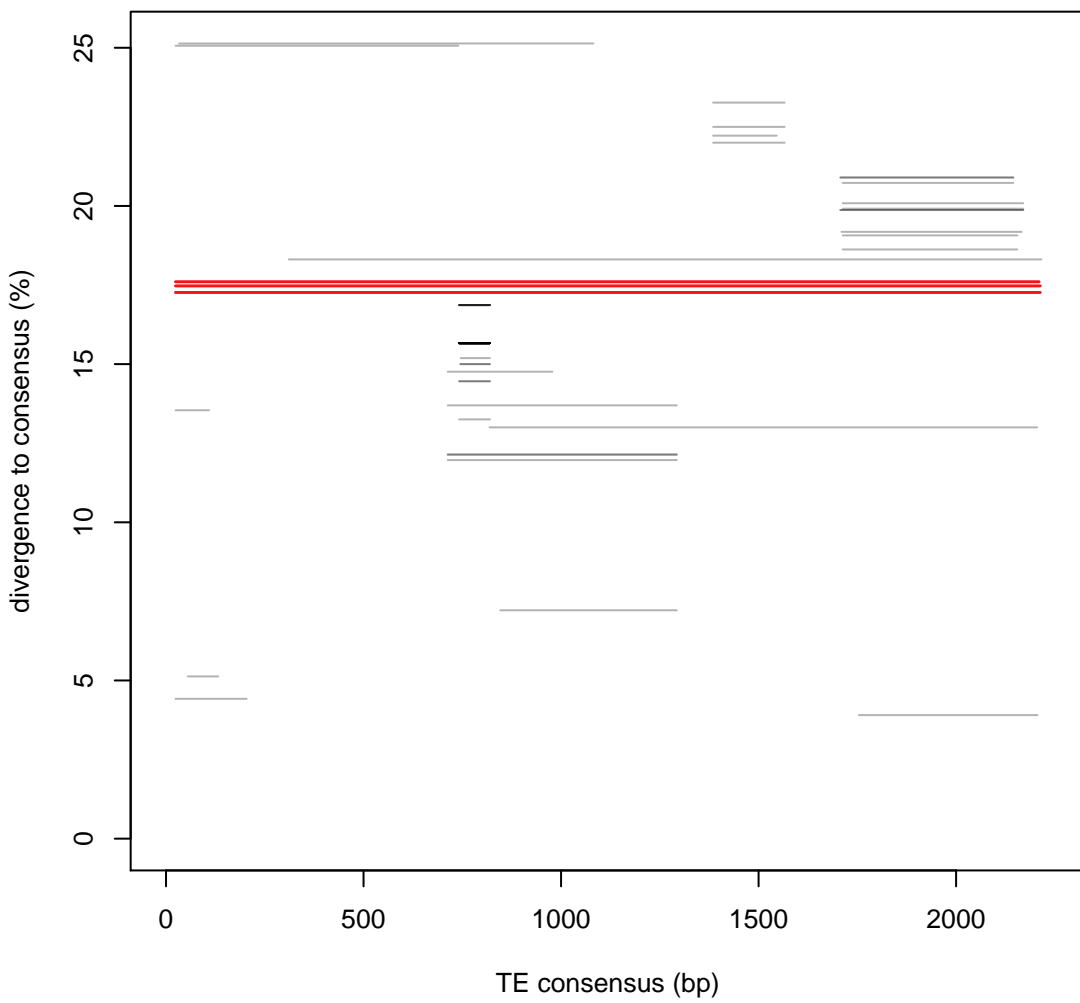
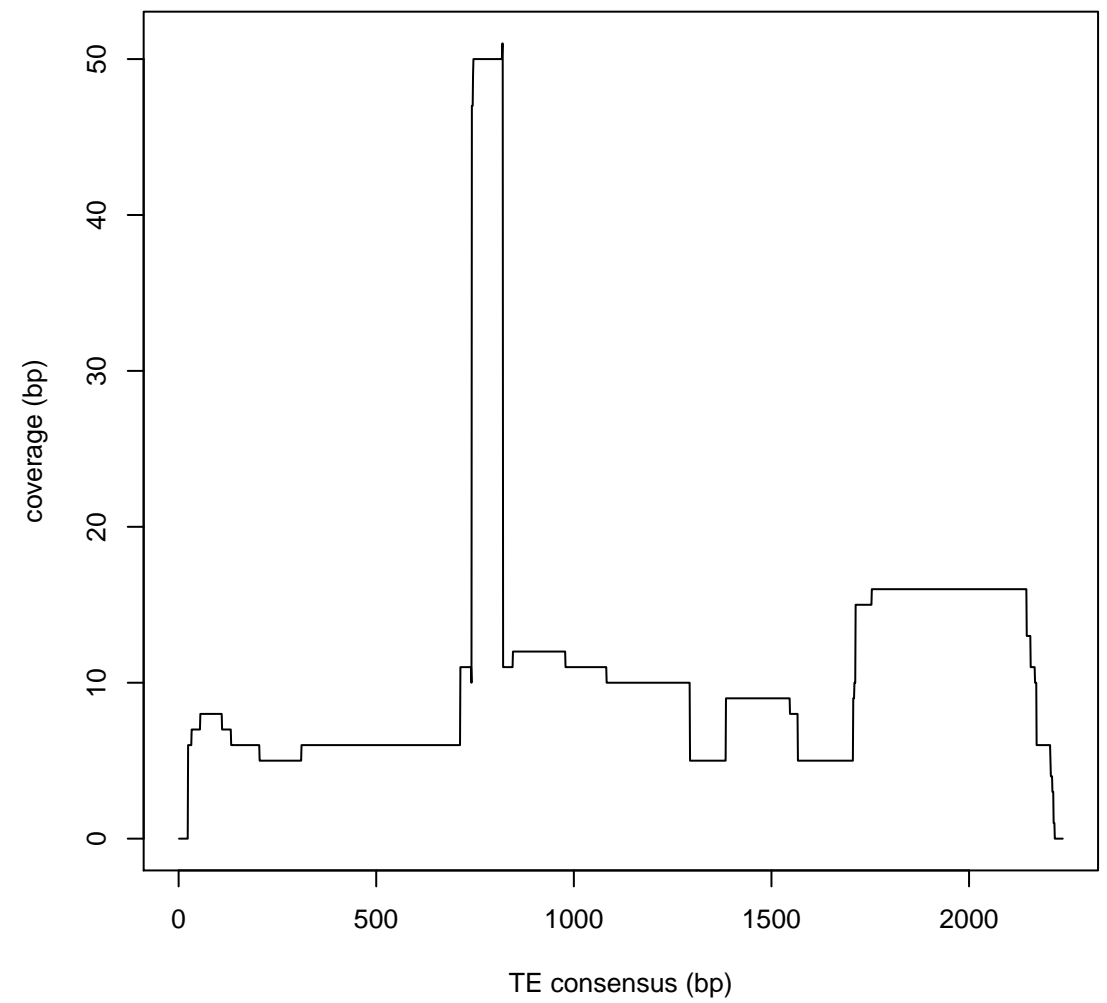


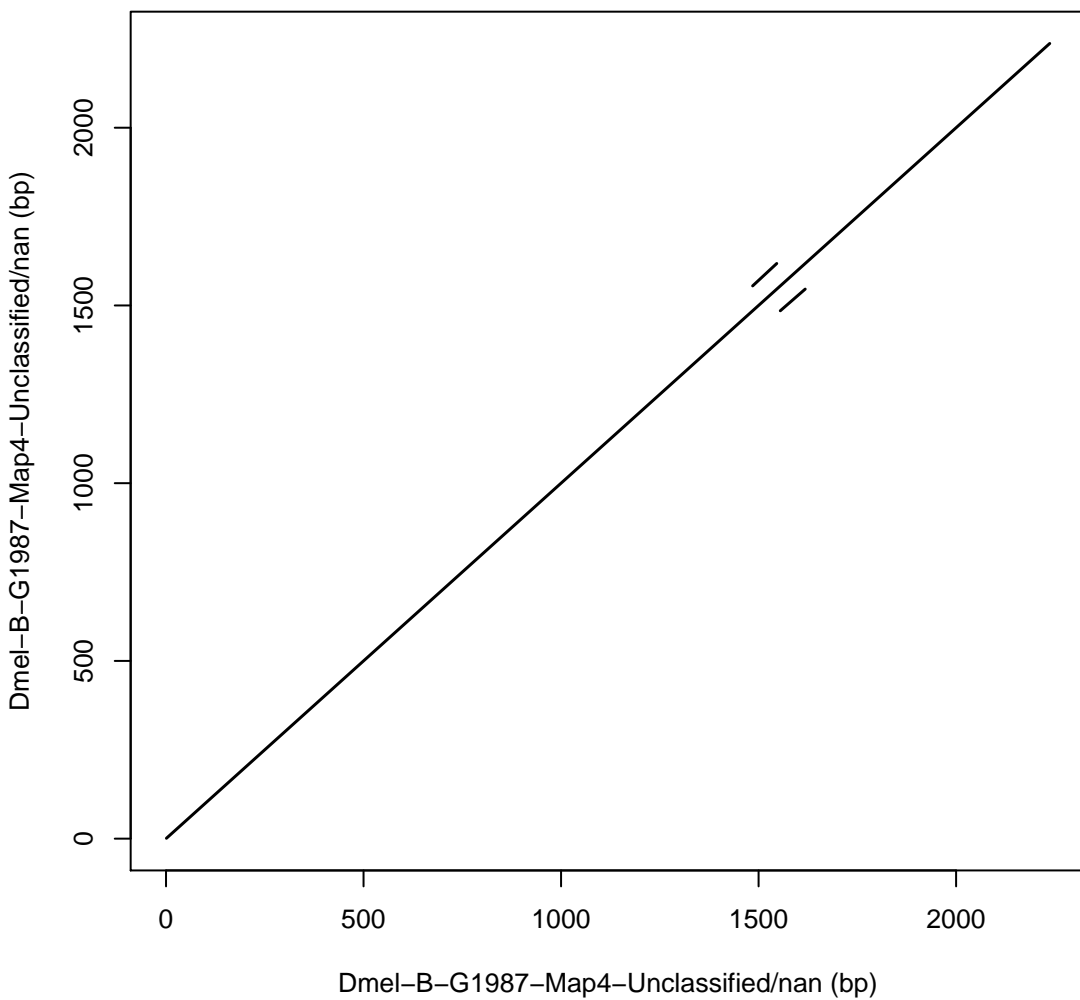
TE: Dmel-B-G1987-Map4-Unclassified/nan  
consensus size: 2237bp; fragments: 71; full length: 3 (>=2013.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

