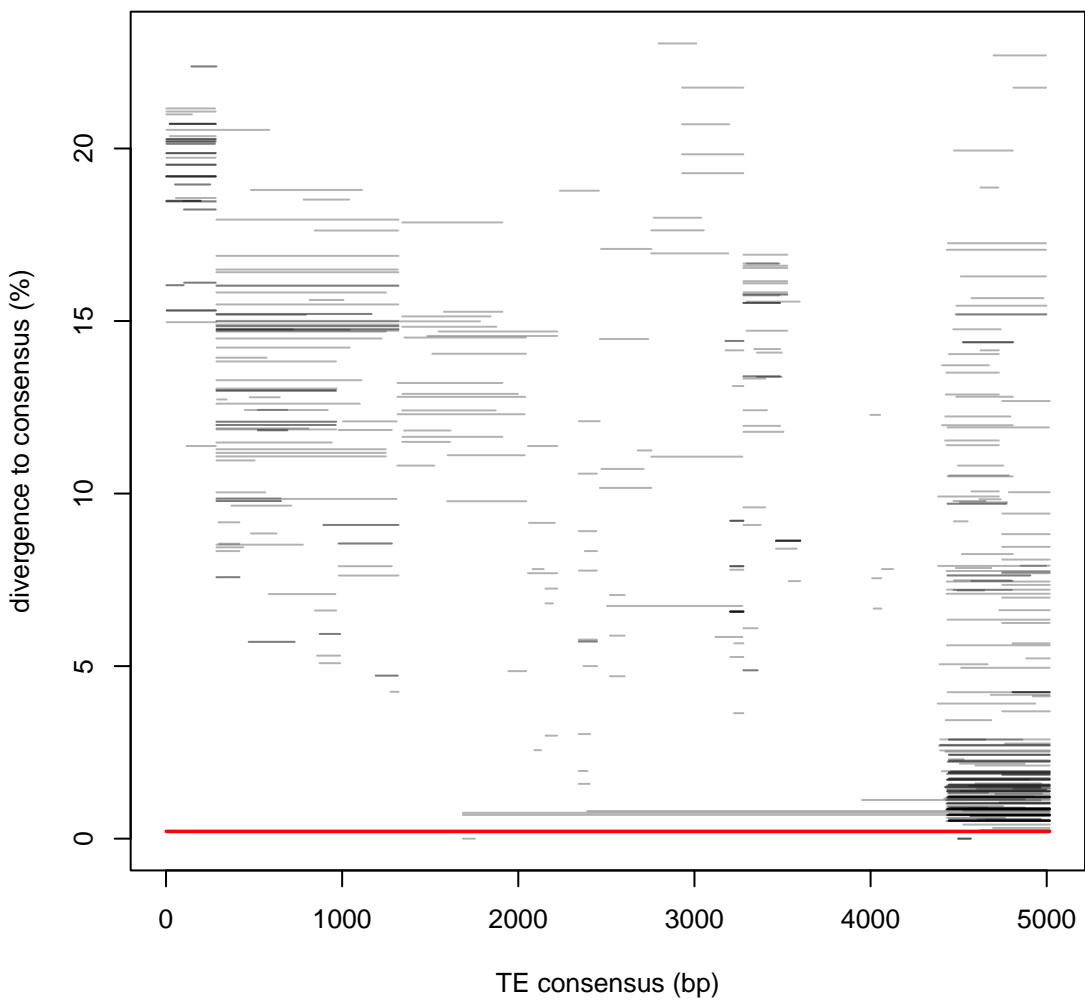
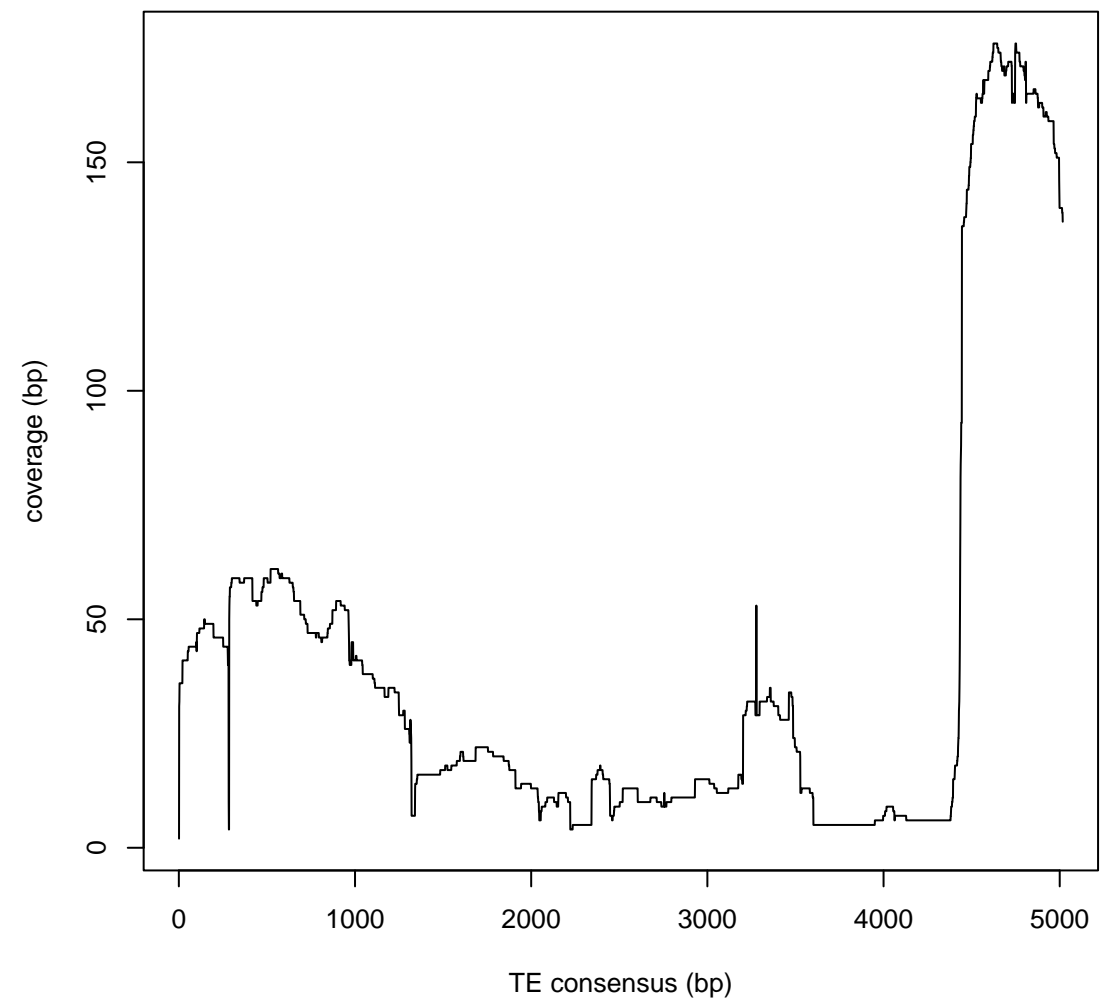


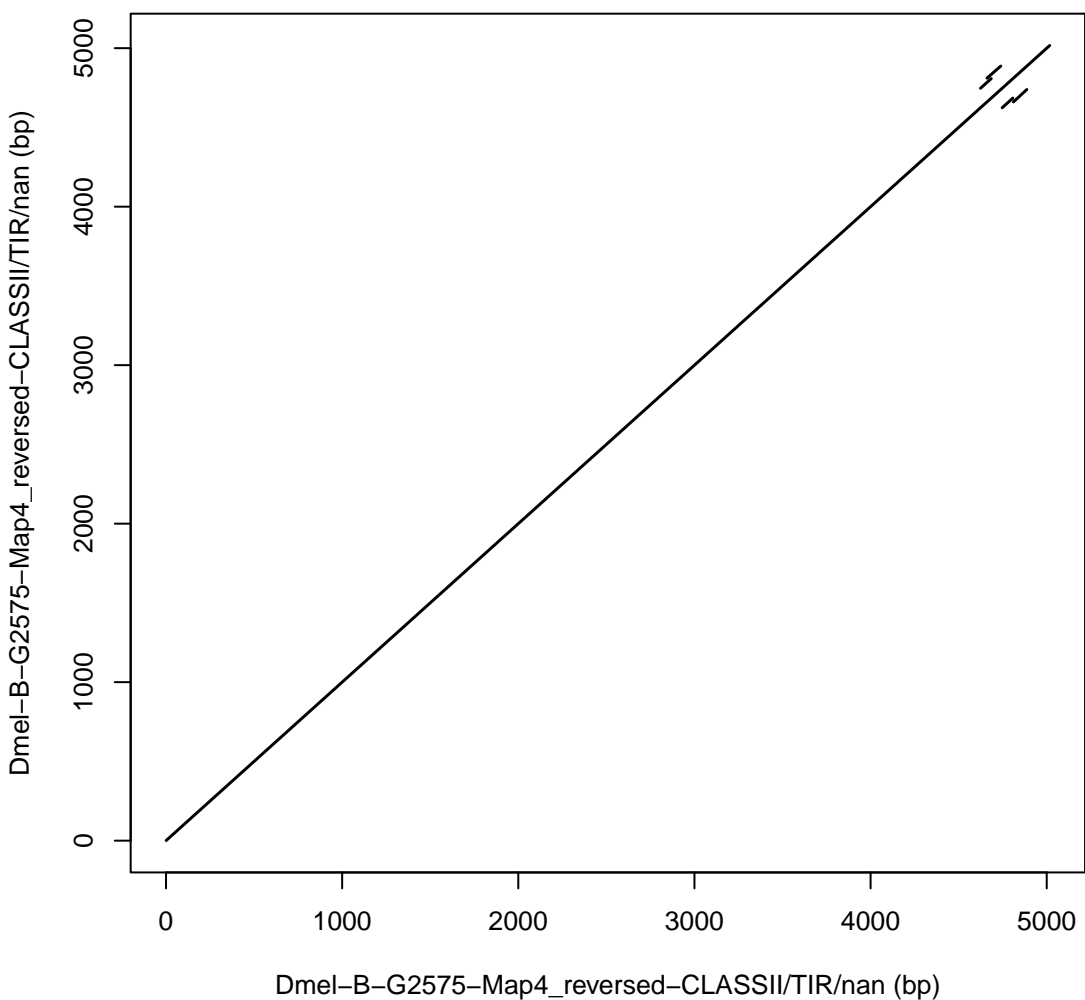
TE: Dmel-B-G2575-Map4_reversed-CLASSII/TIR/nan
consensus size: 5017bp; fragments: 473; full length: 2 (>=4515.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

