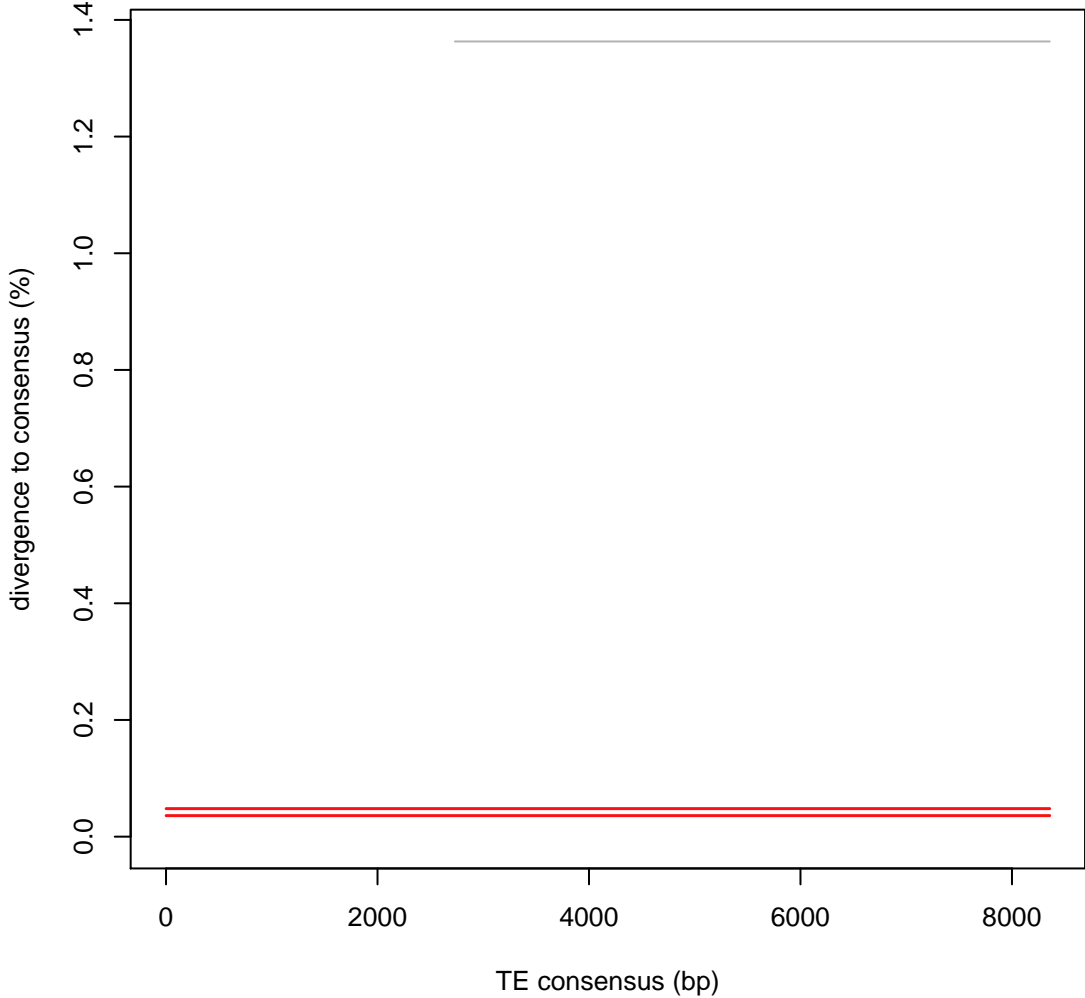
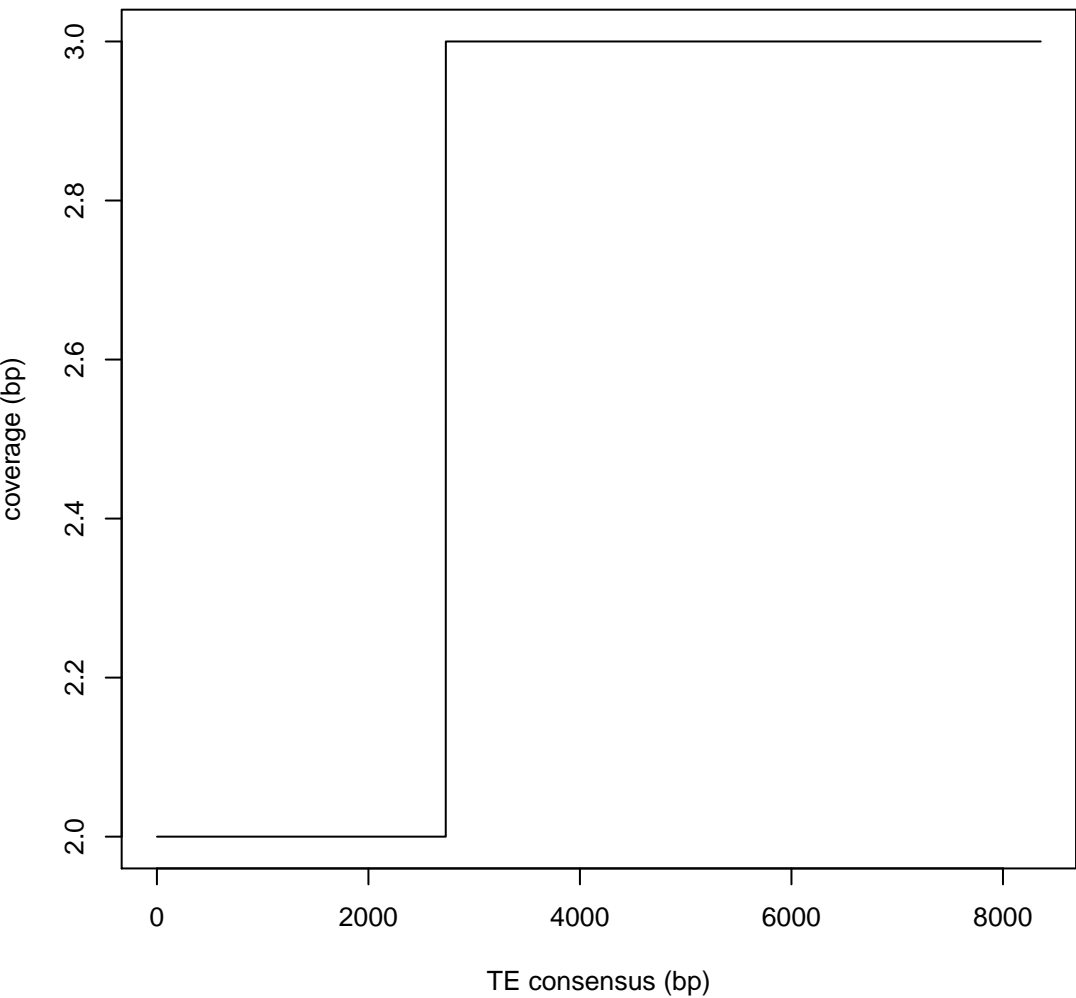


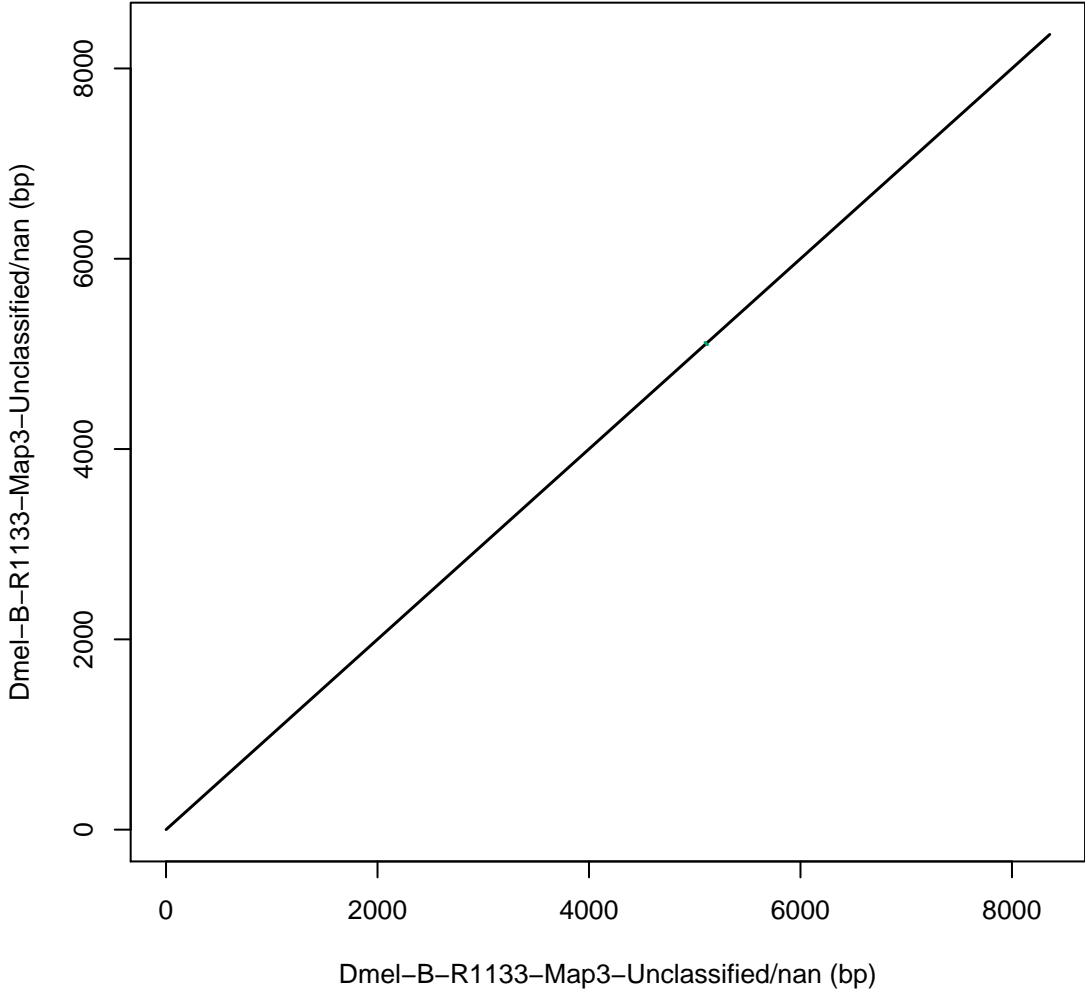
TE: Dmel-B-R1133-Map3-Unclassified/nan
consensus size: 8357bp; fragments: 3; full length: 2 (>=7521.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

