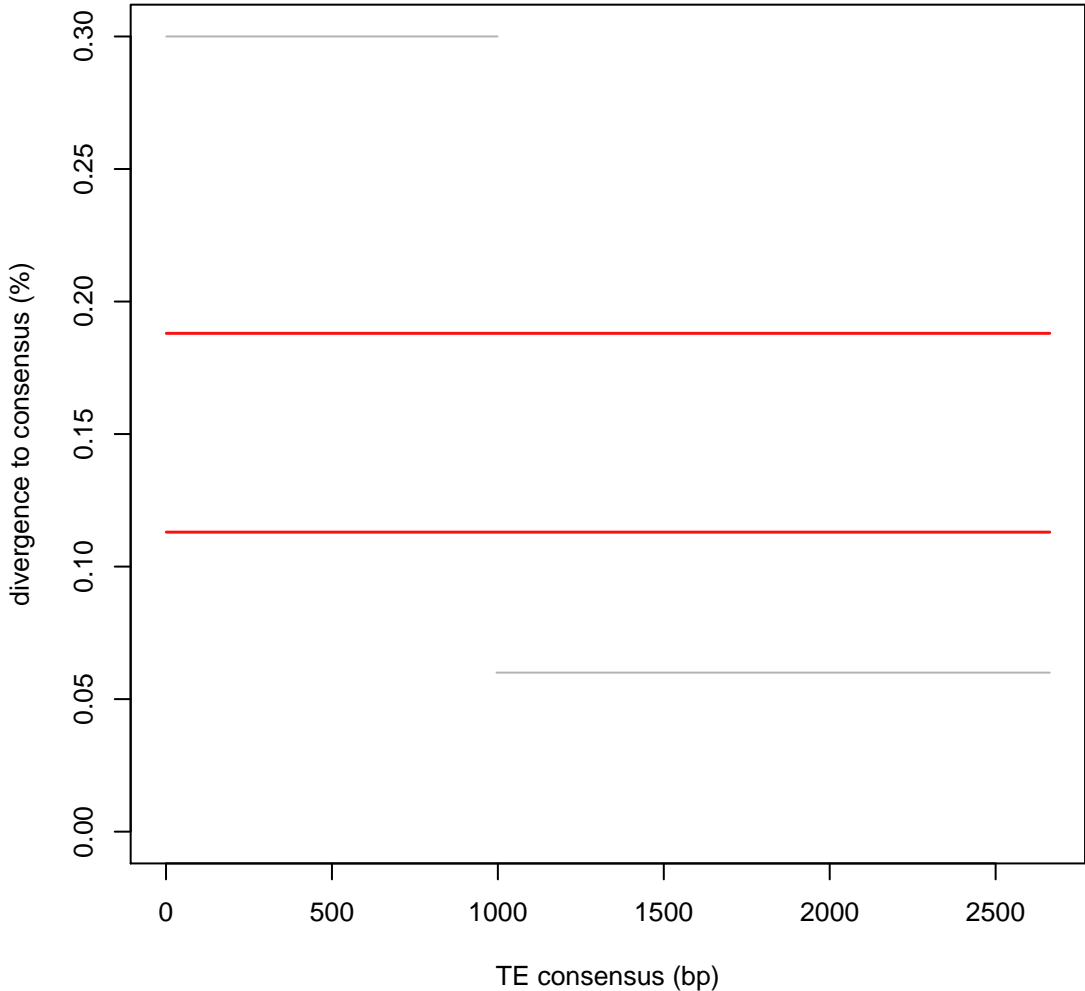
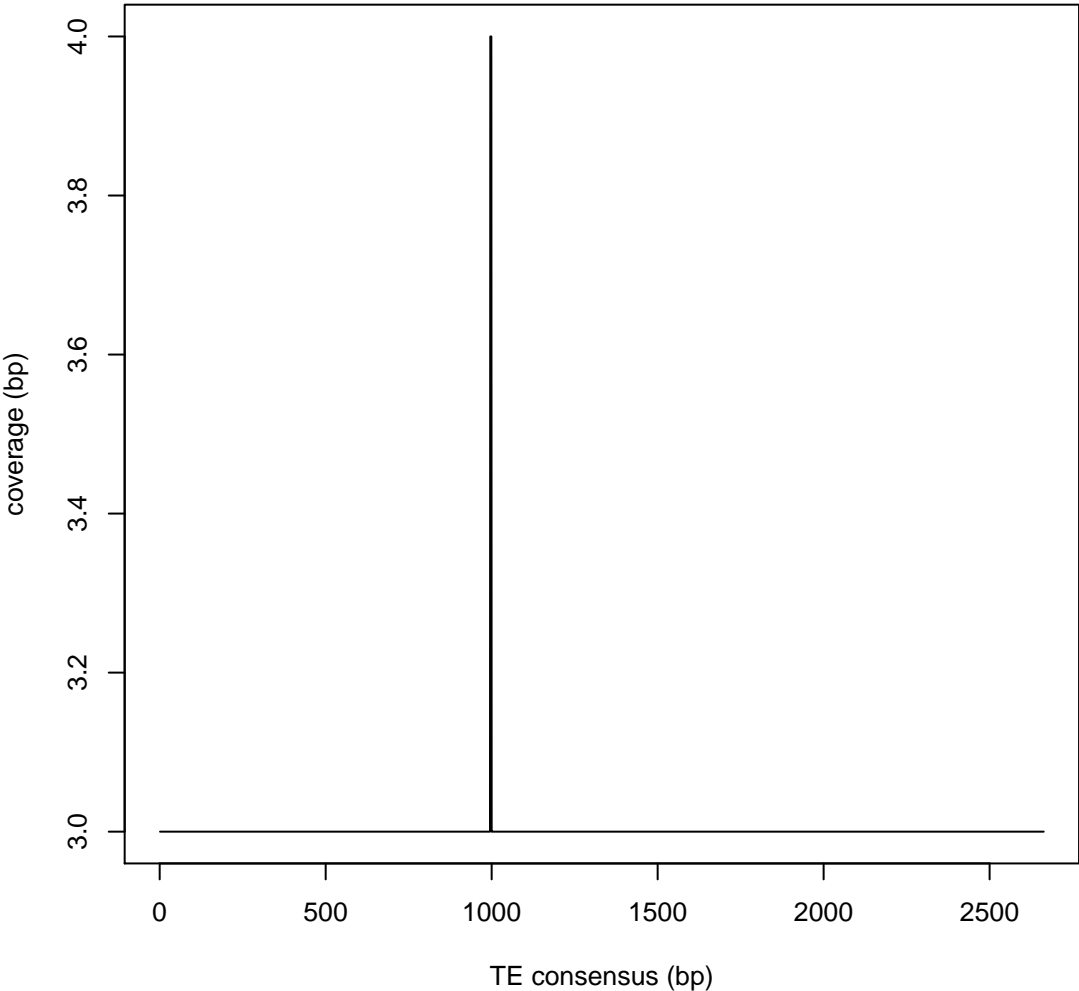


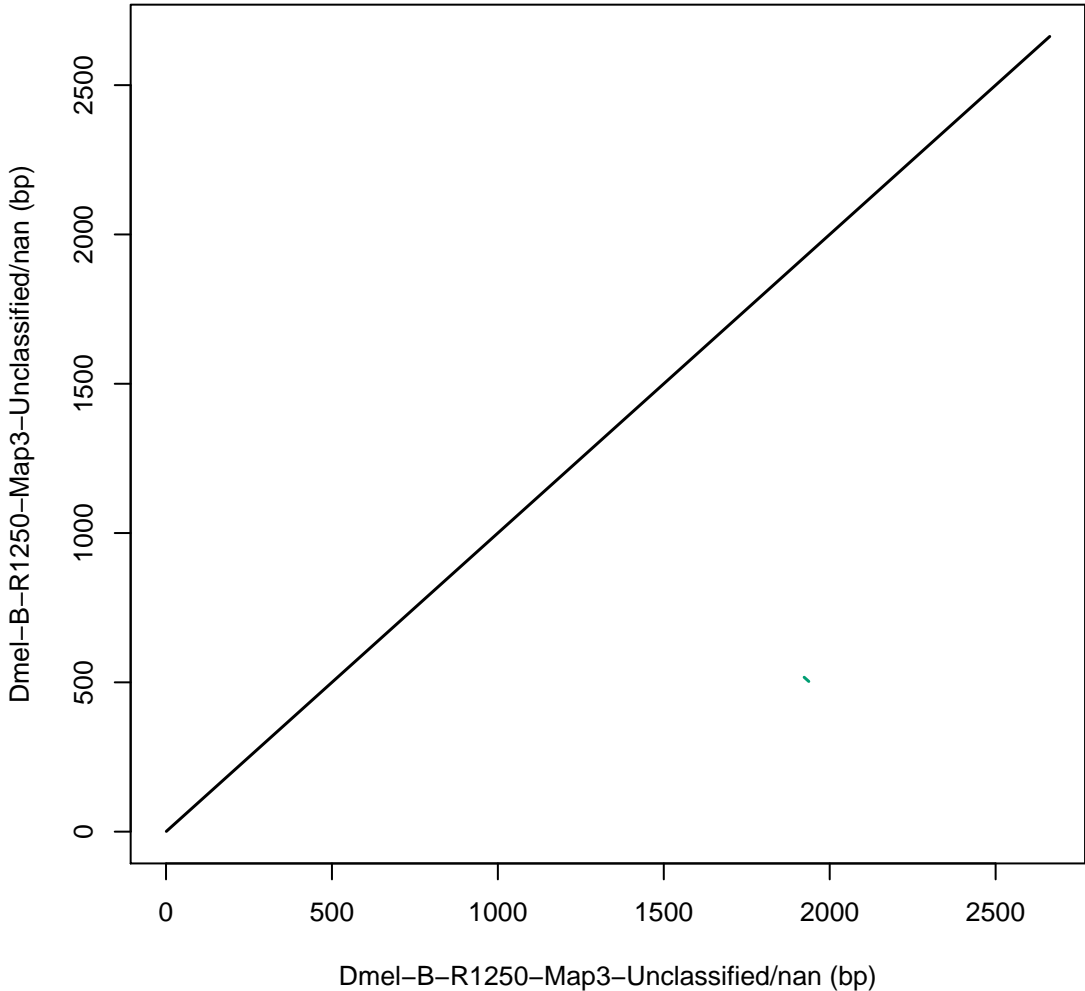
TE: Dmel-B-R1250-Map3-Unclassified/nan
consensus size: 2663bp; fragments: 4; full length: 2 (>=2396.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

