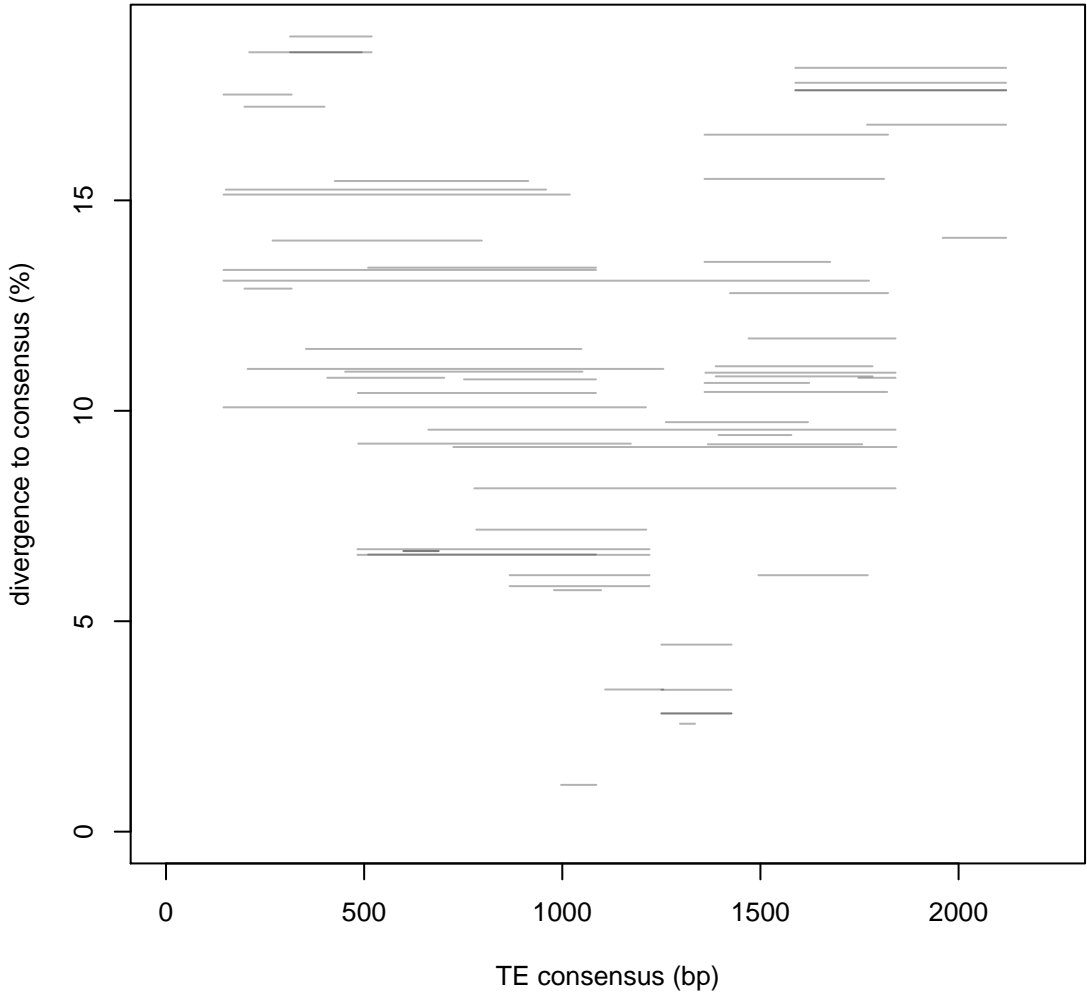
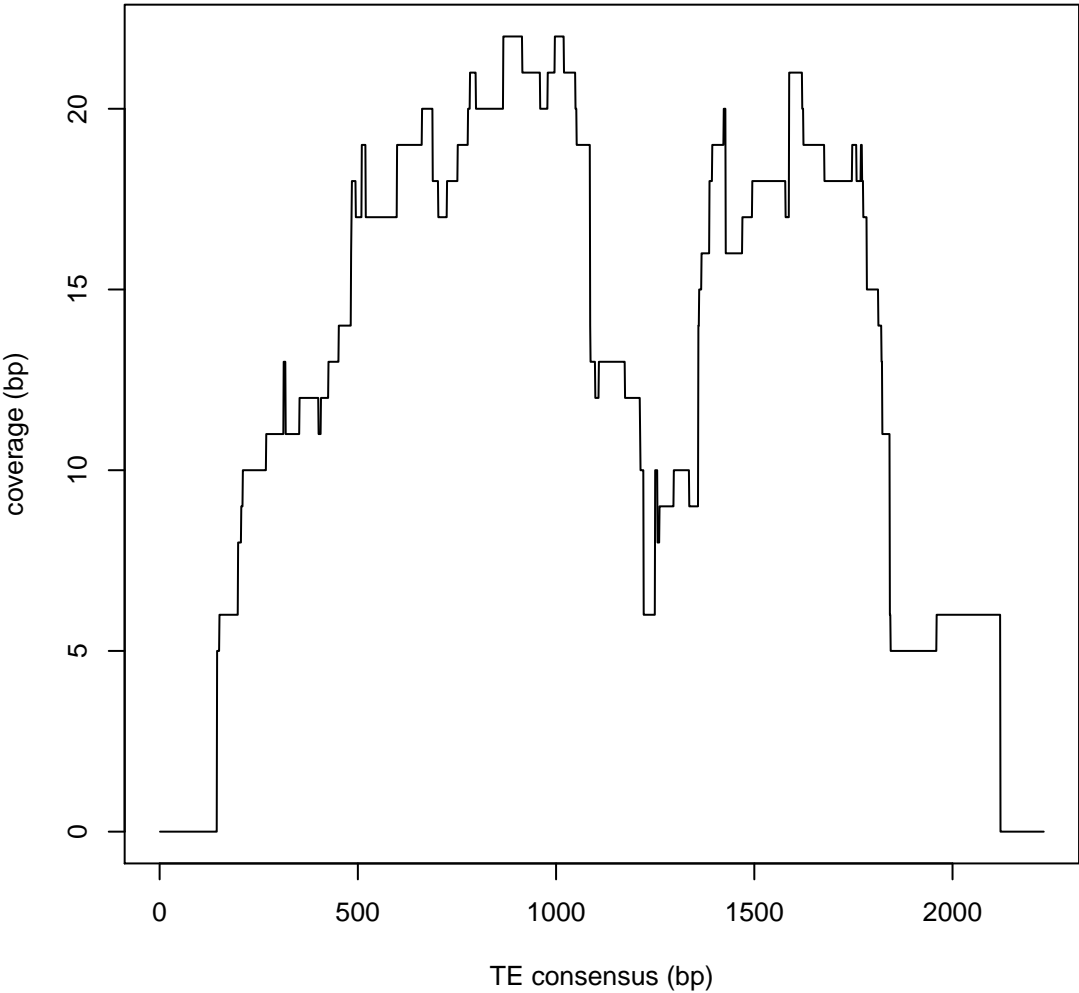


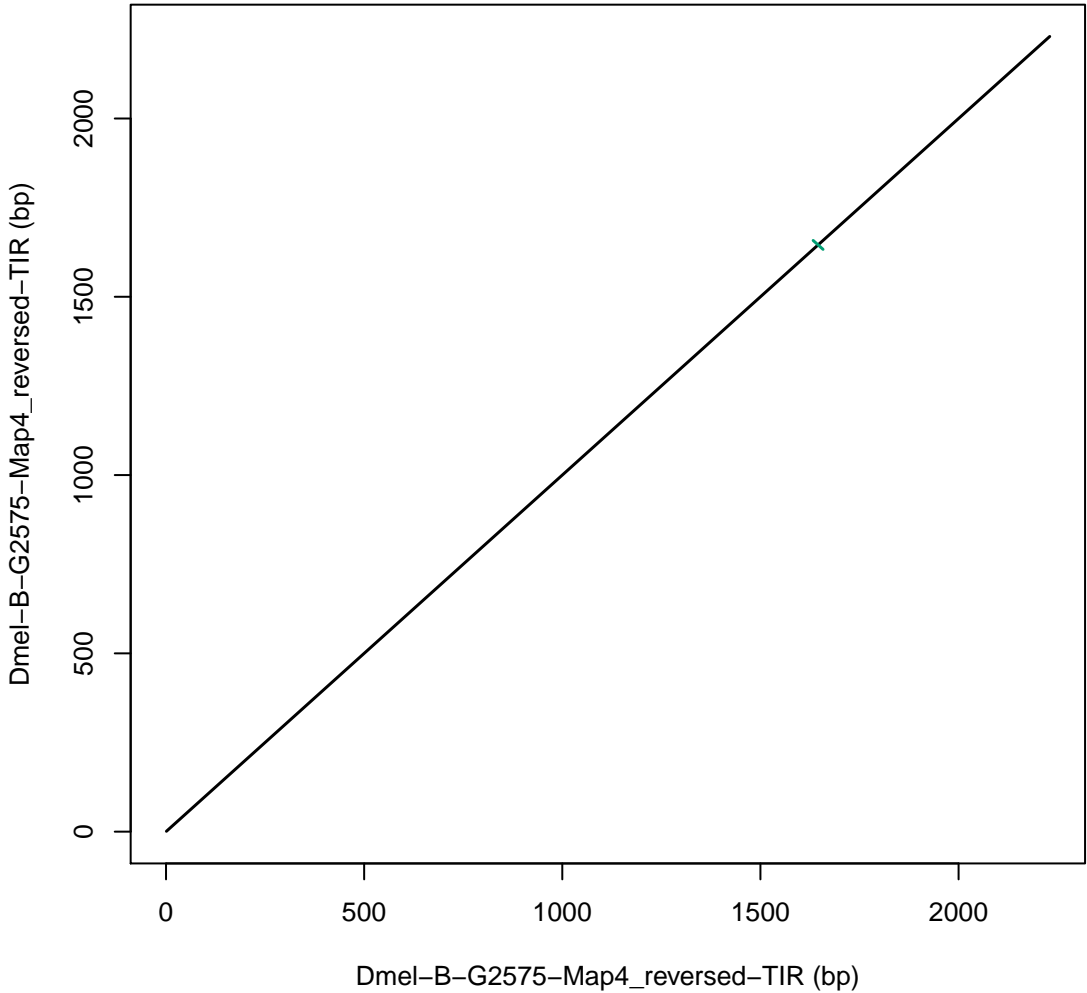
TE: Dmel-B-G2575-Map4_reversed-TIR
consensus size: 2230bp; fragments: 61; full length: 0 (>=2007bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

