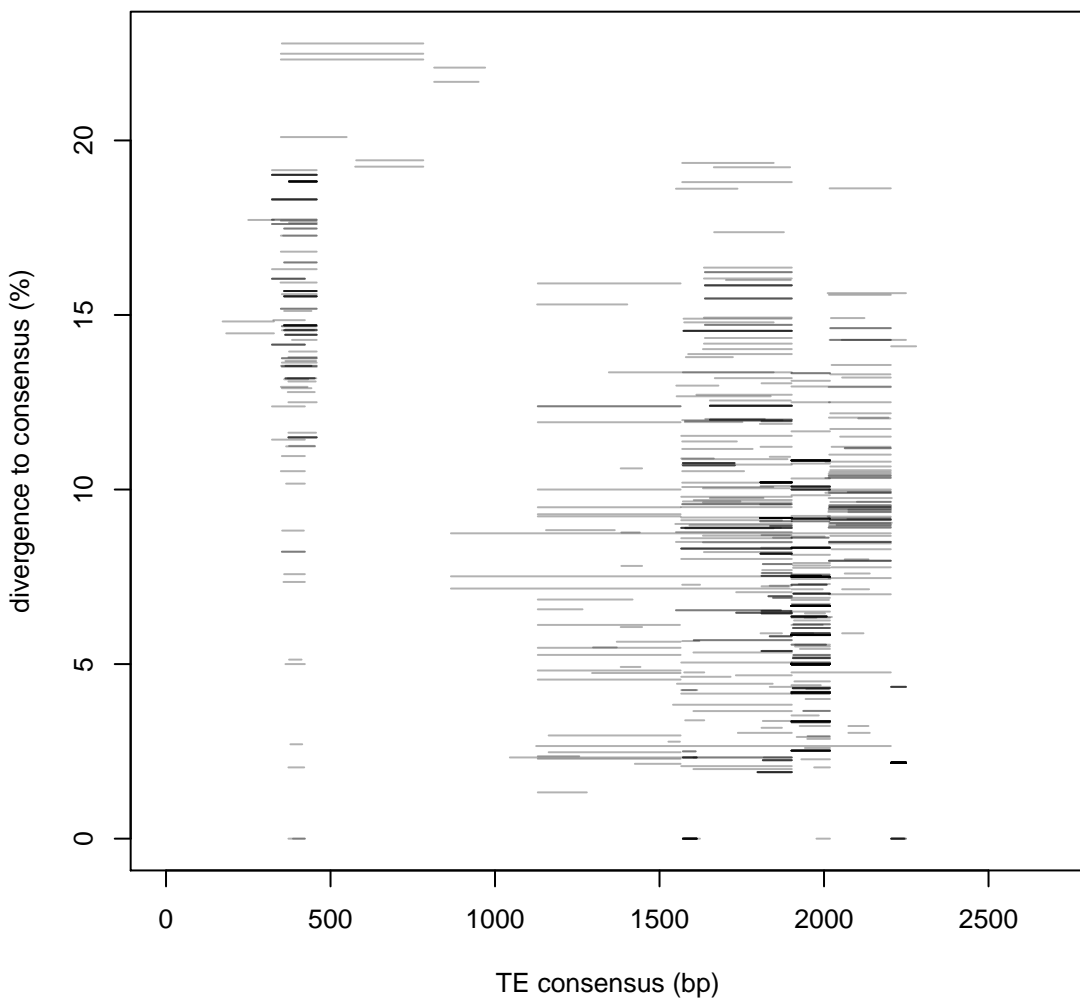
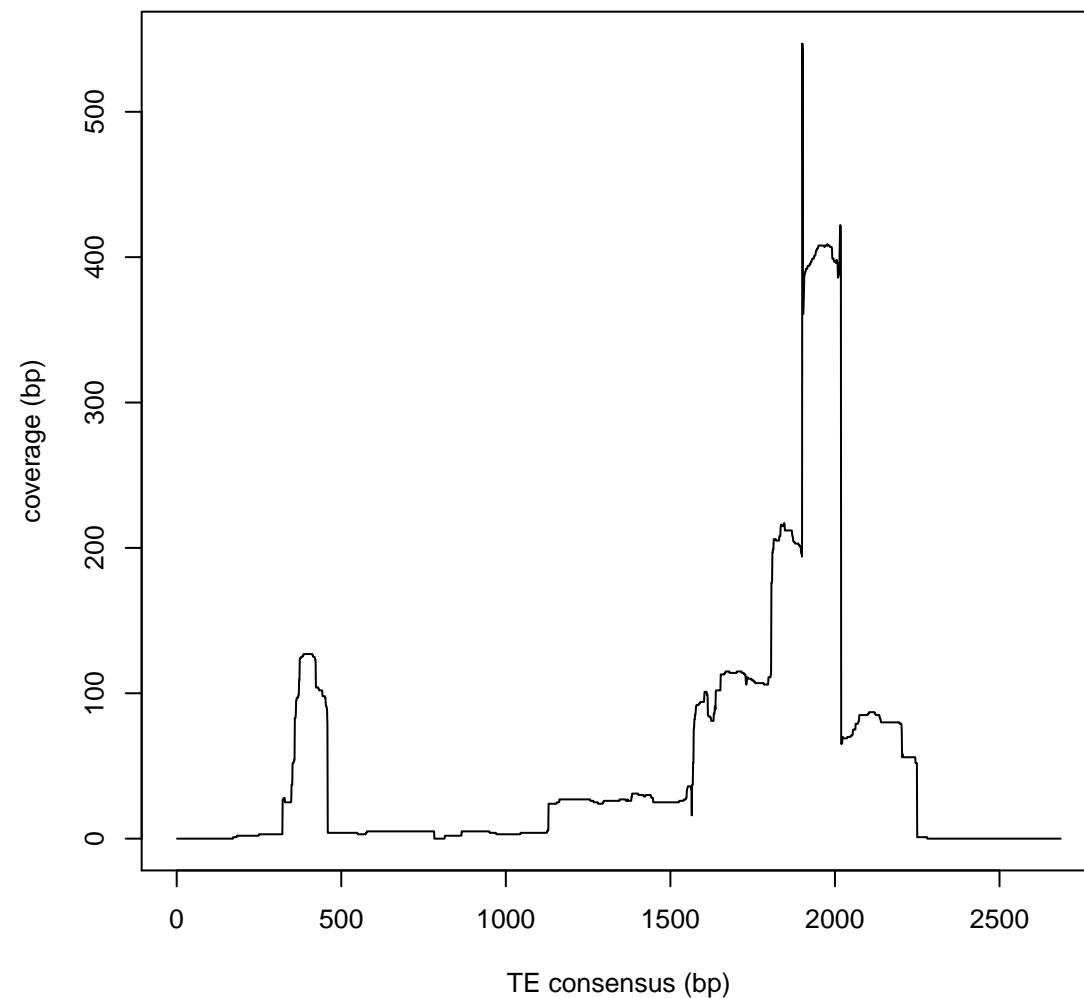


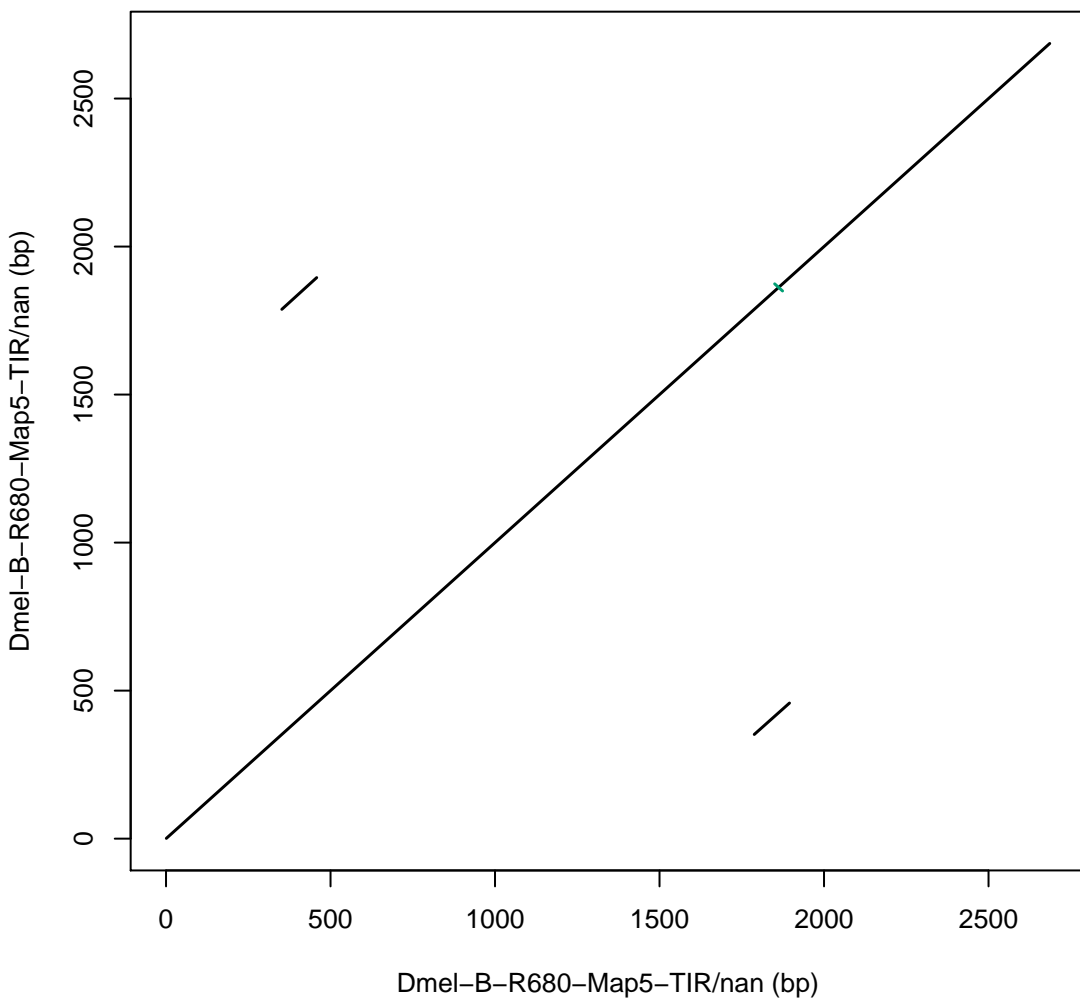
TE: Dmel-B-R680-Map5-TIR/nan  
consensus size: 2686bp; fragments: 972; full length: 0 (>=2417.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

