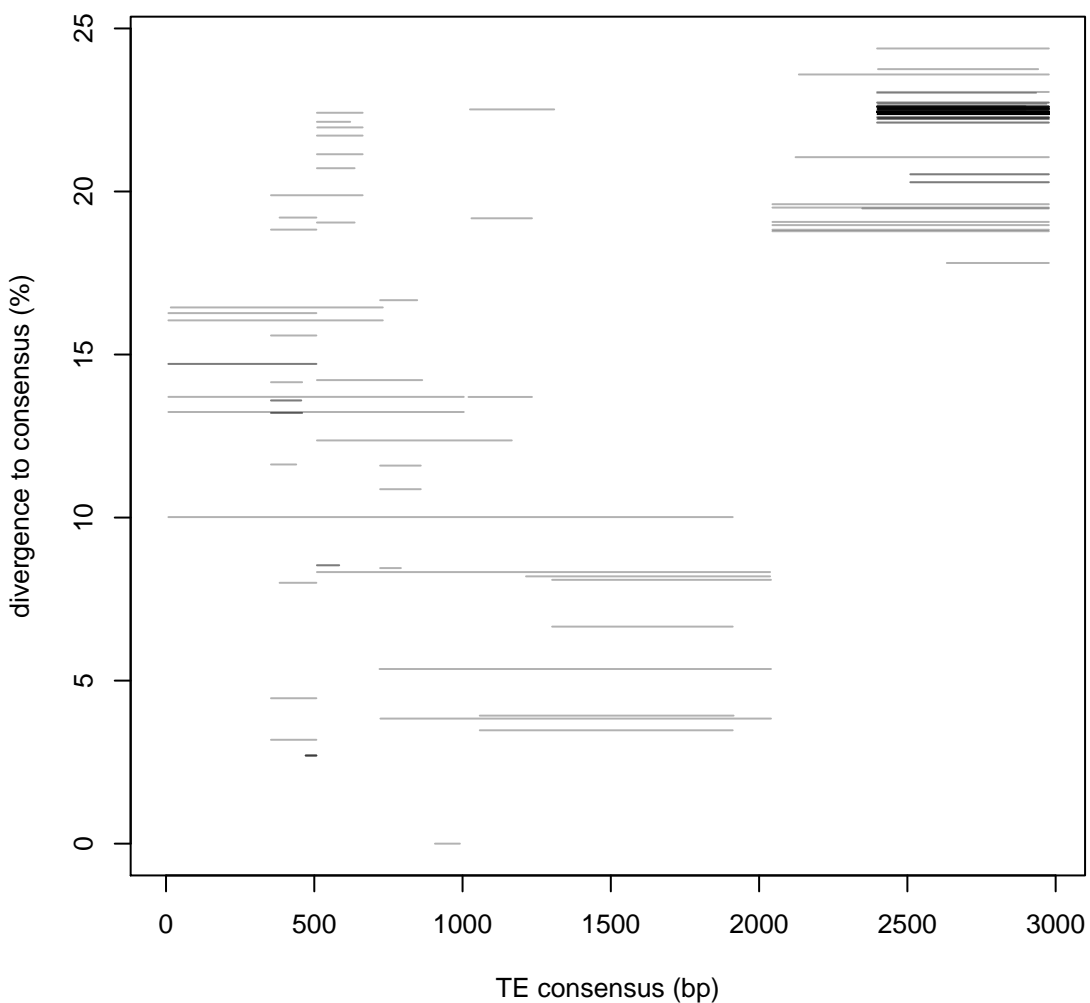
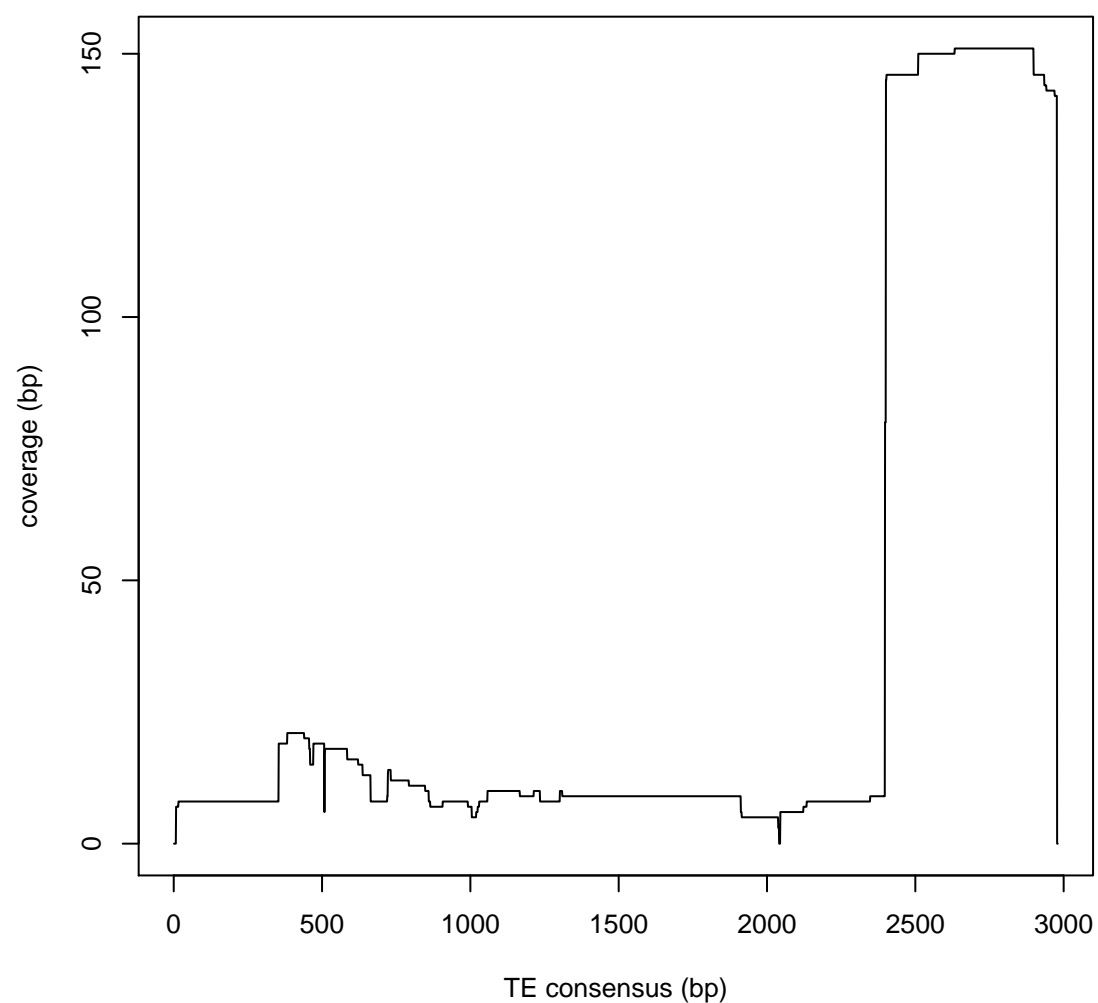


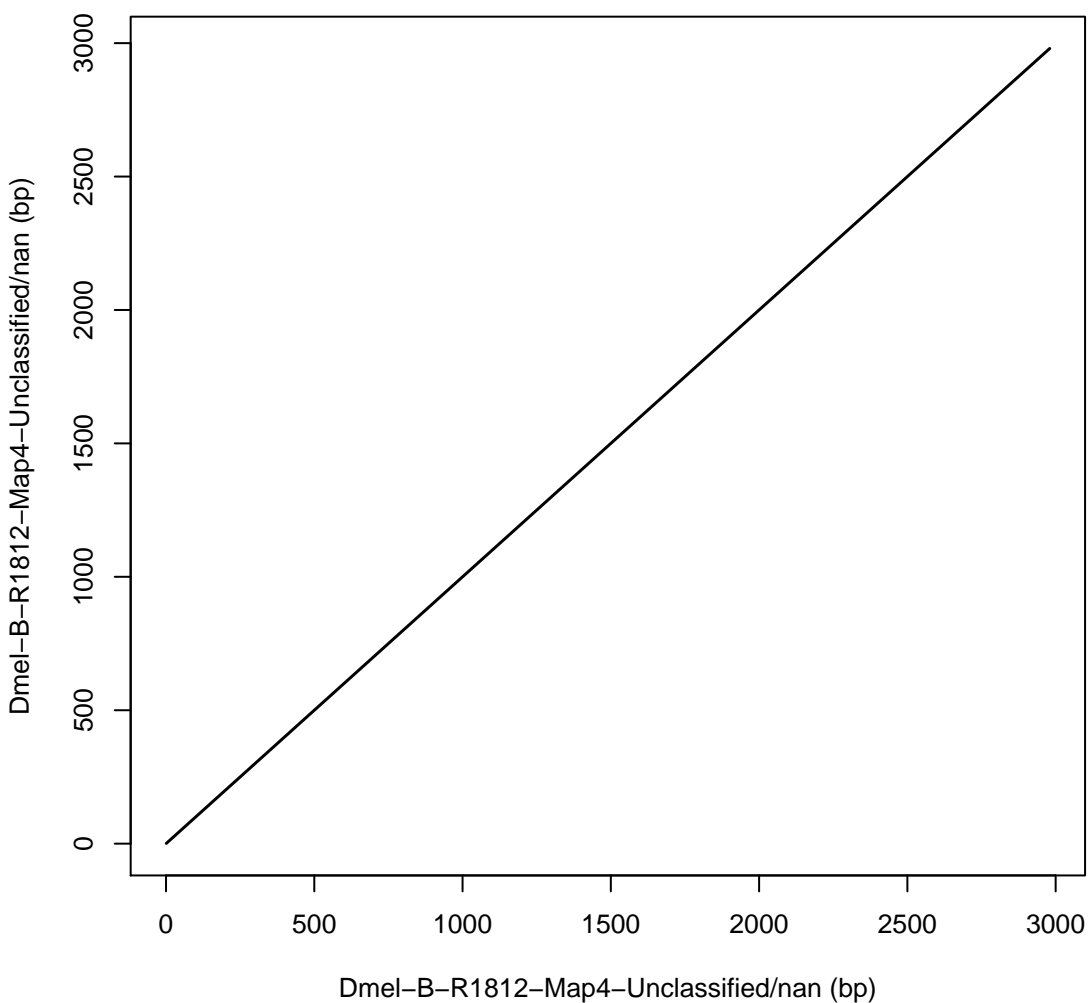
TE: Dmel-B-R1812-Map4-Unclassified/nan  
consensus size: 2980bp; fragments: 203; full length: 0 (>=2682bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

