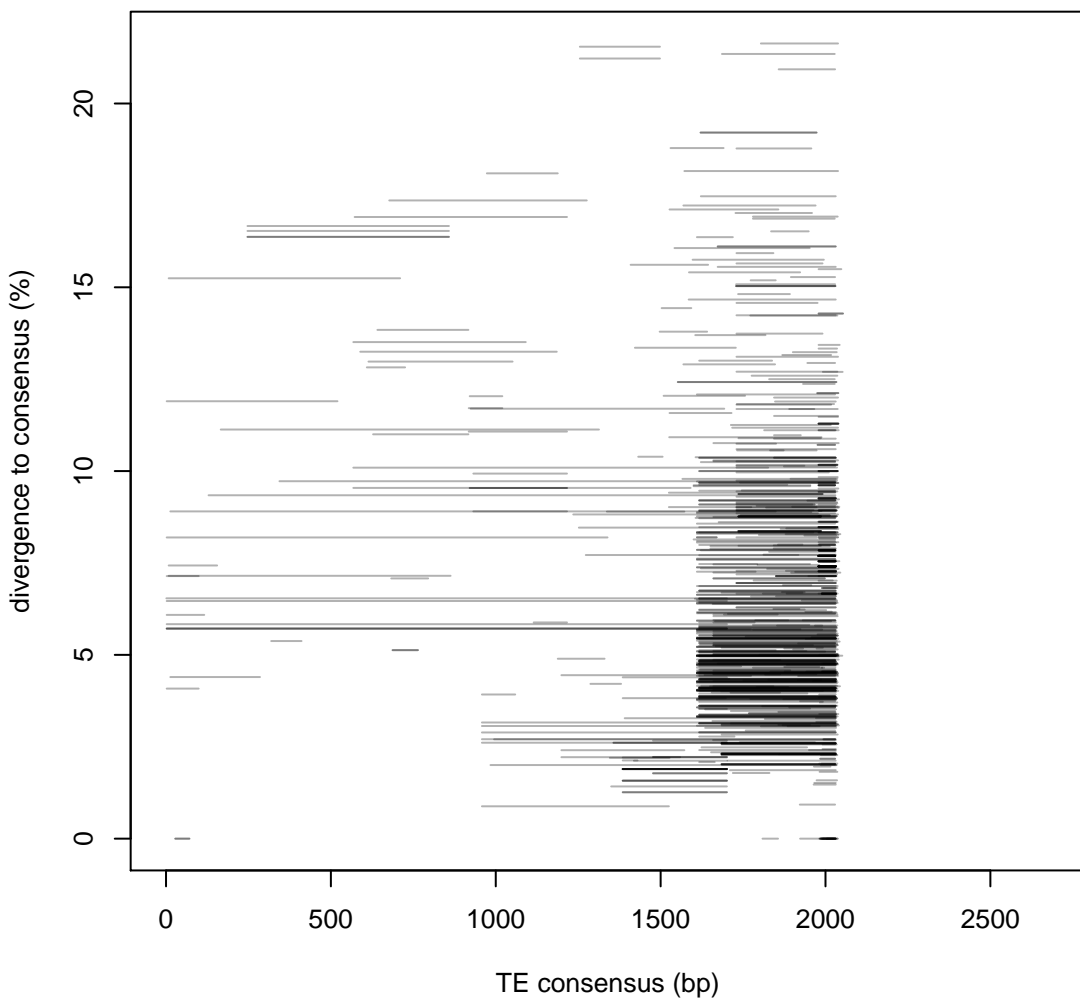
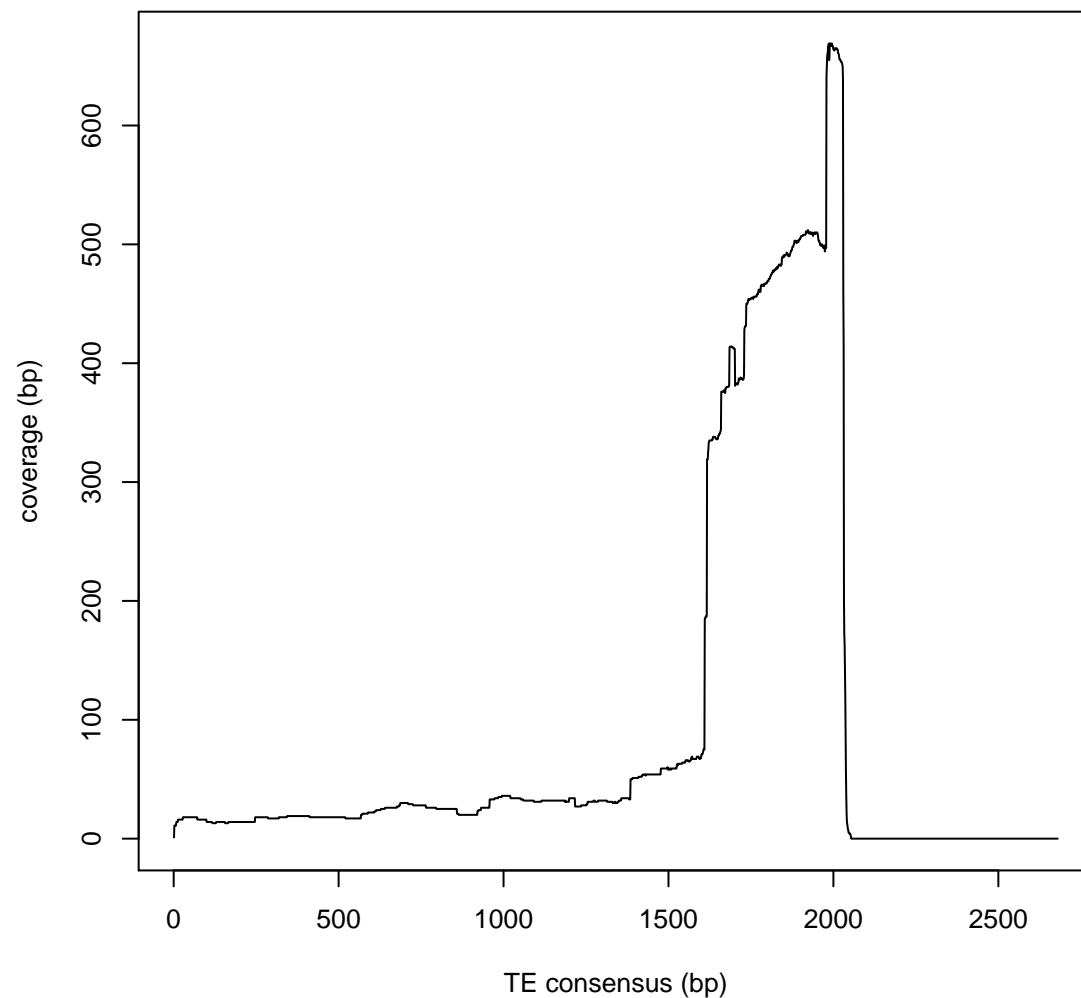


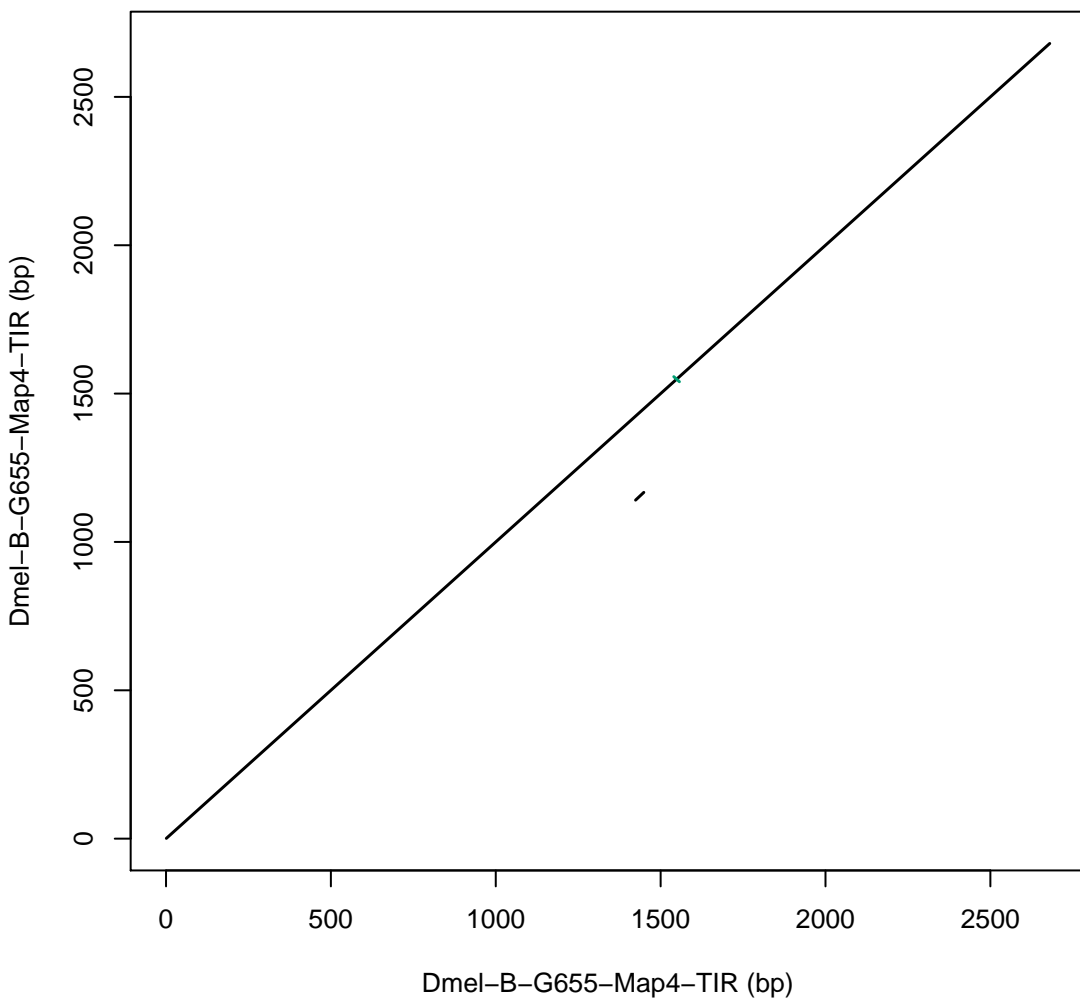
TE: Dmel-B-G655-Map4-TIR  
consensus size: 2680bp; fragments: 872; full length: 0 (>=2412bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

