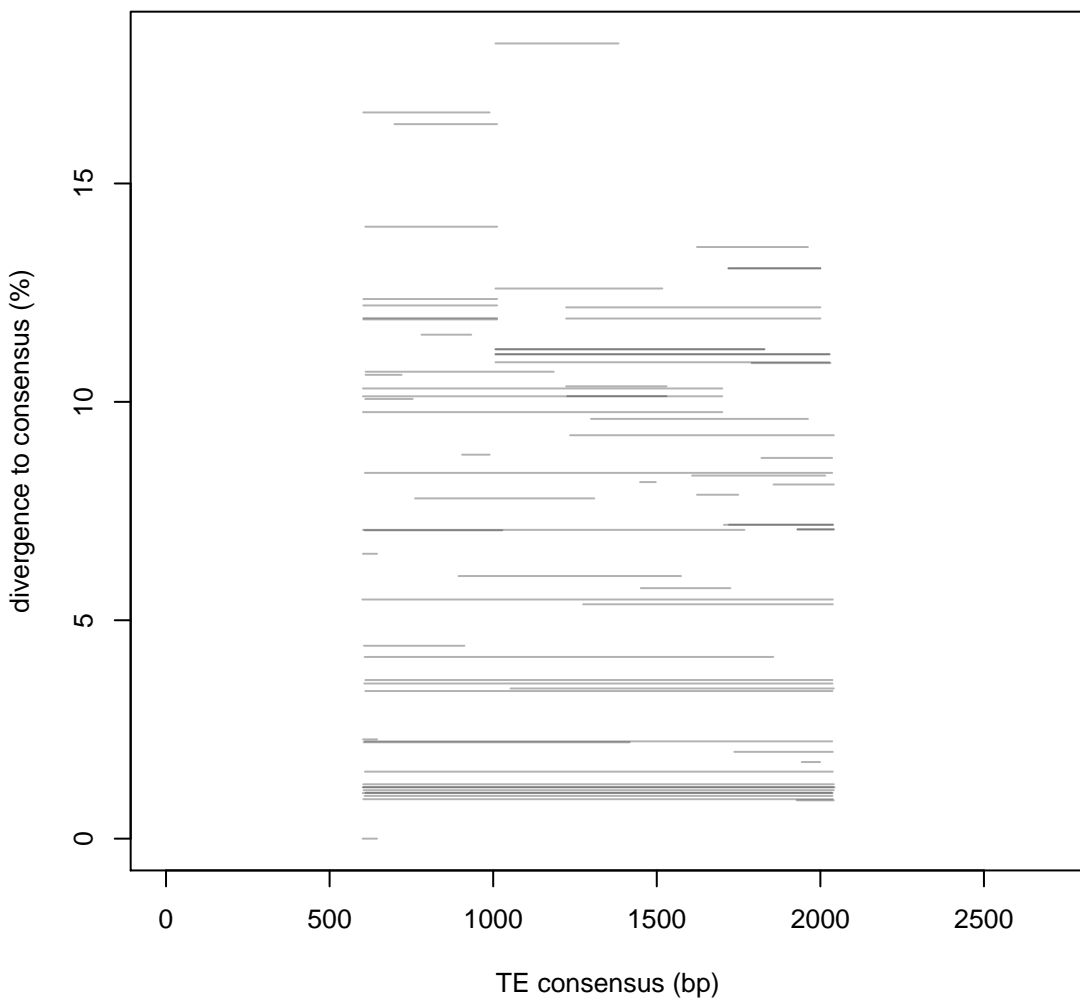
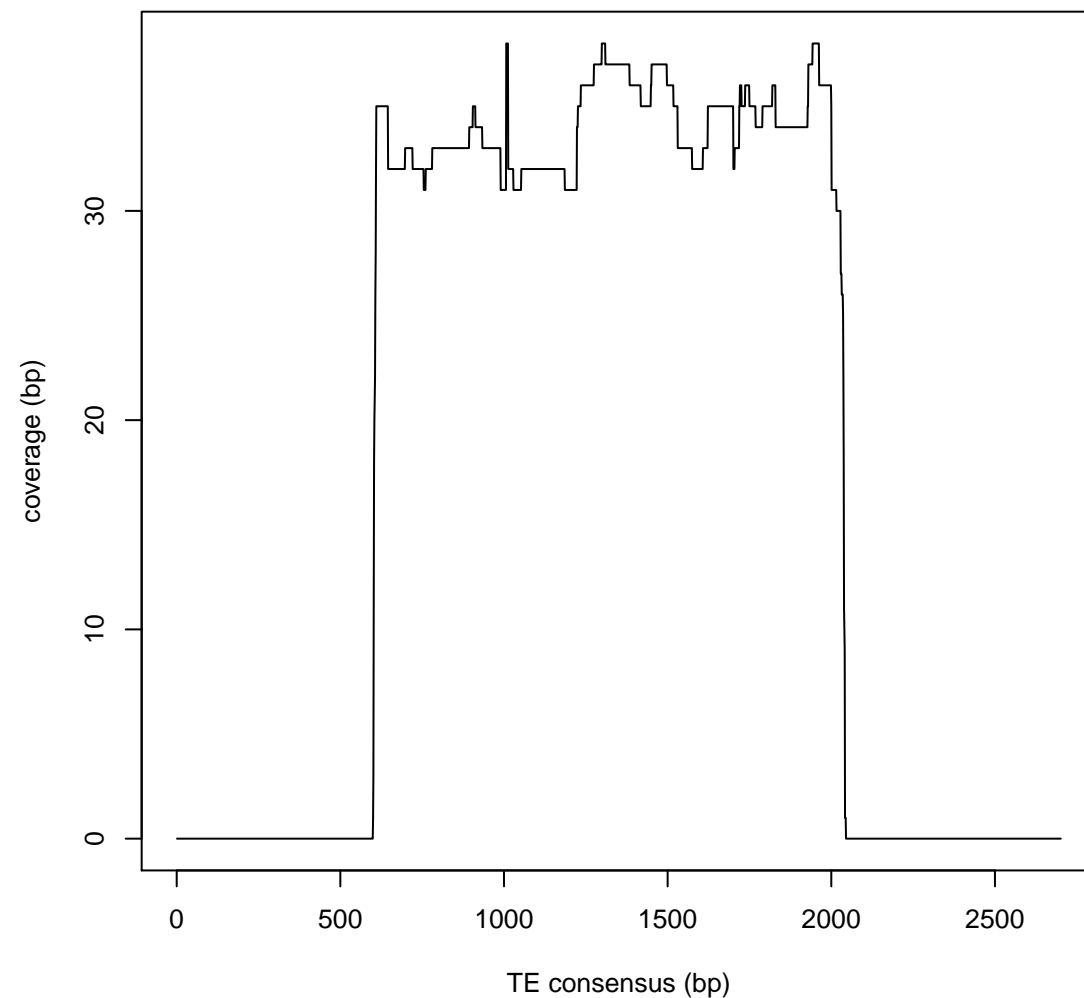


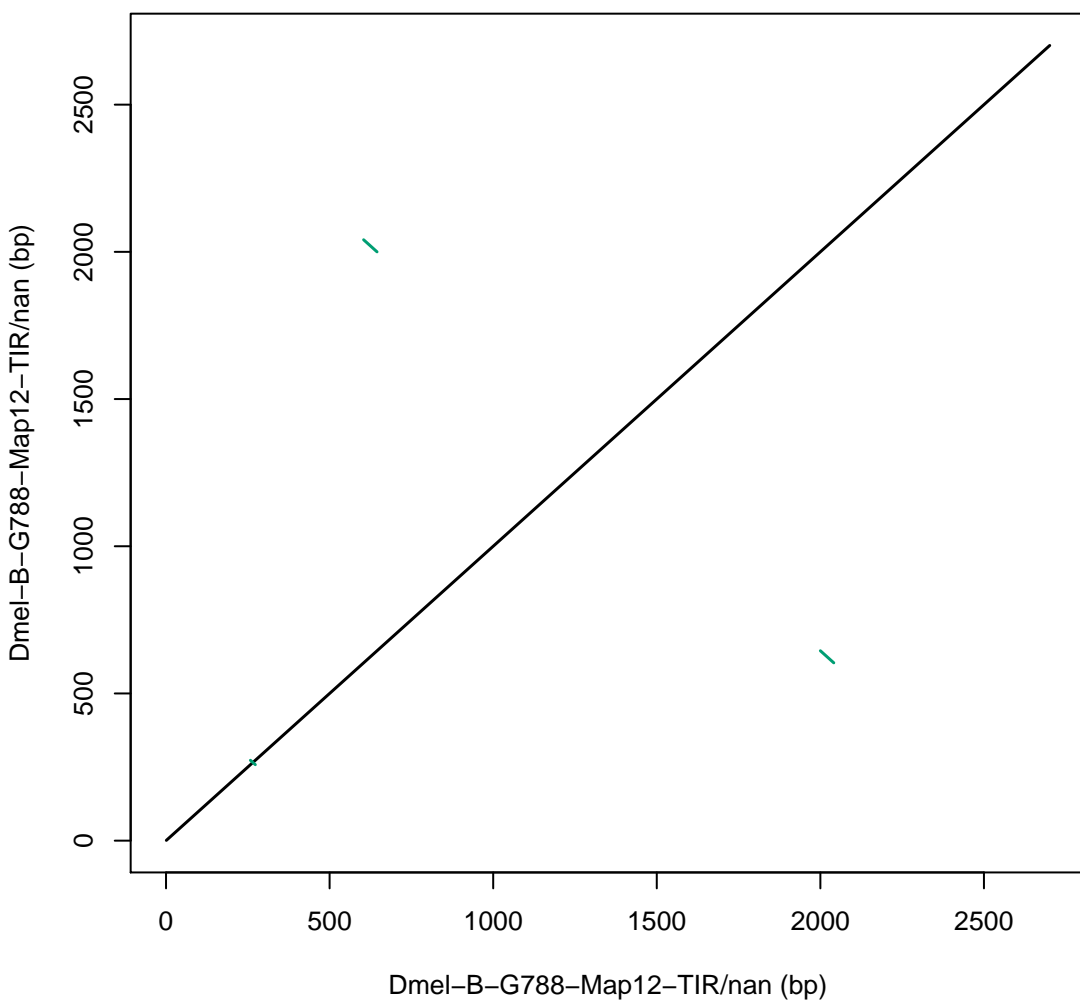
TE: Dmel-B-G788-Map12-TIR/nan  
consensus size: 2701bp; fragments: 72; full length: 0 ( $\geq 2430.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

