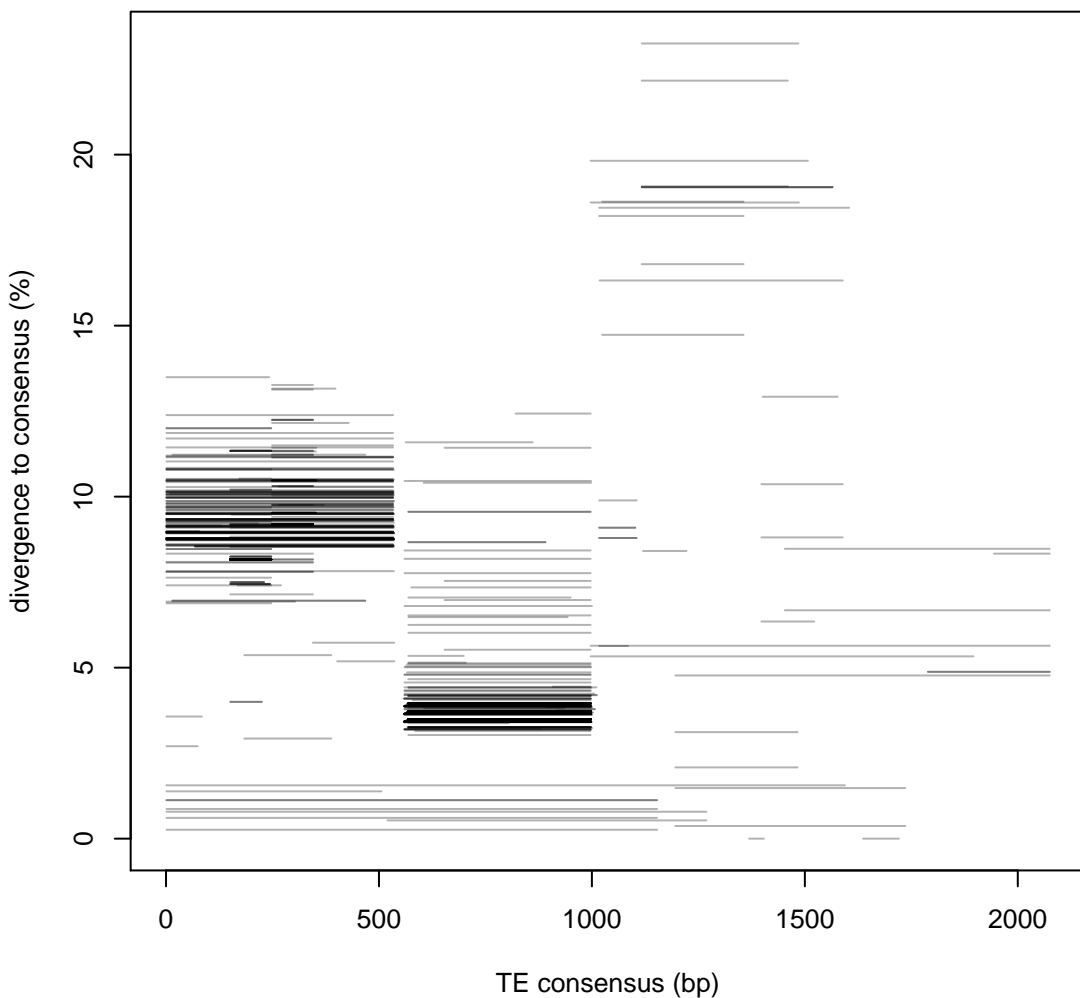
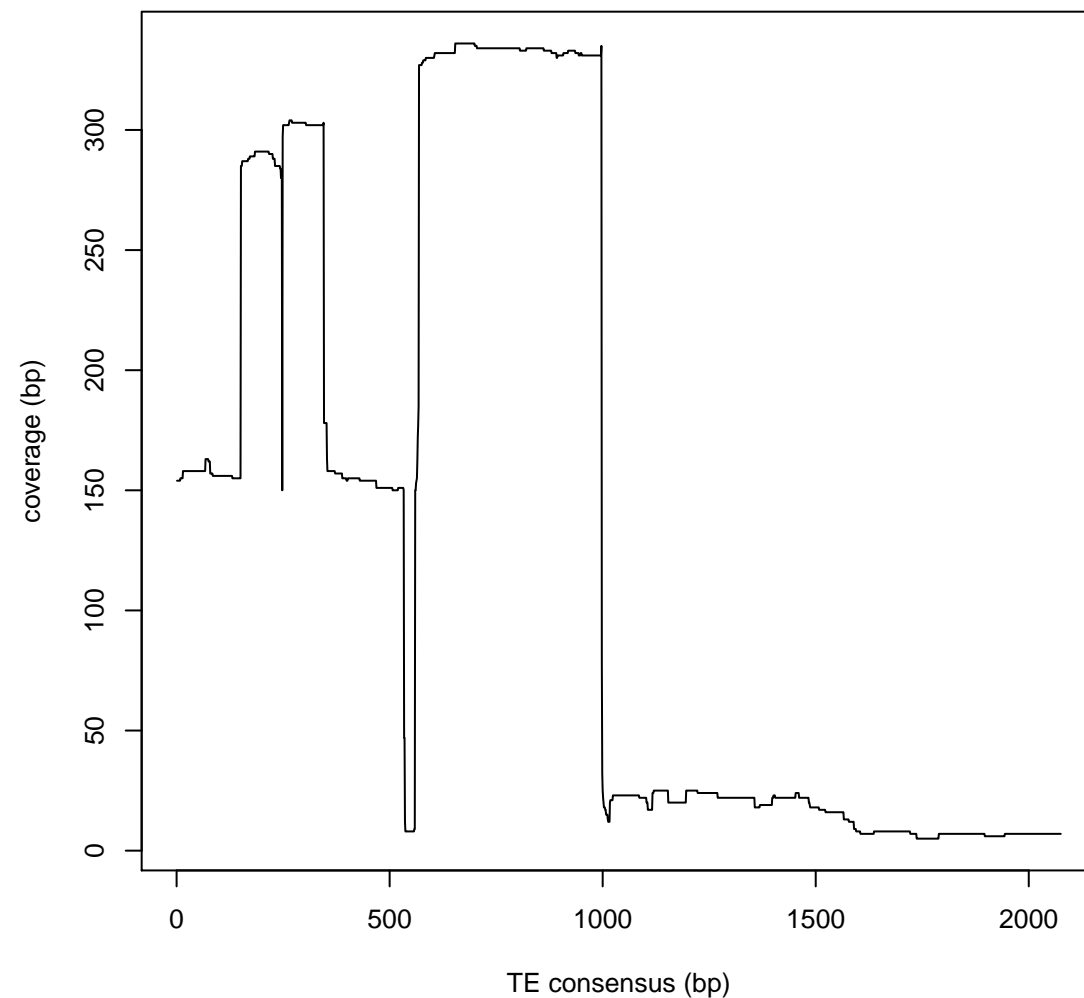


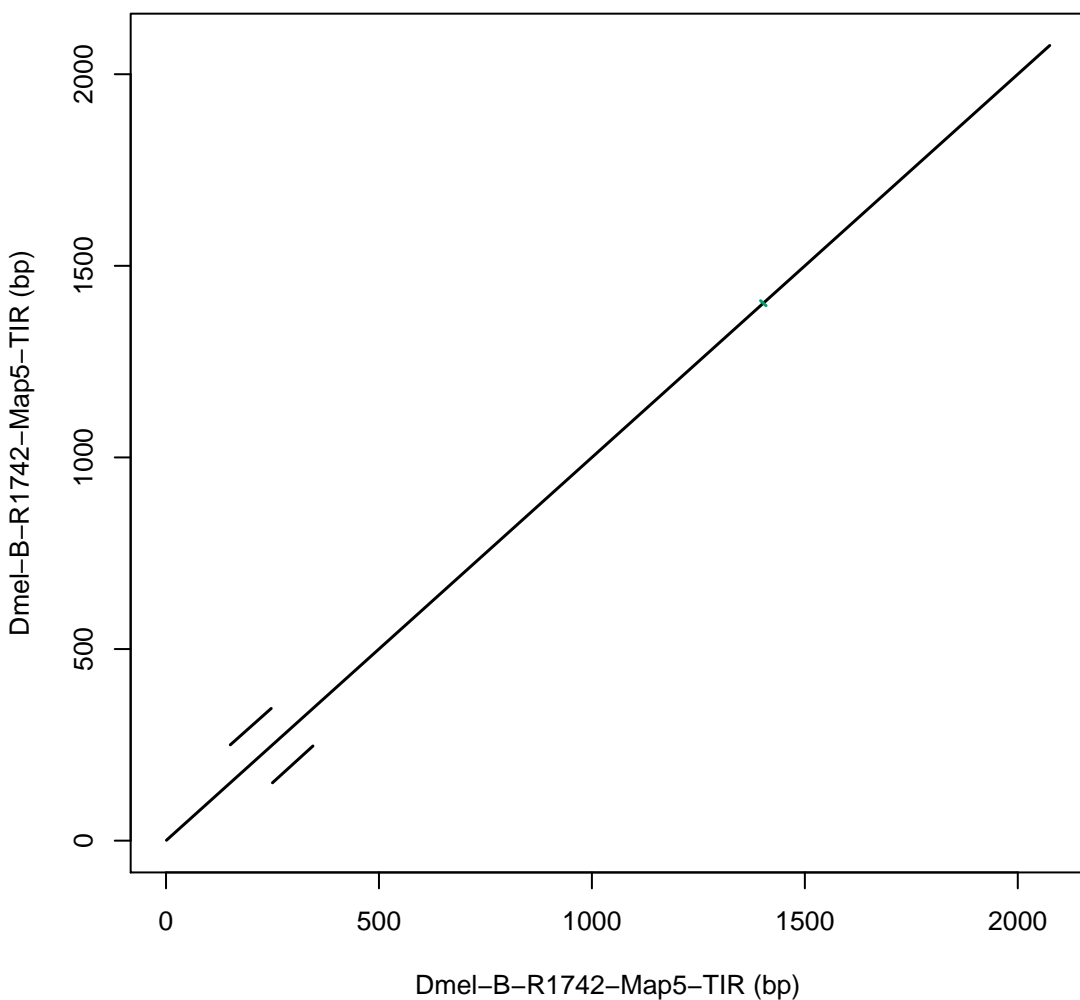
TE: Dmel-B-R1742-Map5-TIR  
consensus size: 2075bp; fragments: 828; full length: 0 (>=1867.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

