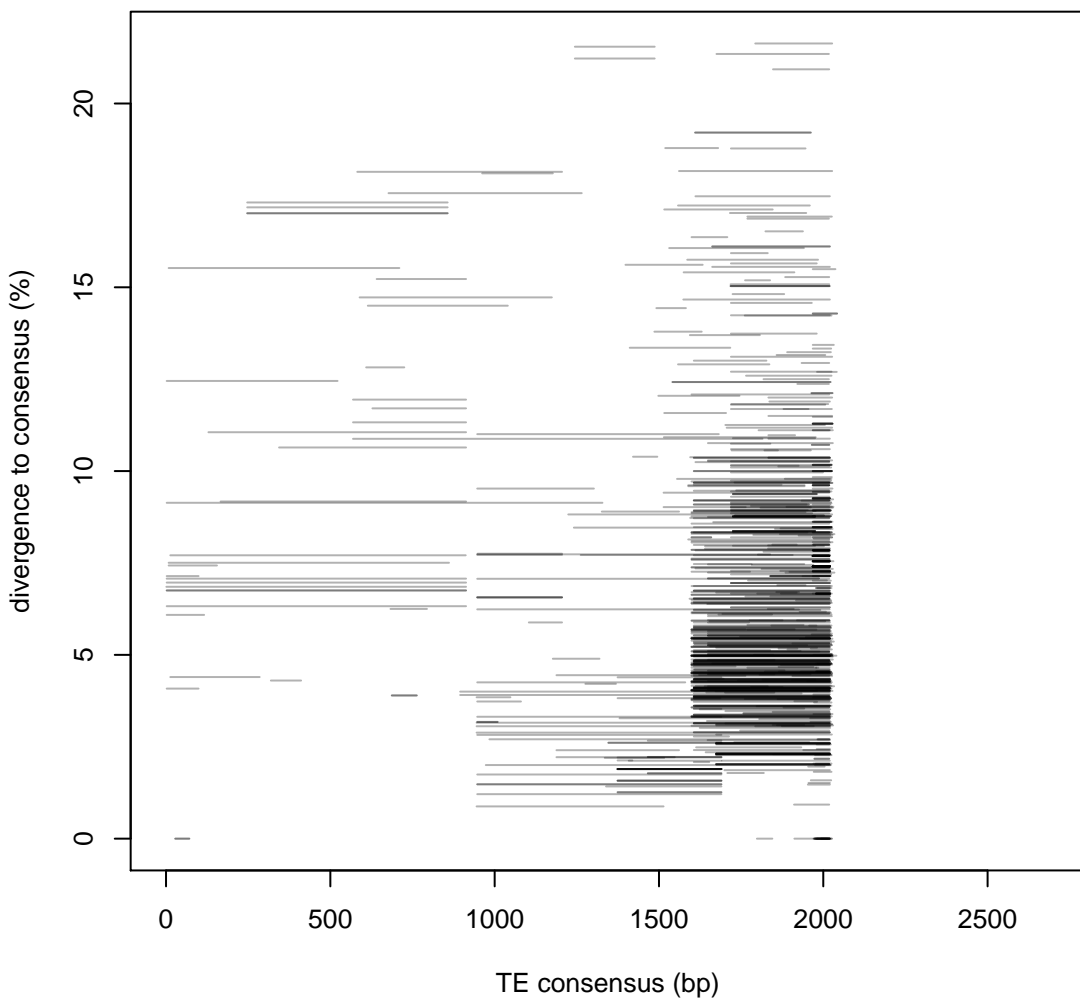
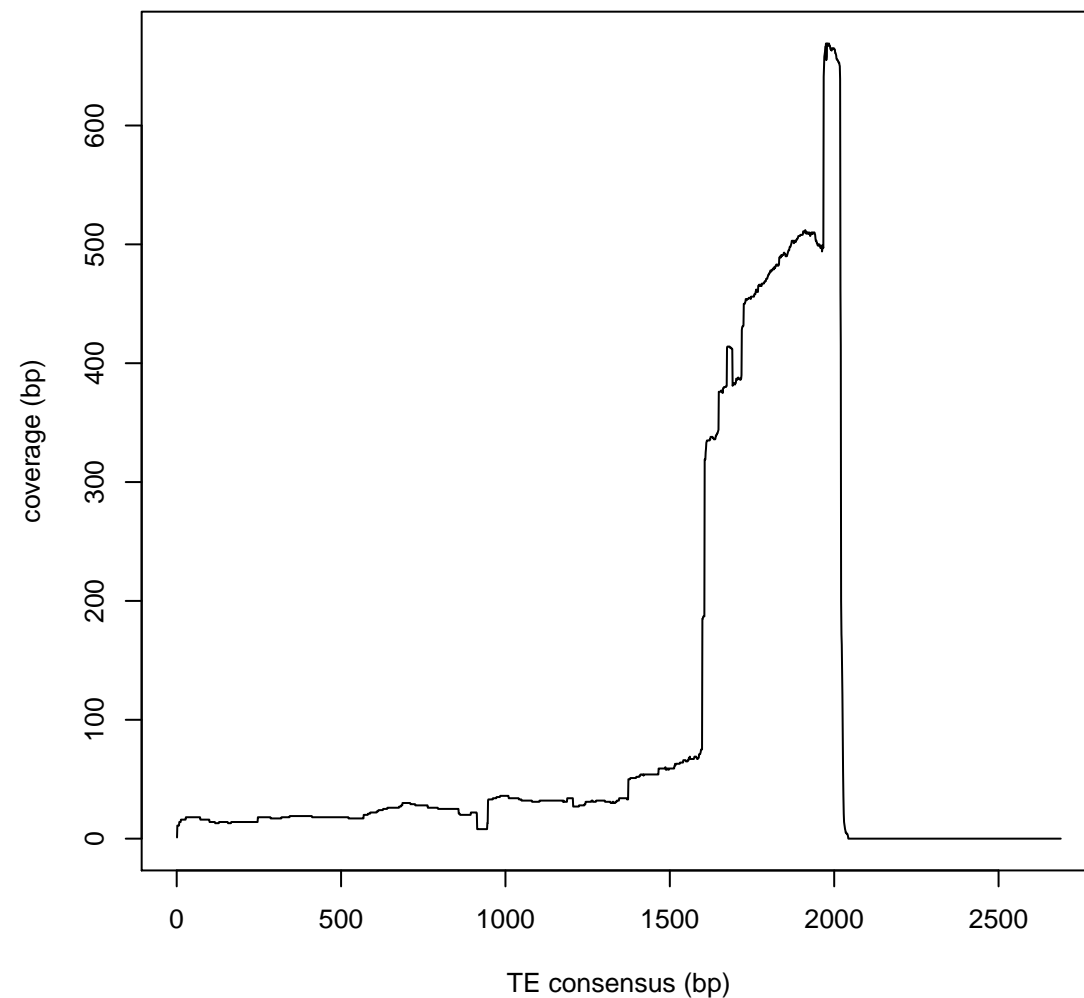


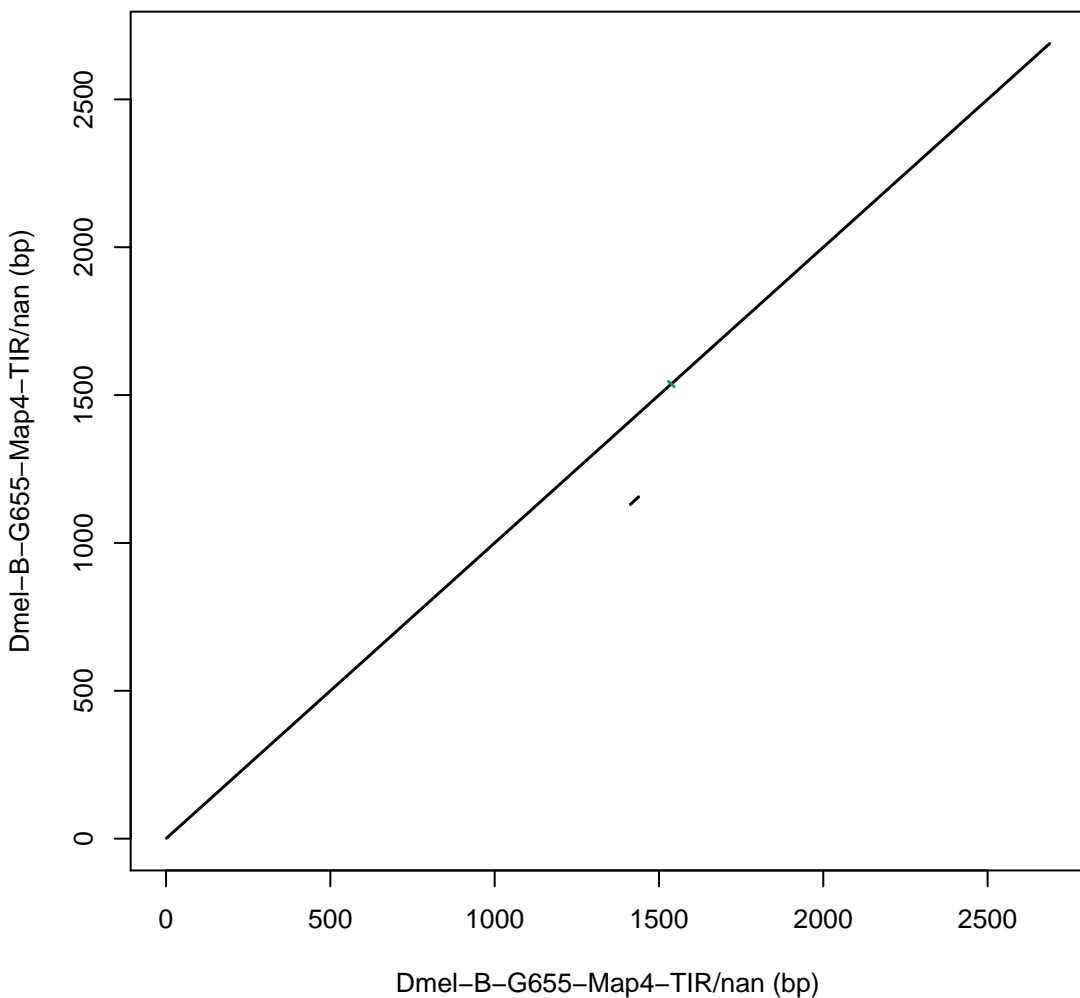
TE: Dmel-B-G655-Map4-TIR/nan
consensus size: 2689bp; fragments: 884; full length: 0 (>=2420.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

