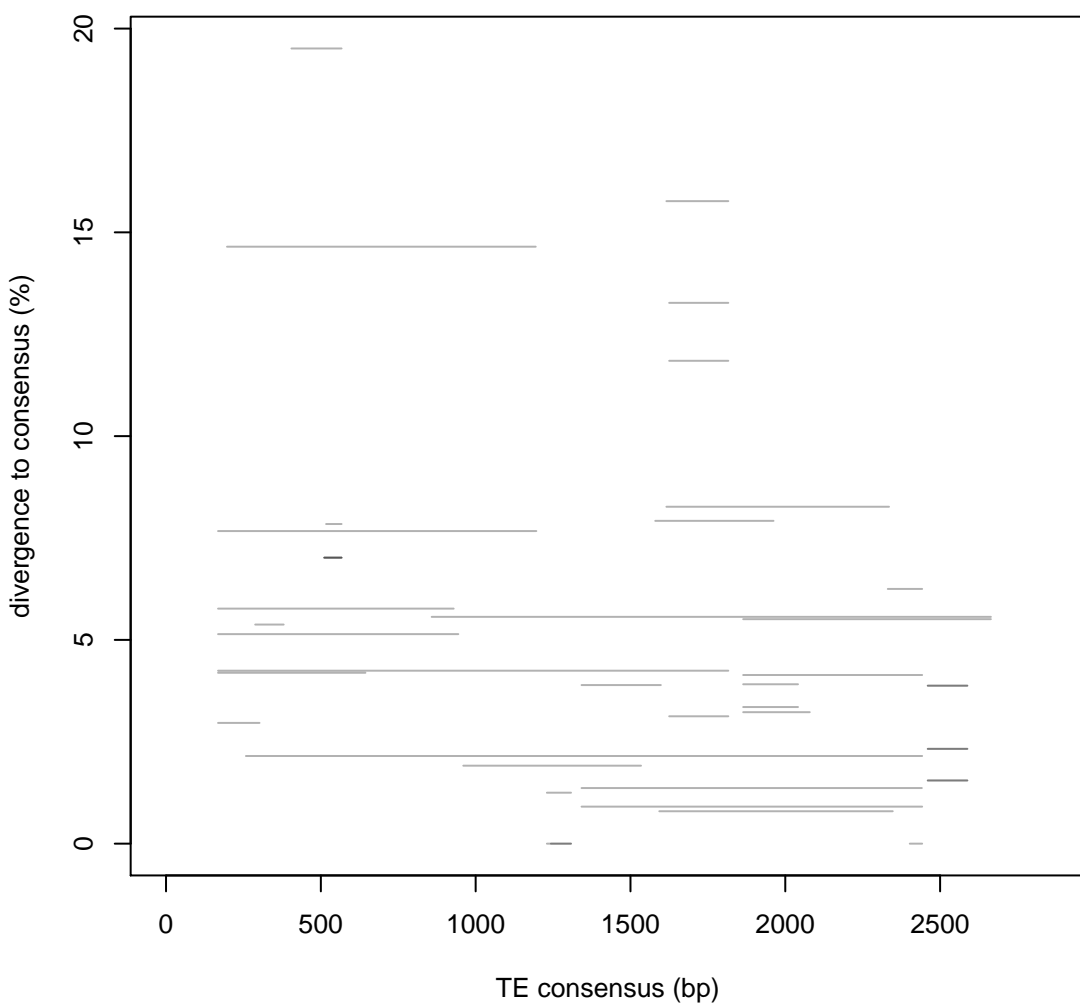
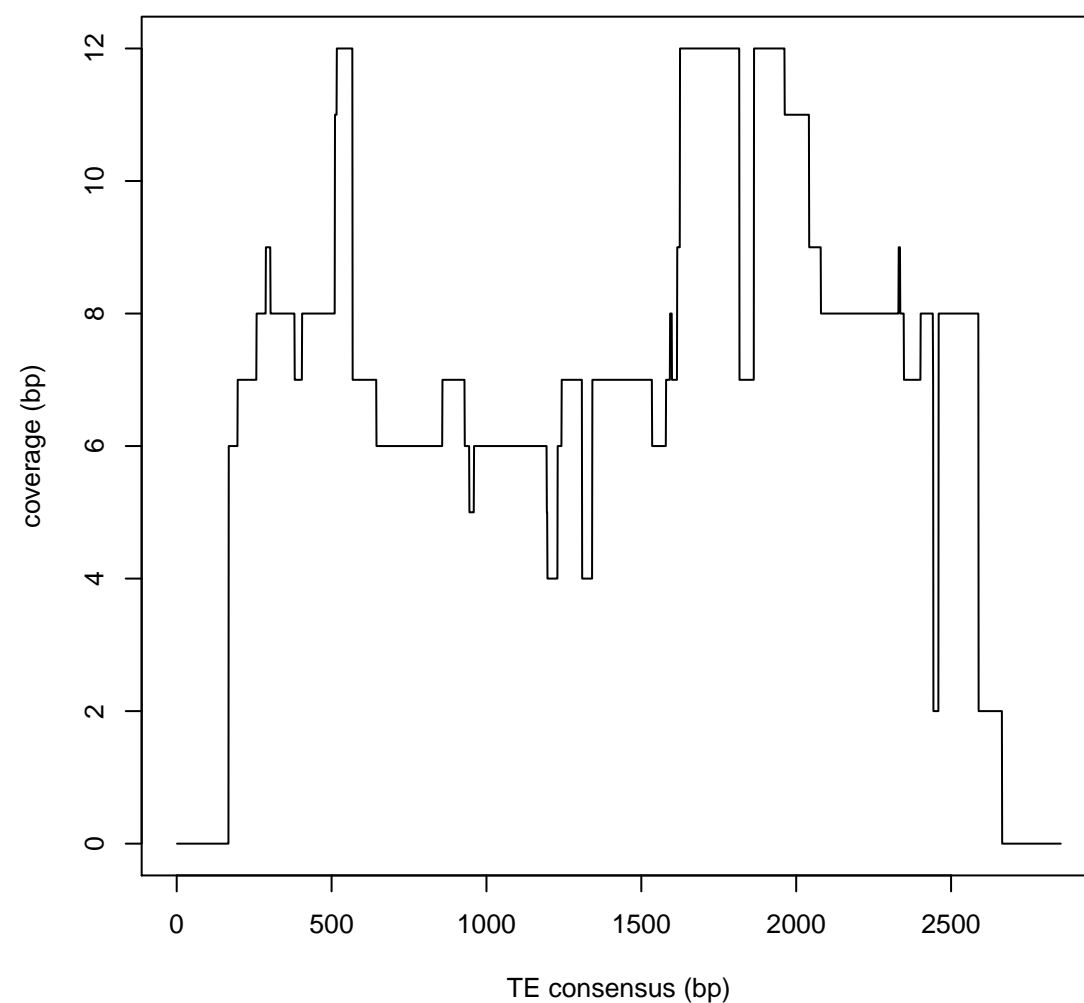


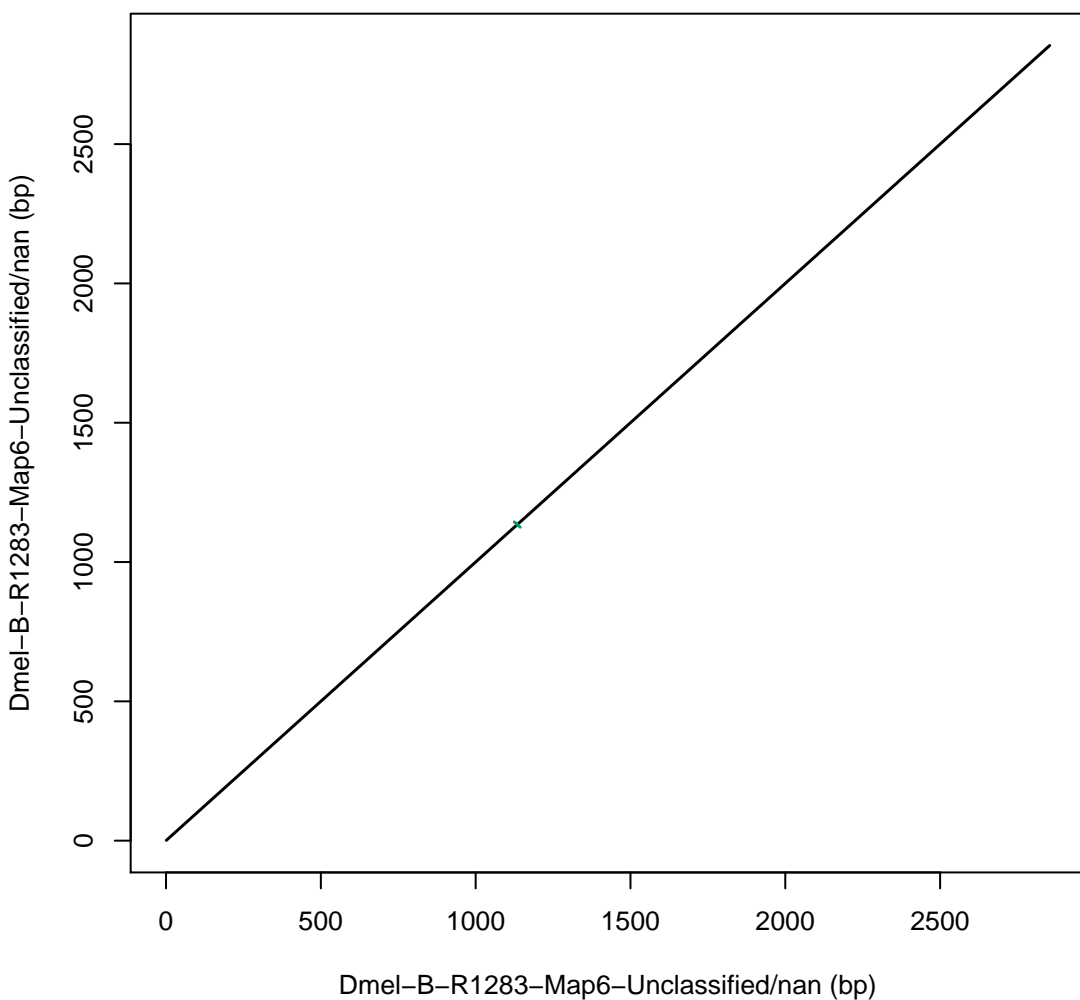
TE: Dmel-B-R1283-Map6-Unclassified/nan  
consensus size: 2854bp; fragments: 42; full length: 0 (>=2568.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

