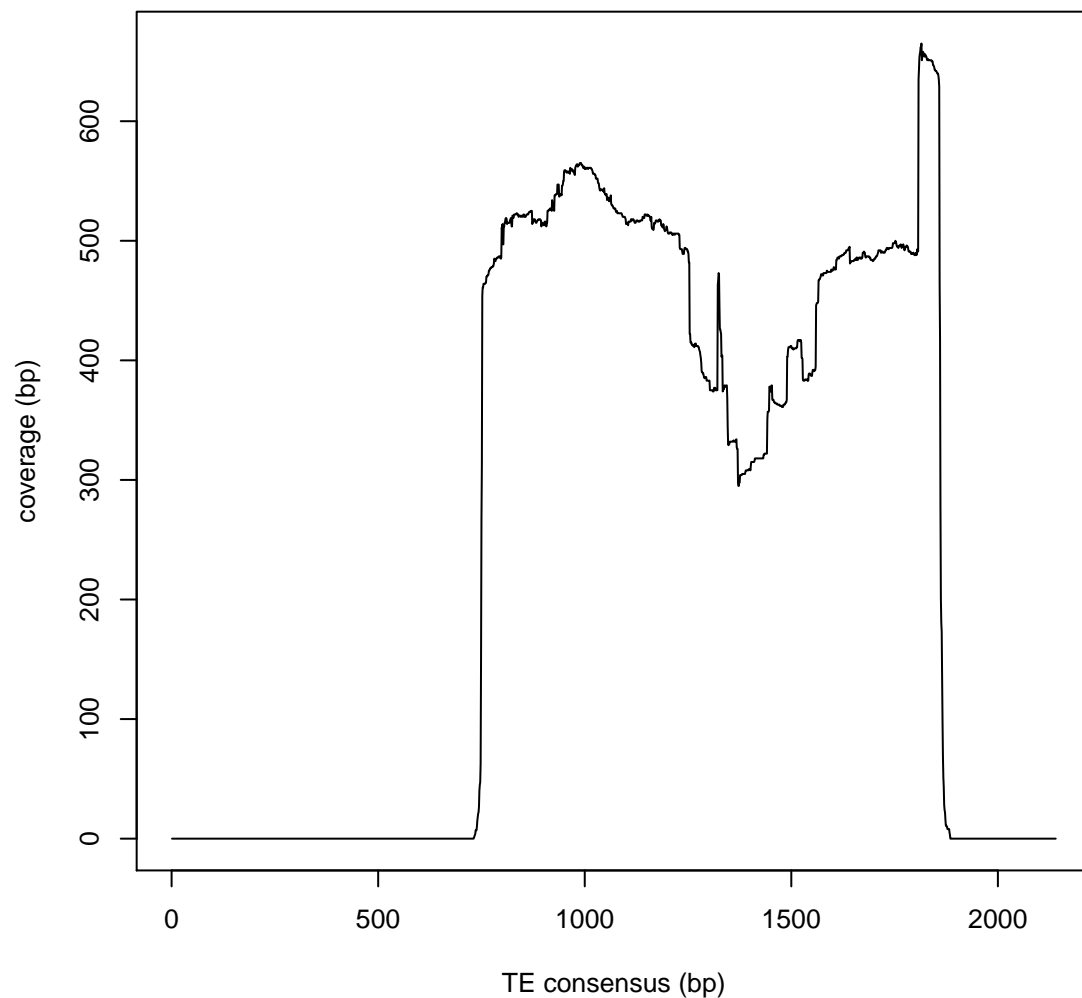


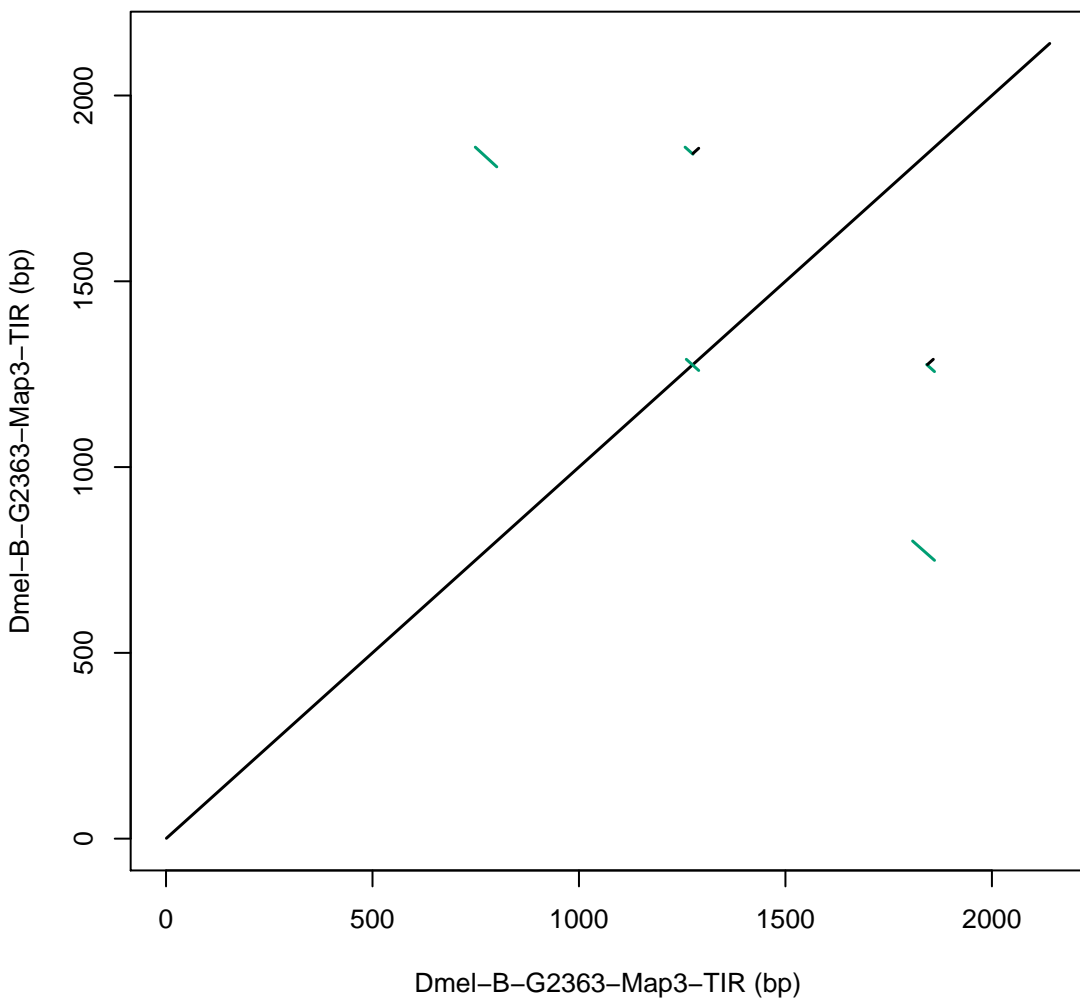
TE: Dmel-B-G2363-Map3-TIR
consensus size: 2140bp; fragments: 1494; full length: 0 (>=1926bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

