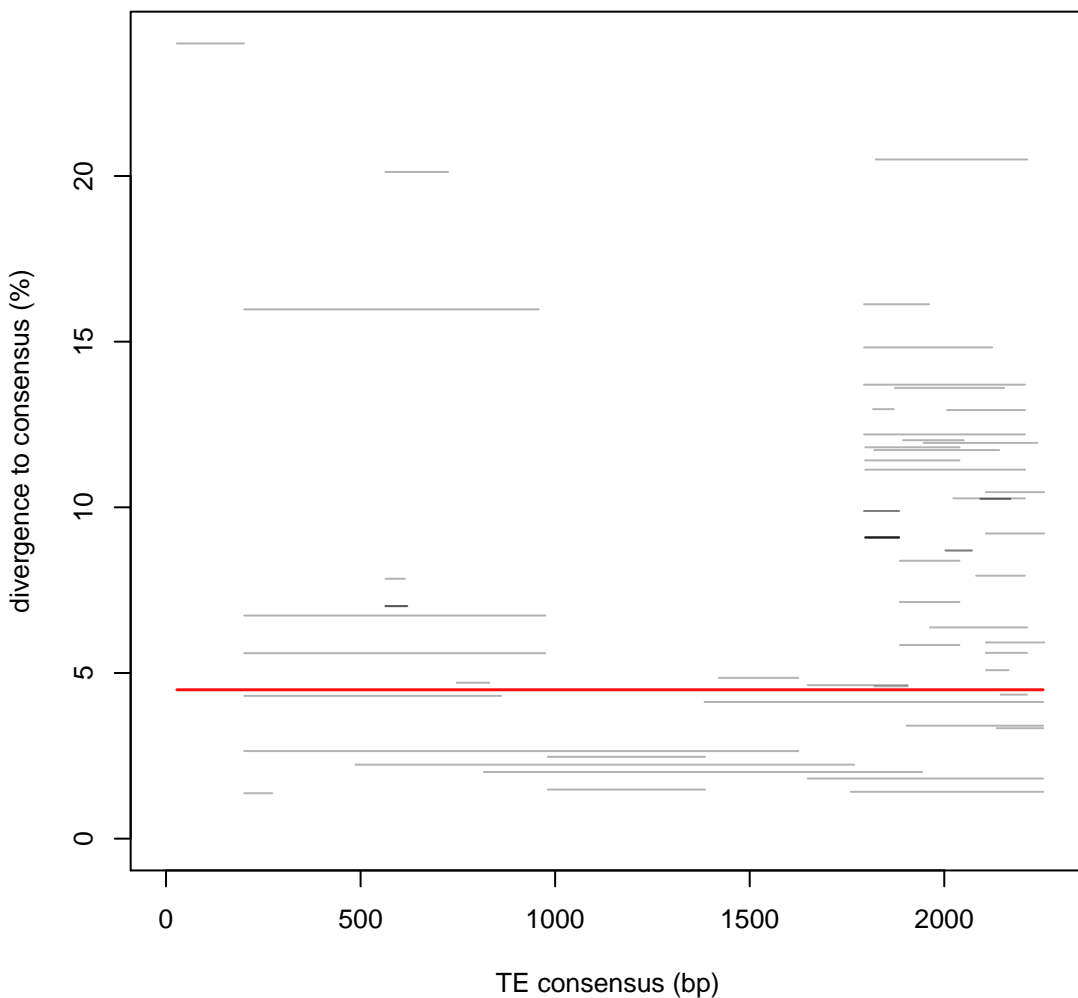
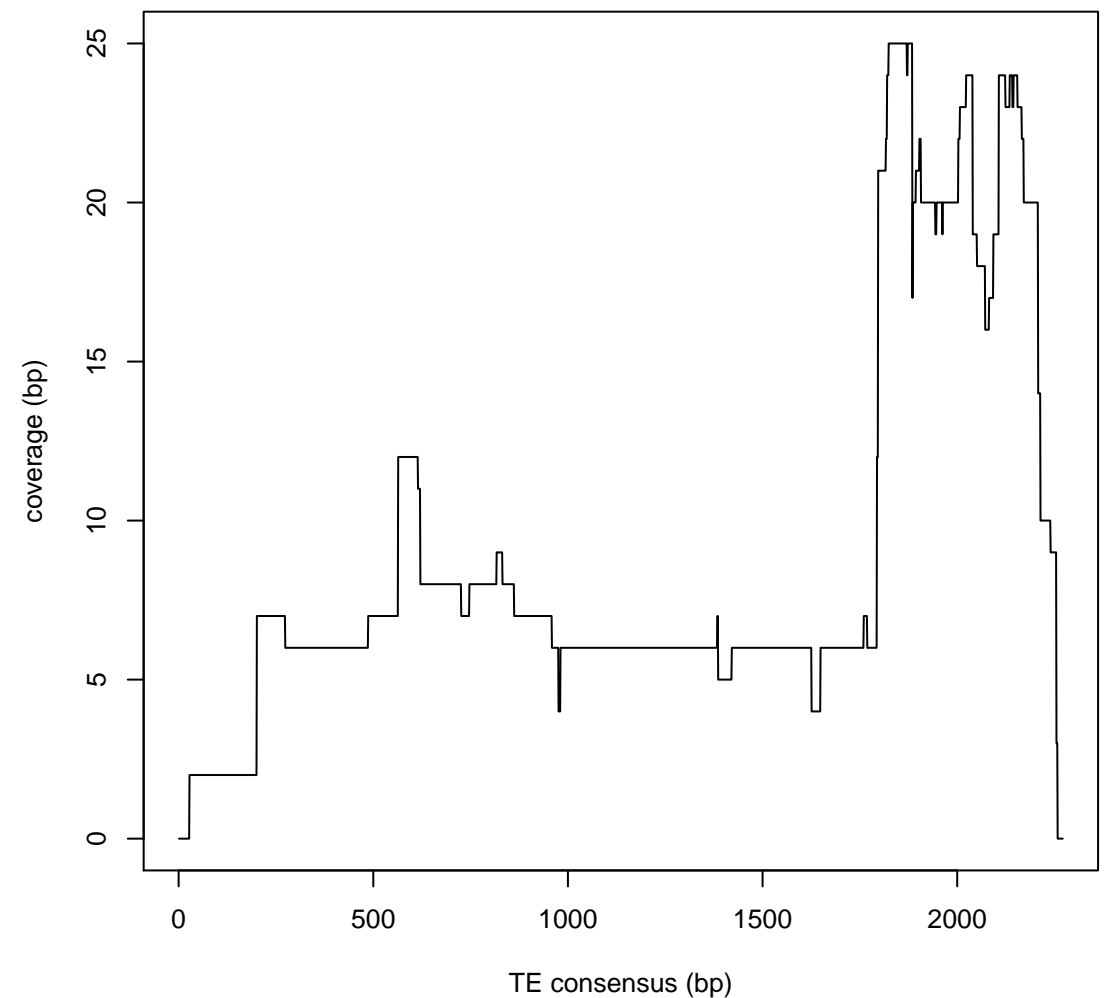


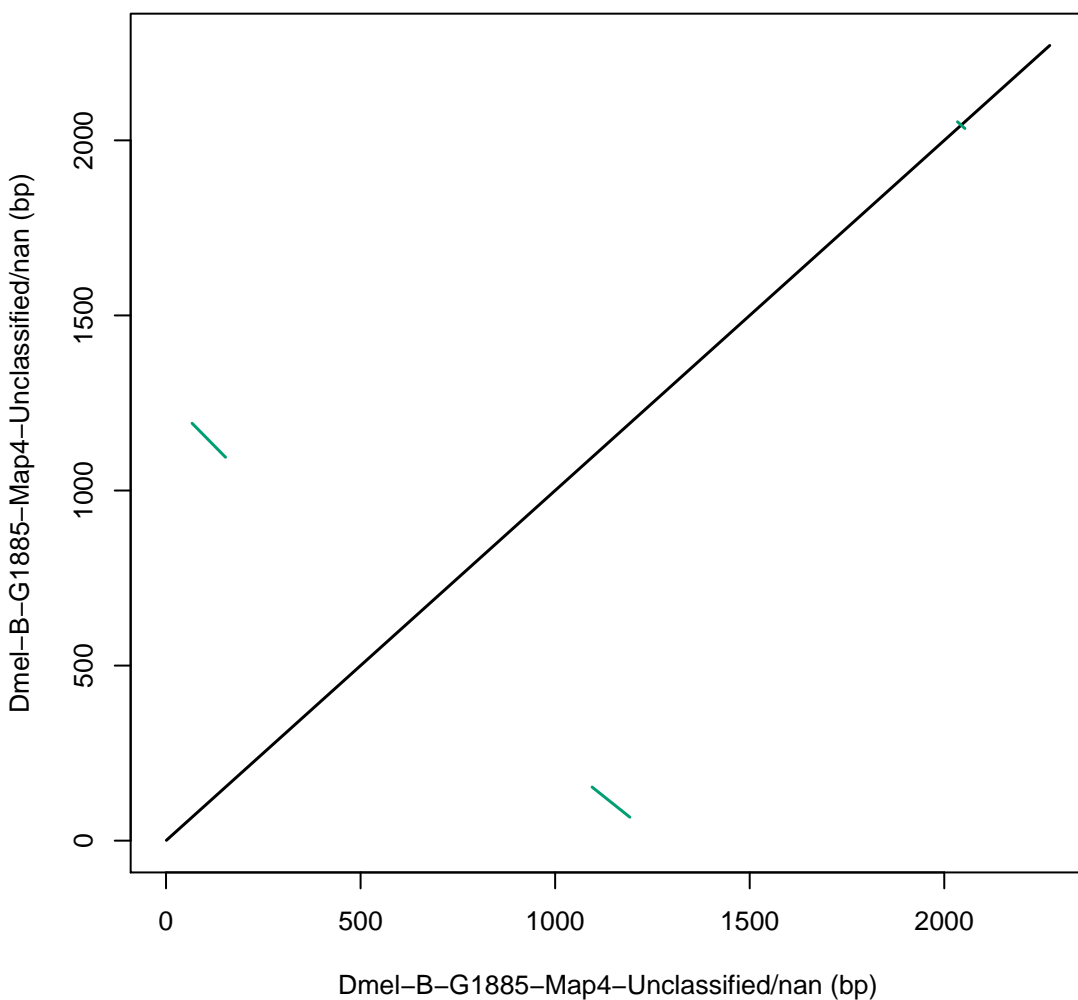
TE: Dmel-B-G1885-Map4-Unclassified/nan  
consensus size: 2271bp; fragments: 64; full length: 1 ( $\geq 2043.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

