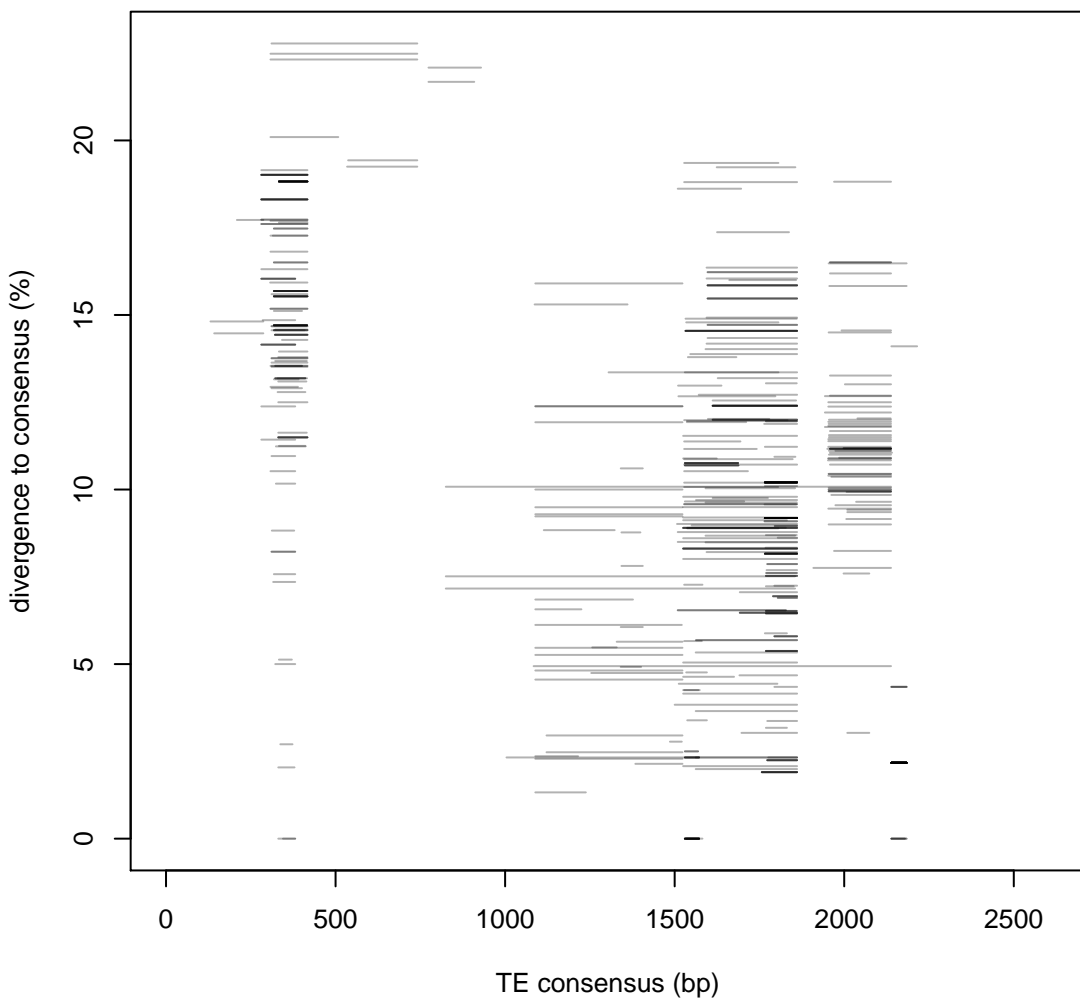
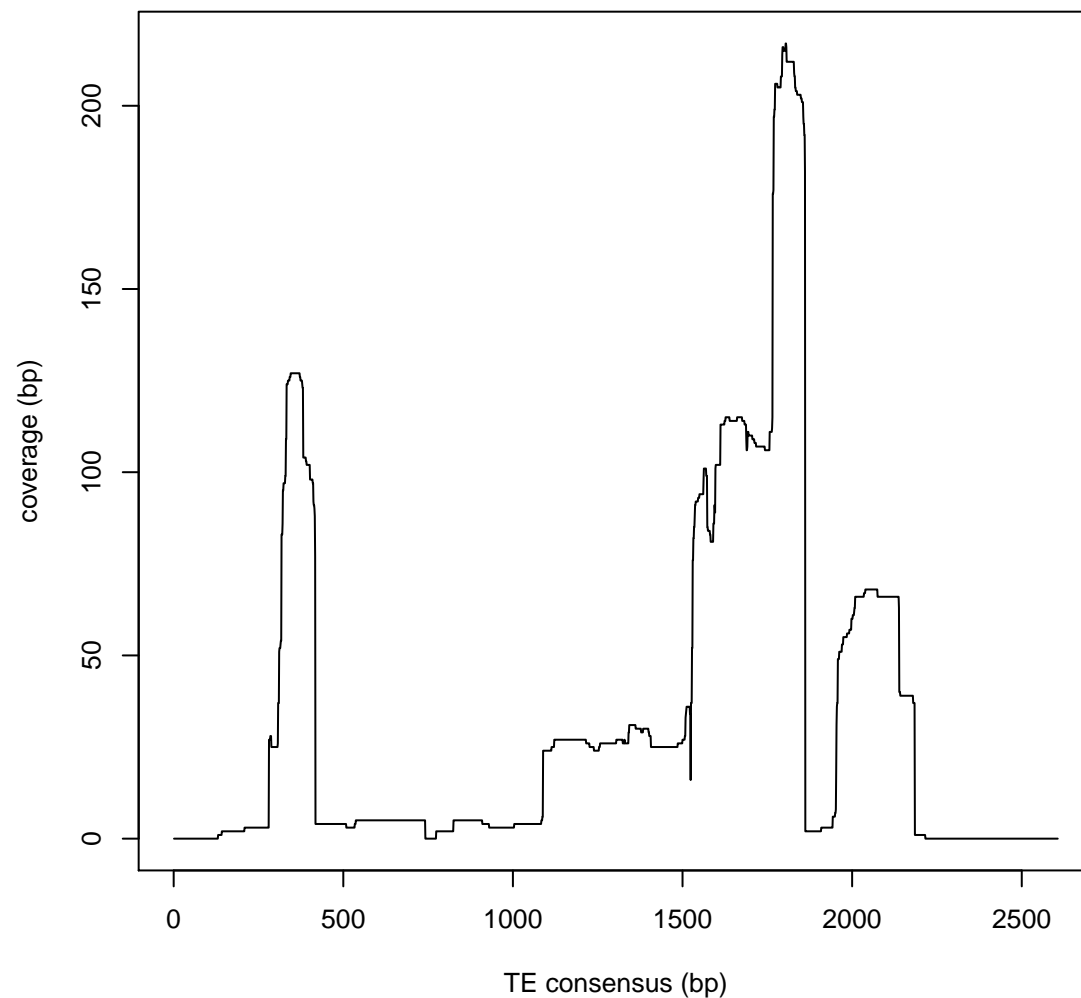


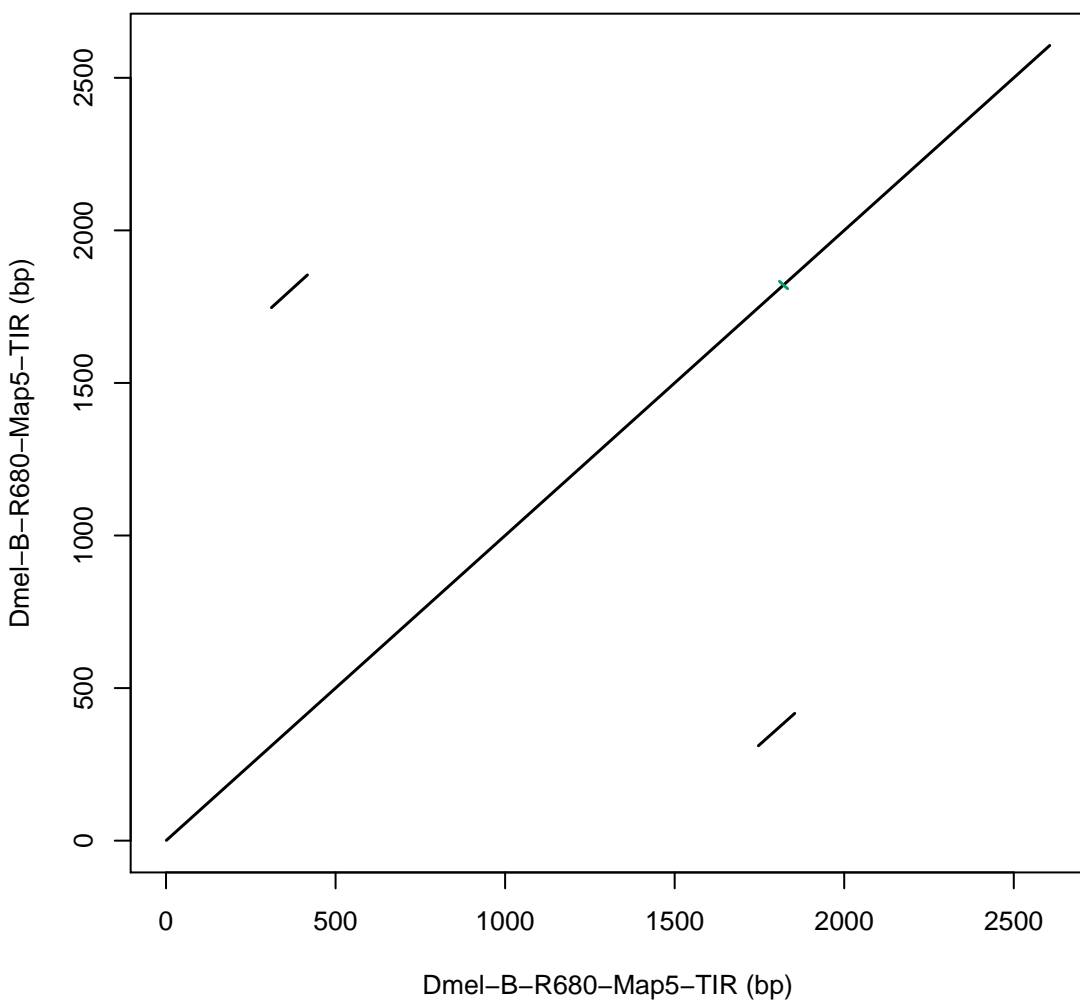
TE: Dmel-B-R680-Map5-TIR  
consensus size: 2606bp; fragments: 530; full length: 0 (>=2345.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

