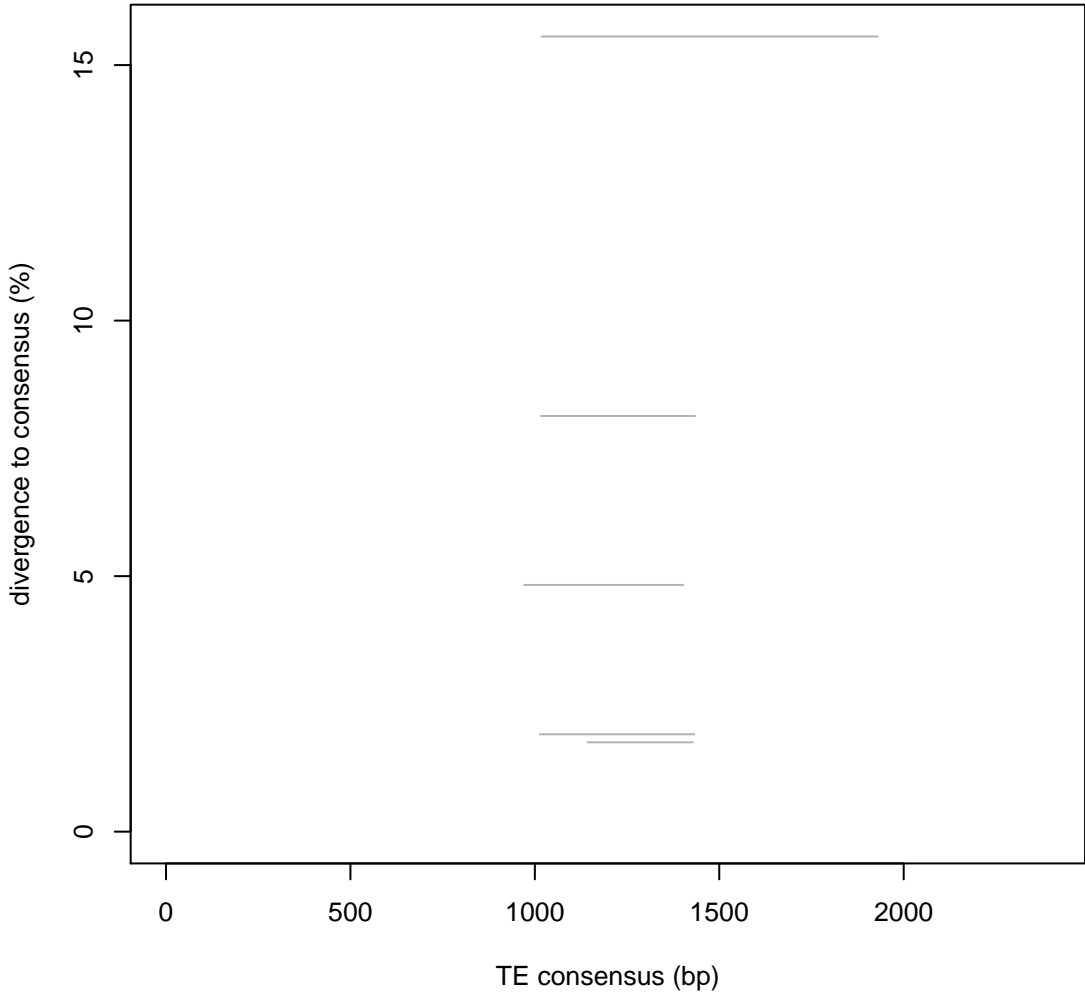
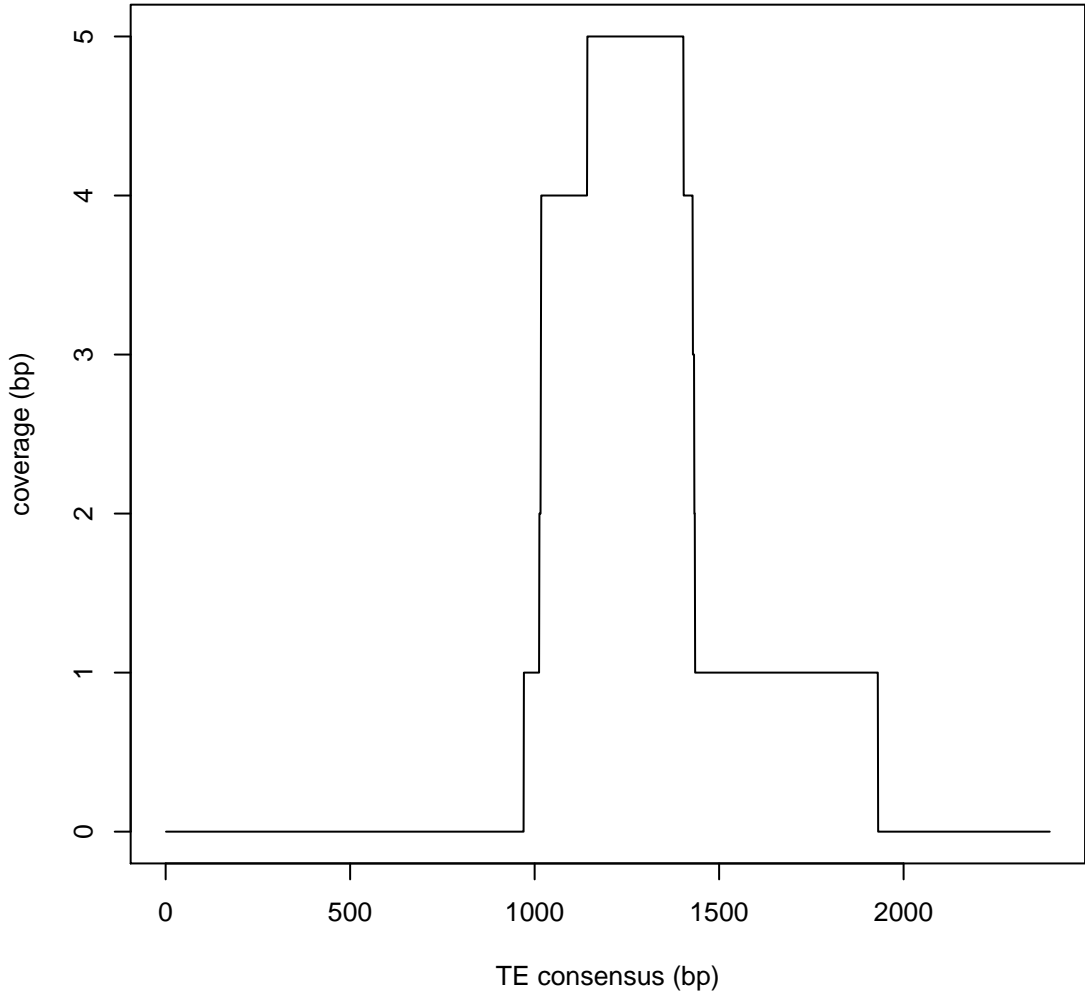


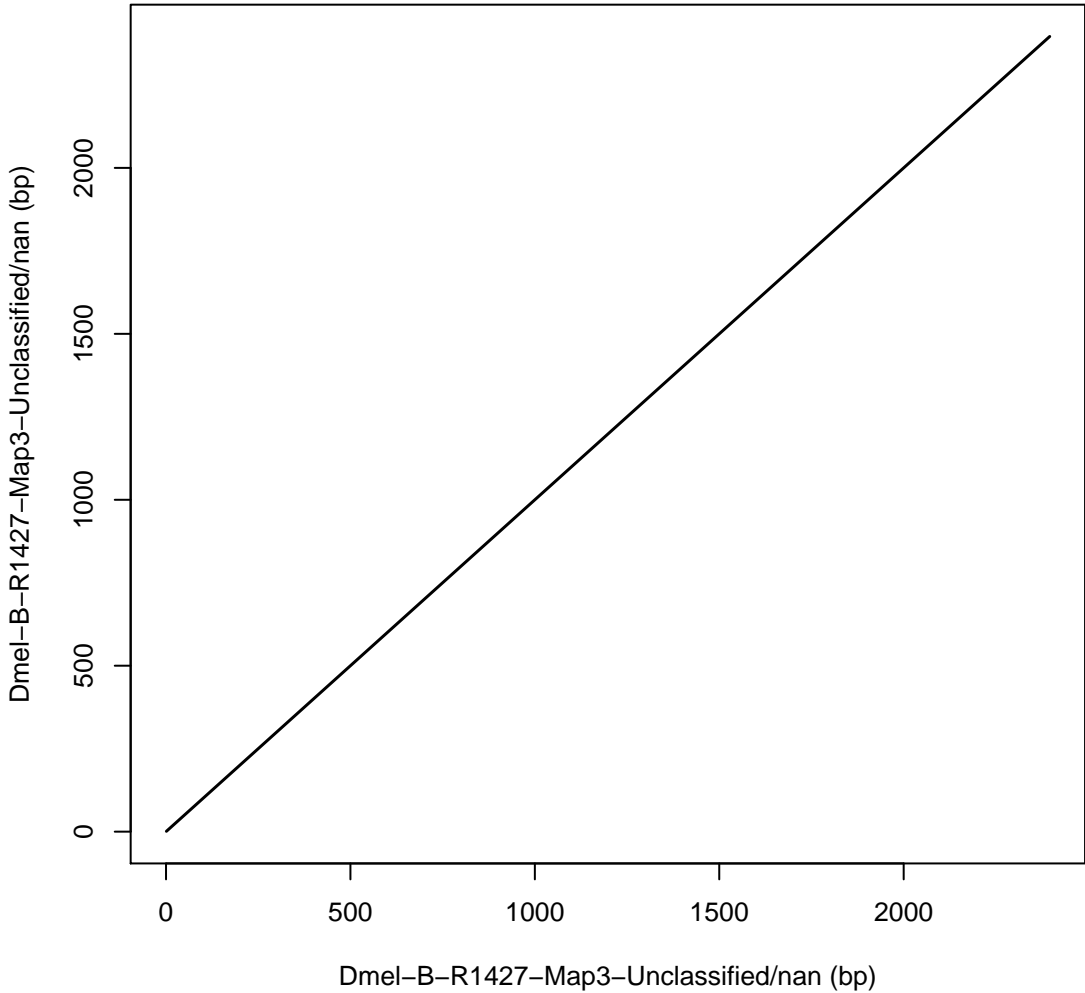
TE: Dmel-B-R1427-Map3-Unclassified/nan
consensus size: 2396bp; fragments: 5; full length: 0 (>=2156.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

