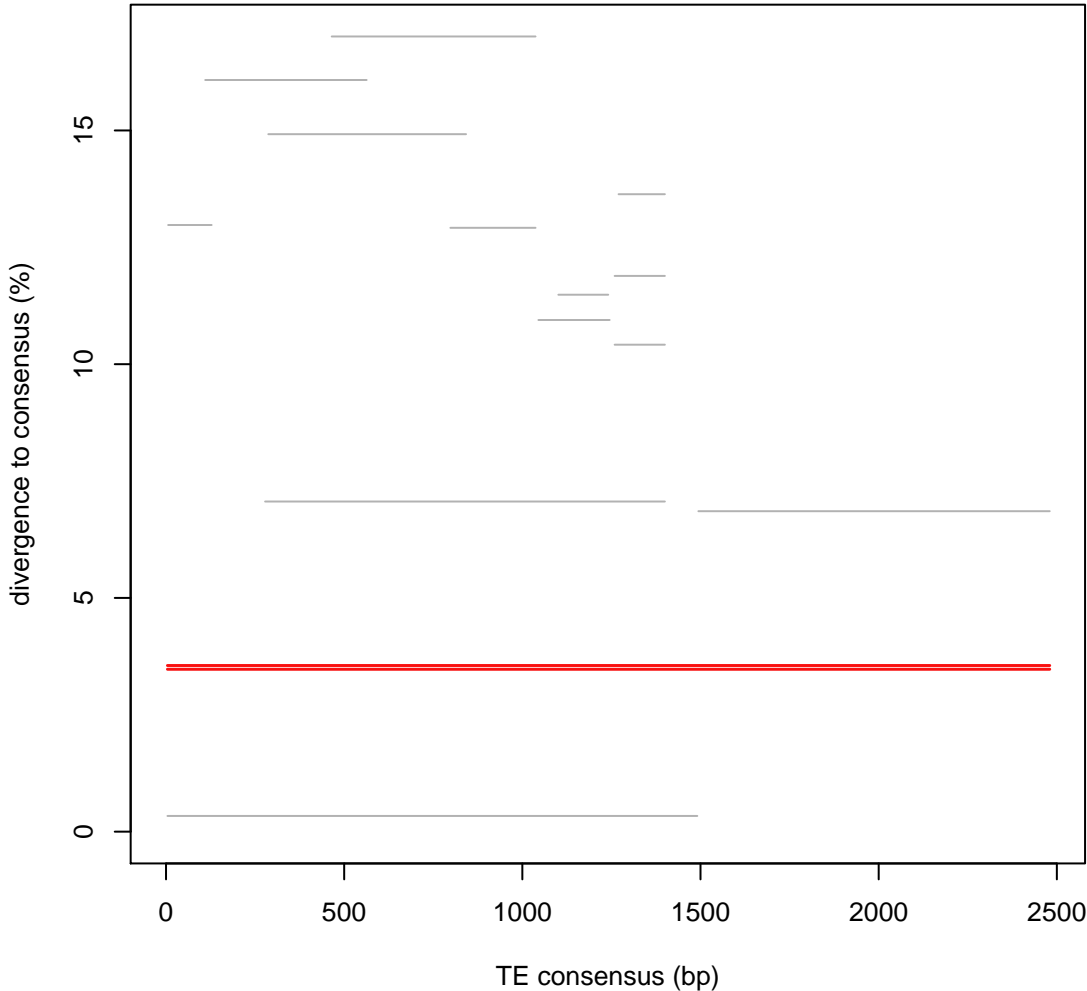
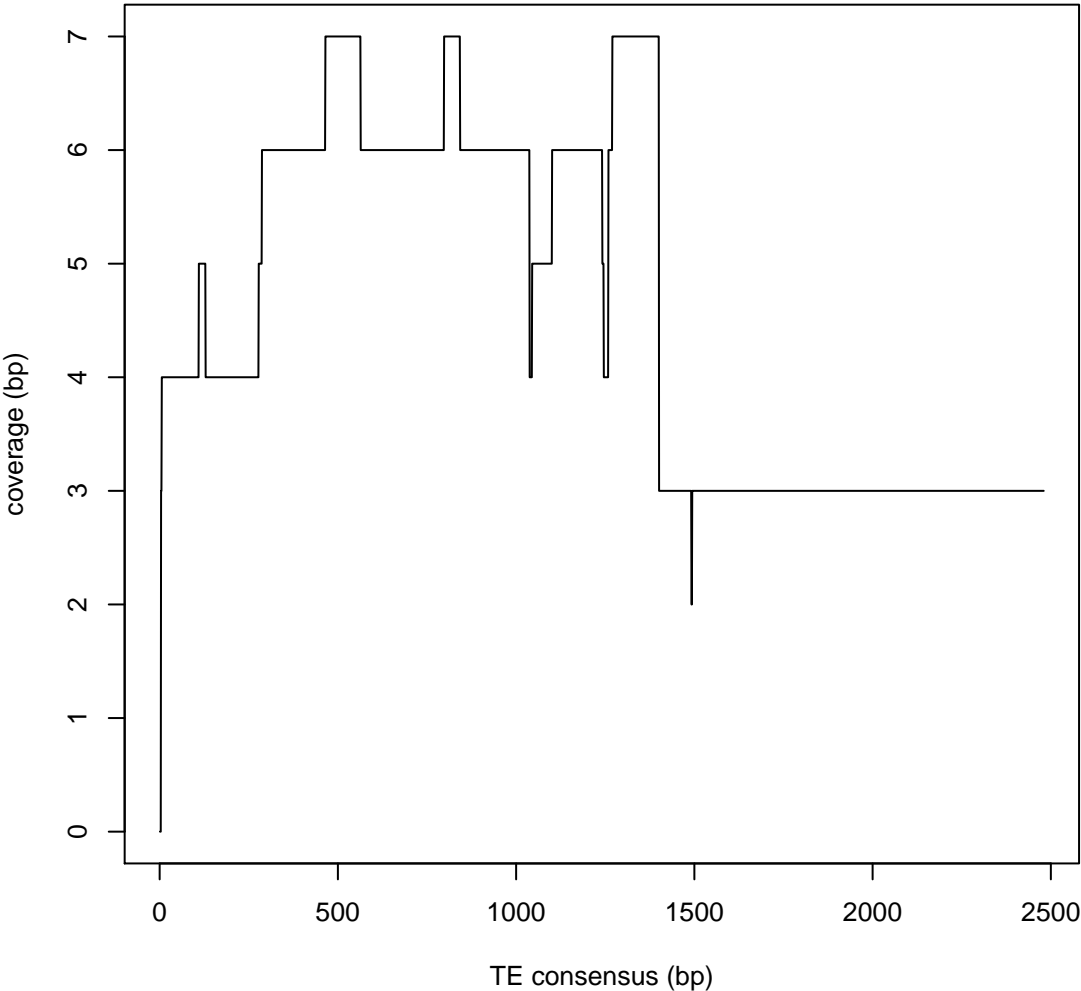


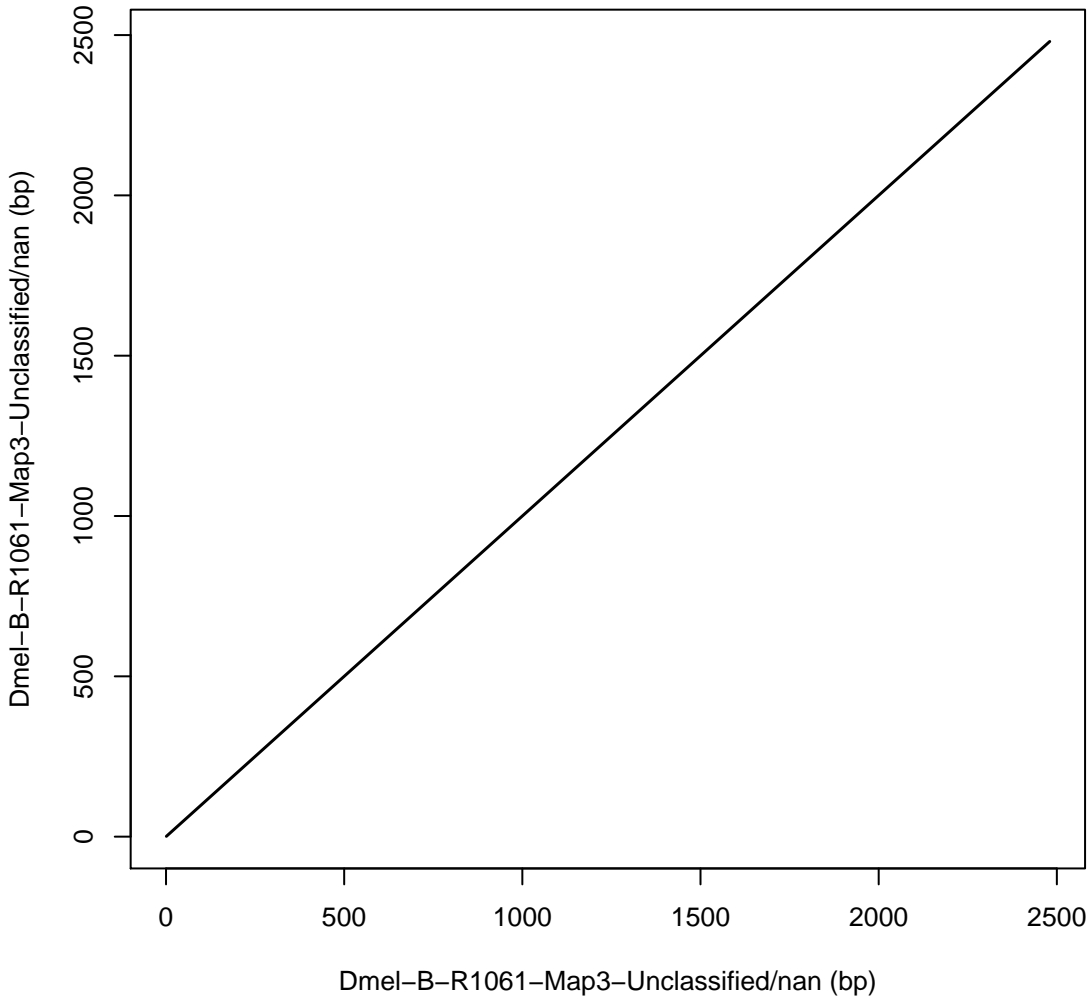
TE: Dmel-B-R1061-Map3-Unclassified/nan
consensus size: 2480bp; fragments: 15; full length: 2 (>=2232bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

