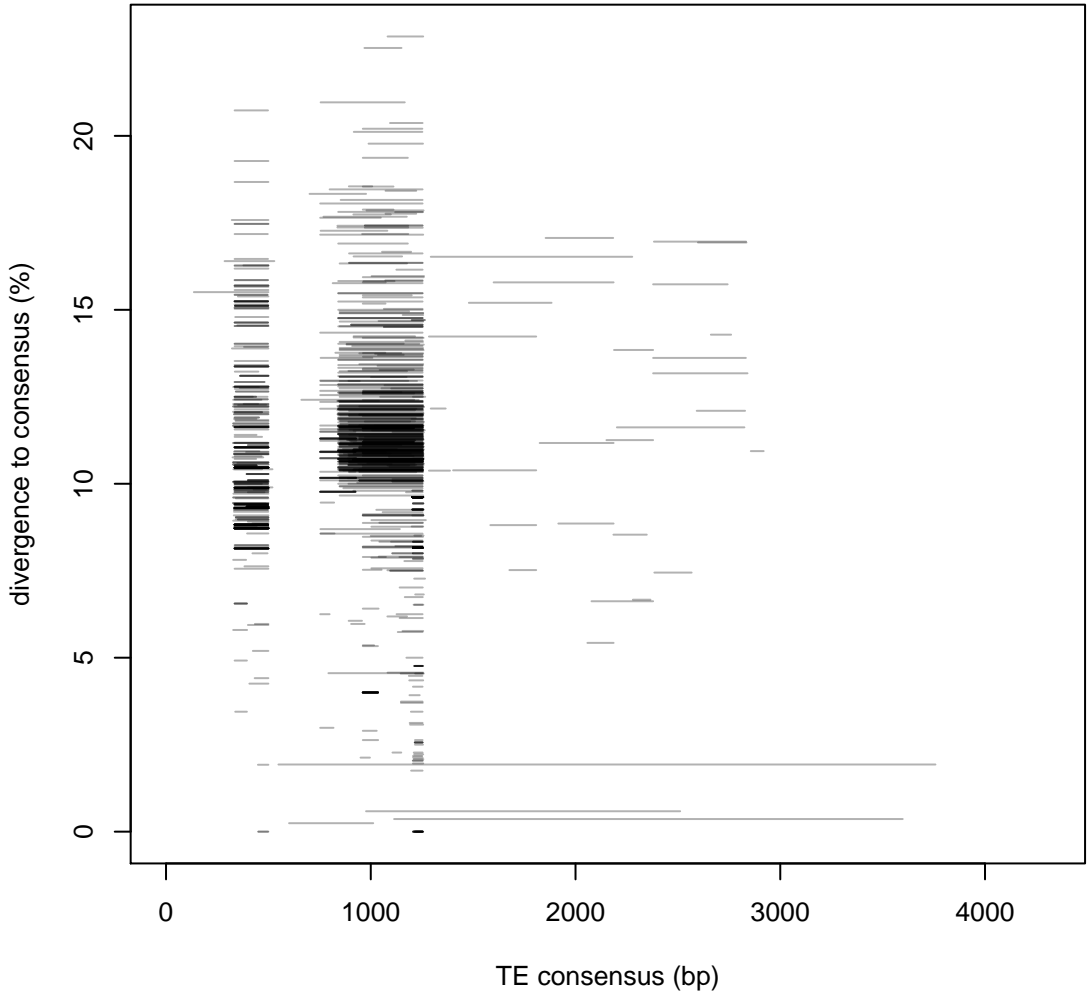
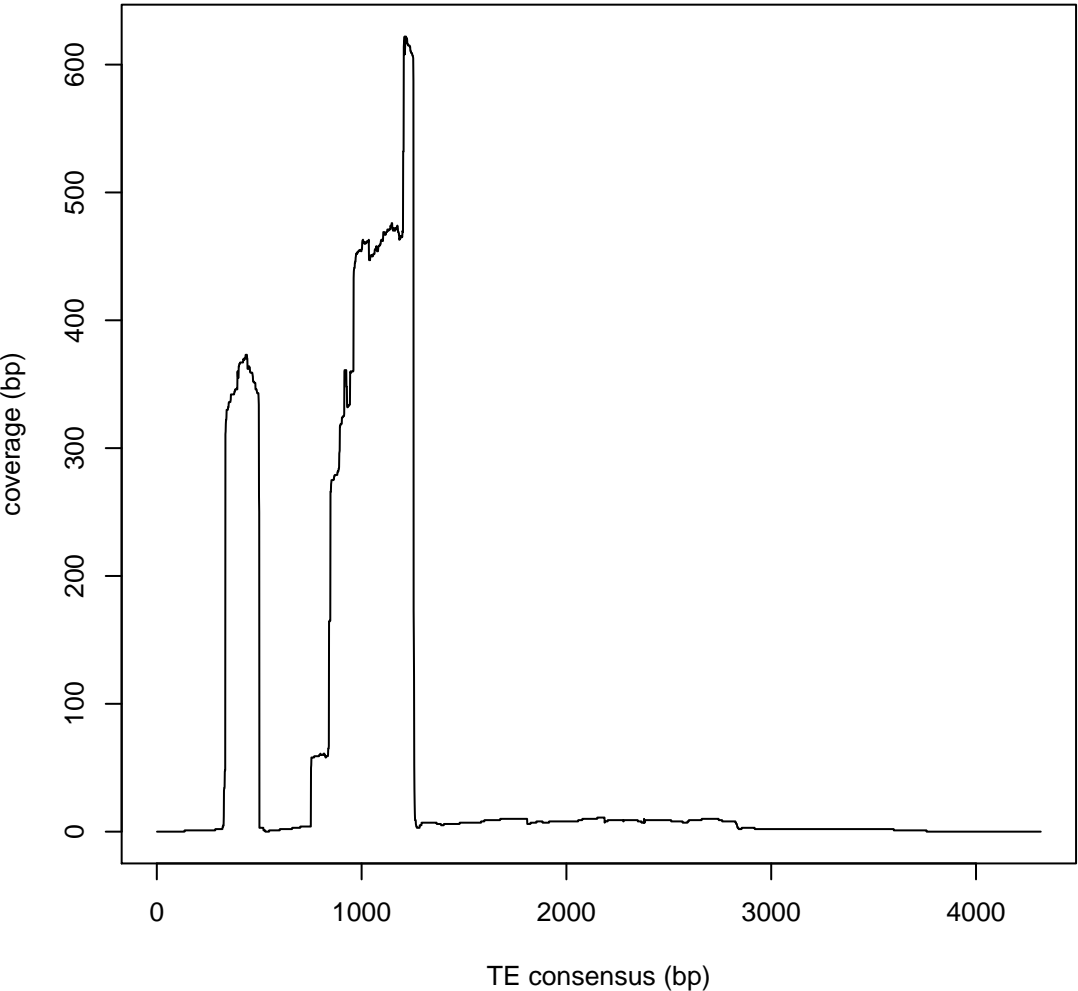


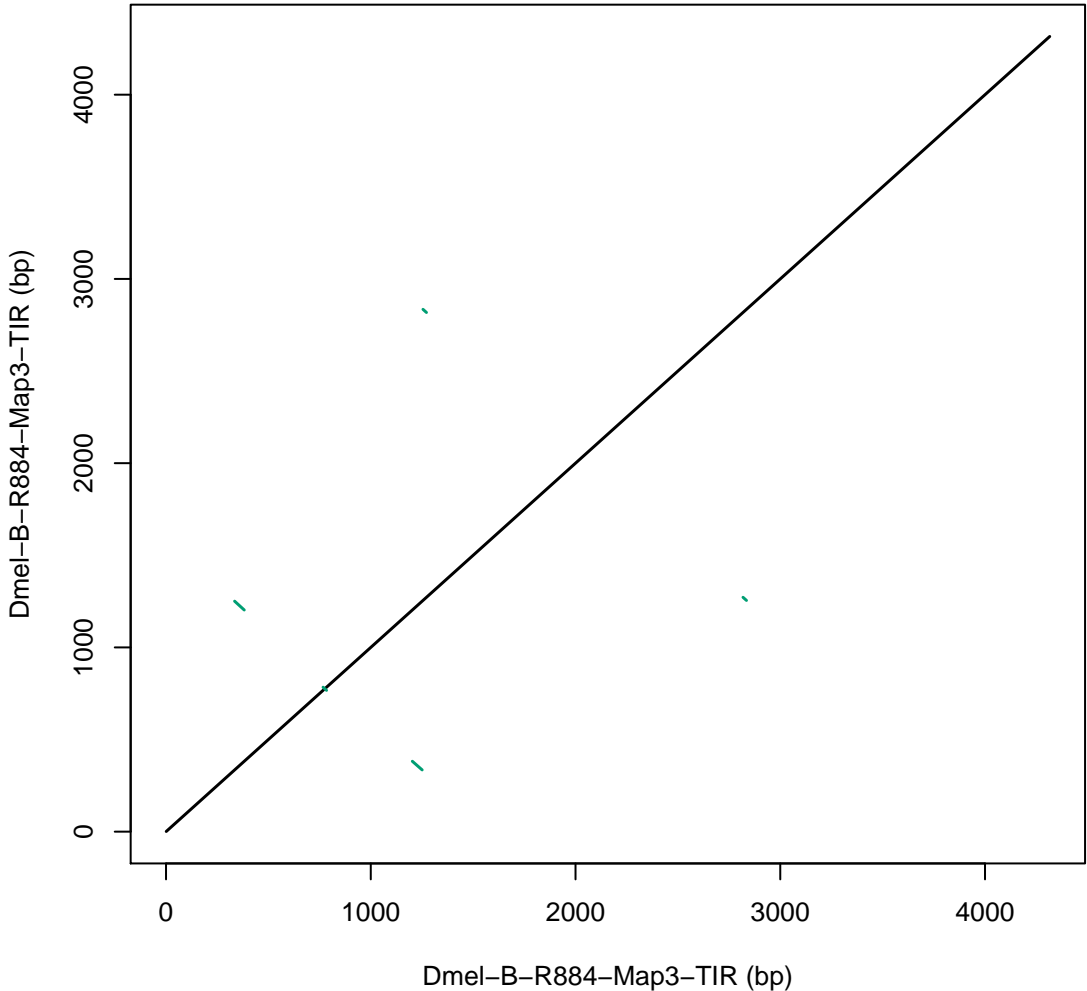
TE: Dmel-B-R884-Map3-TIR
consensus size: 4316bp; fragments: 1162; full length: 0 (>=3884.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

