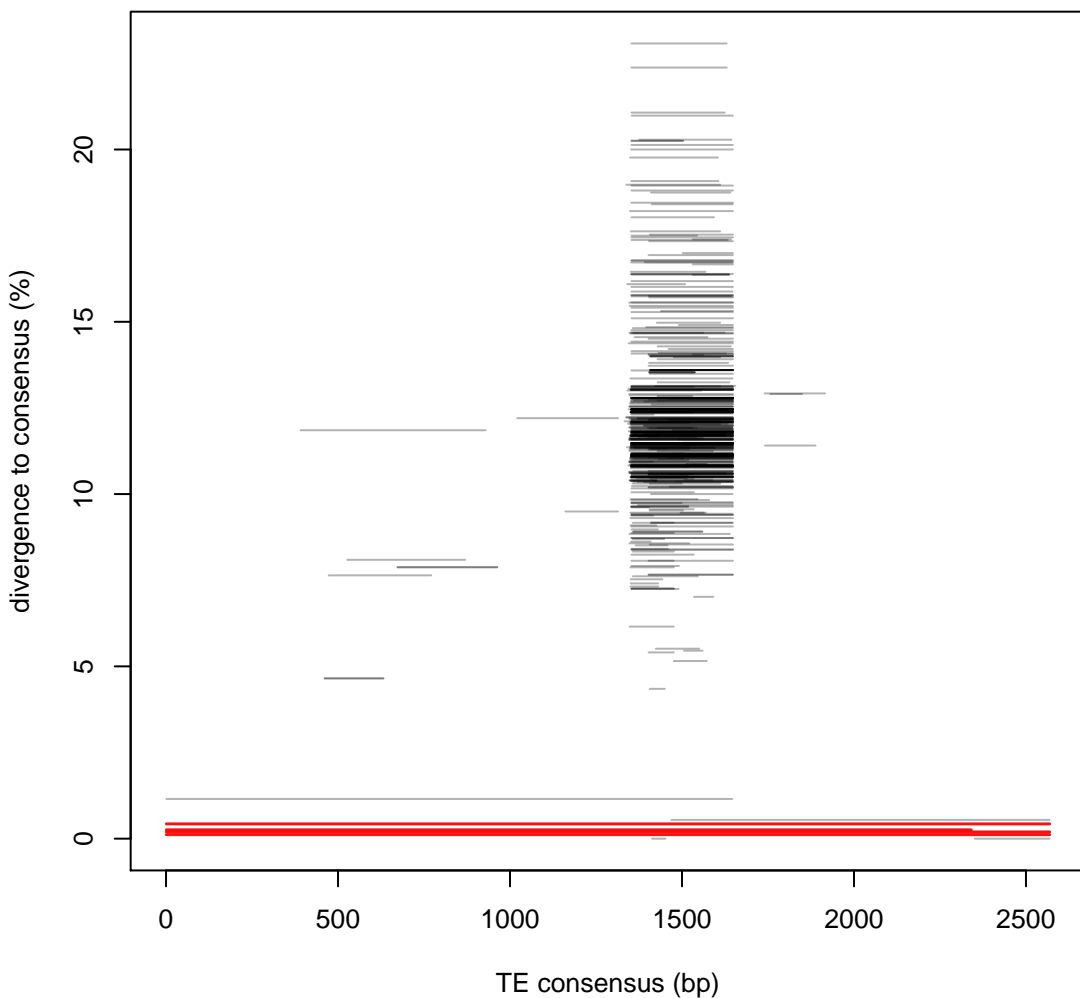
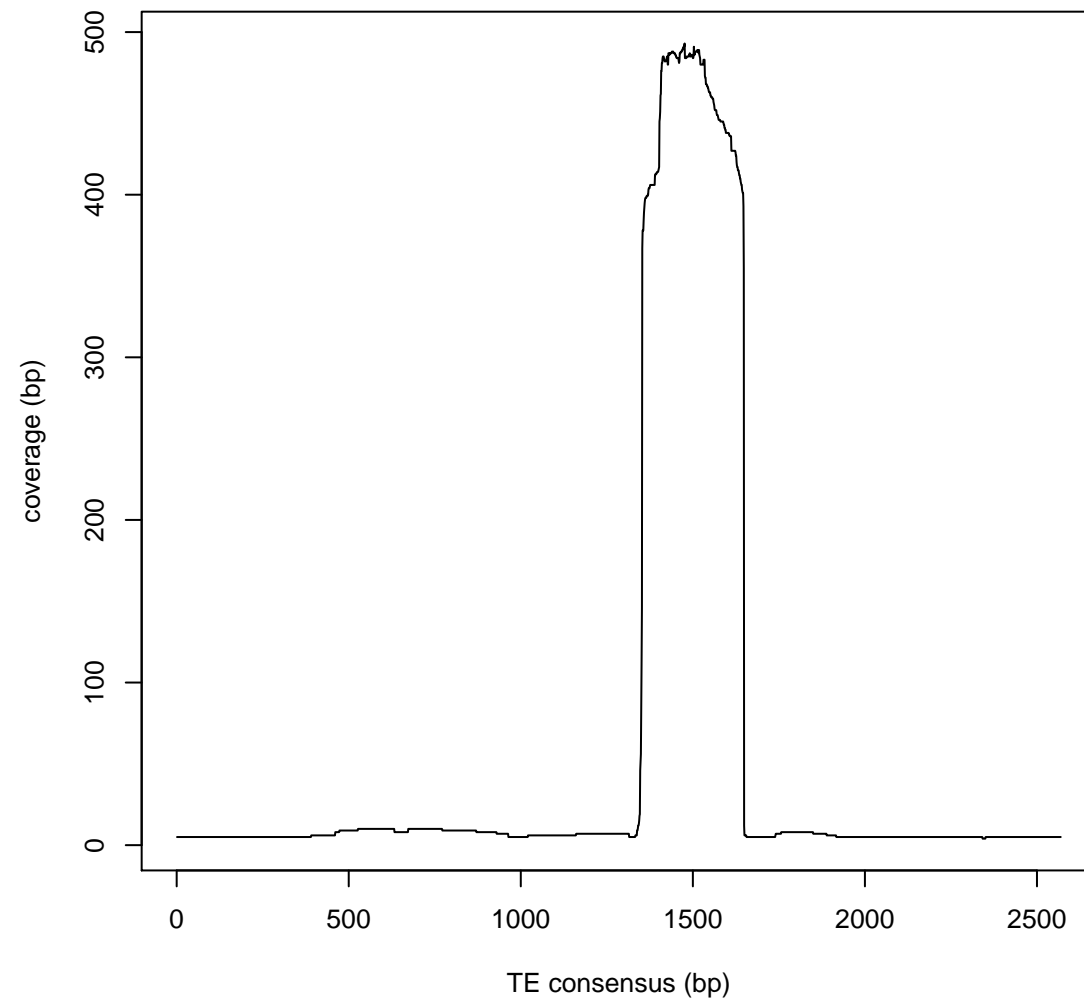


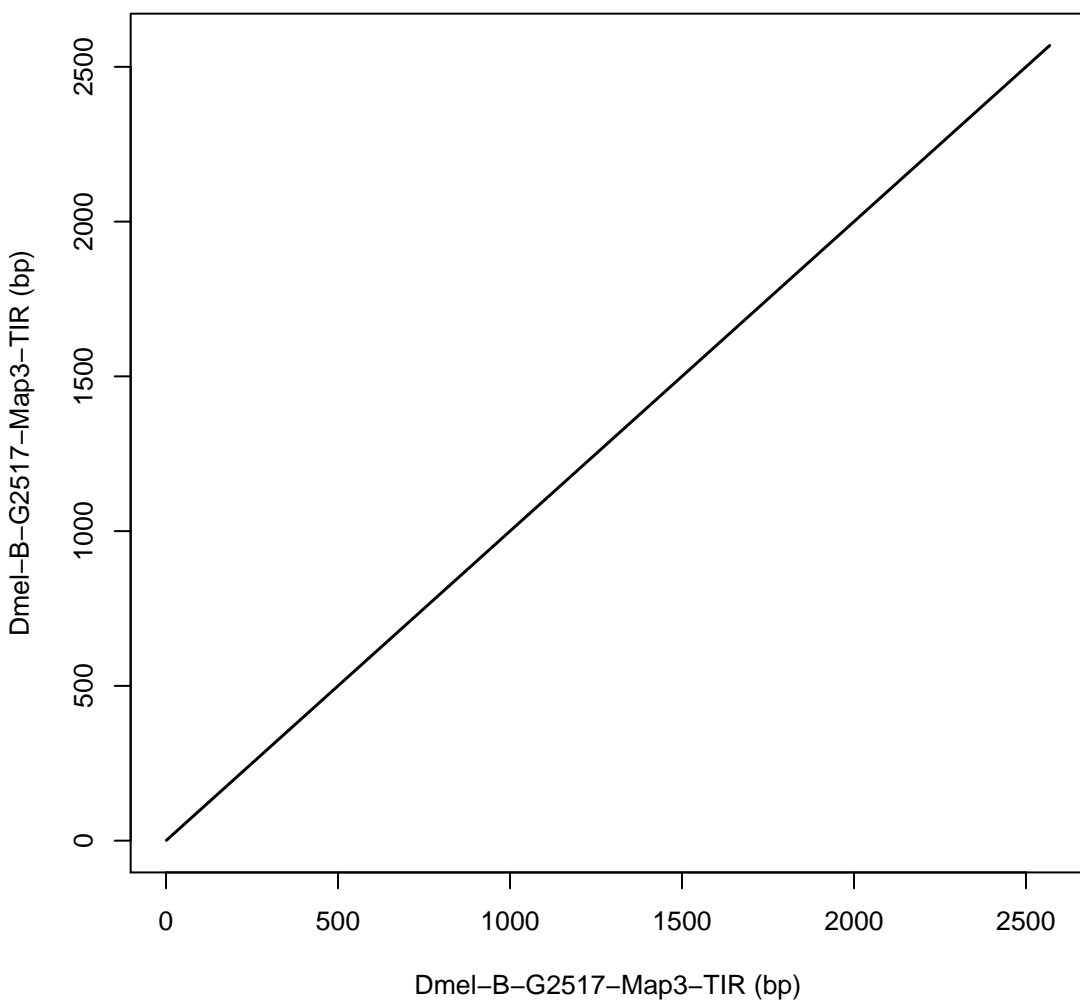
TE: Dmel-B-G2517-Map3-TIR
consensus size: 2569bp; fragments: 553; full length: 4 (≥ 2312.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

