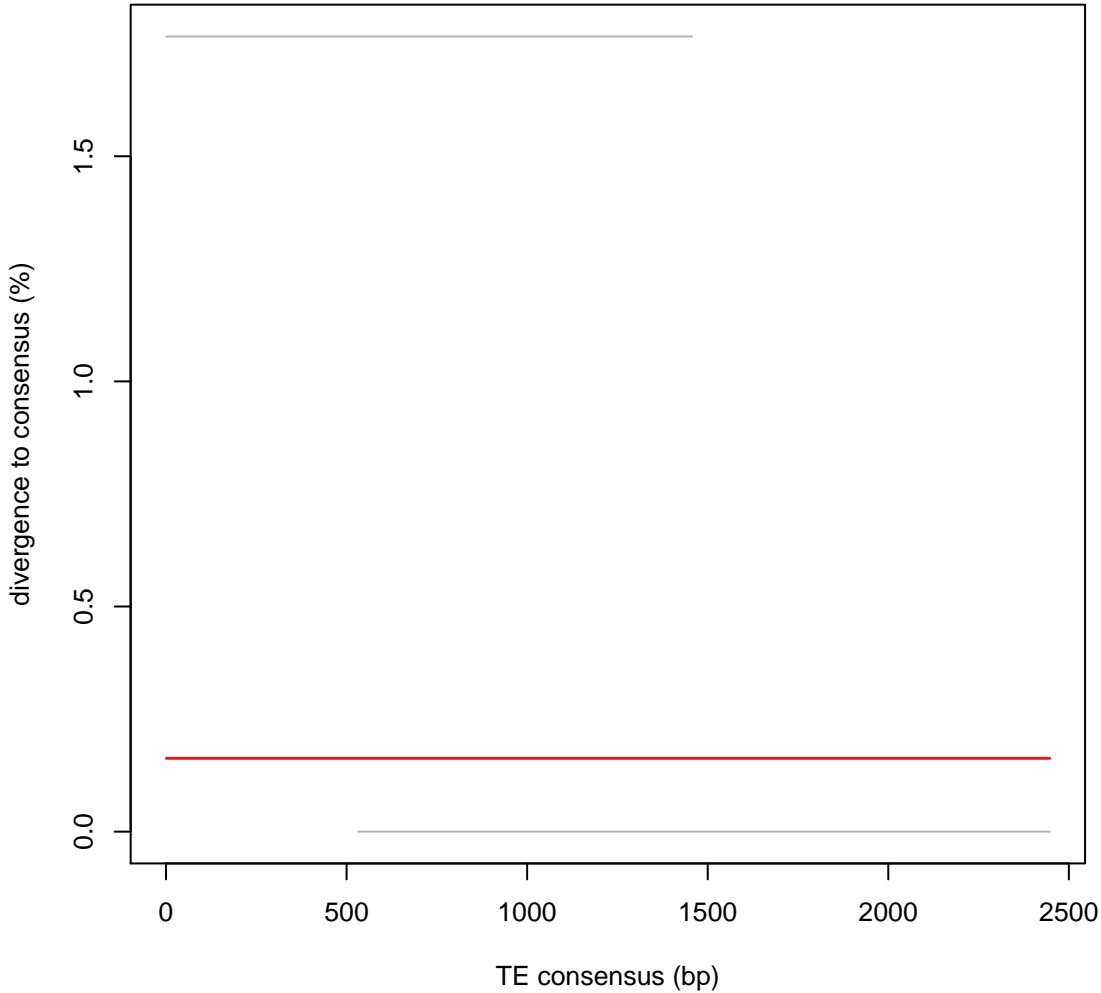
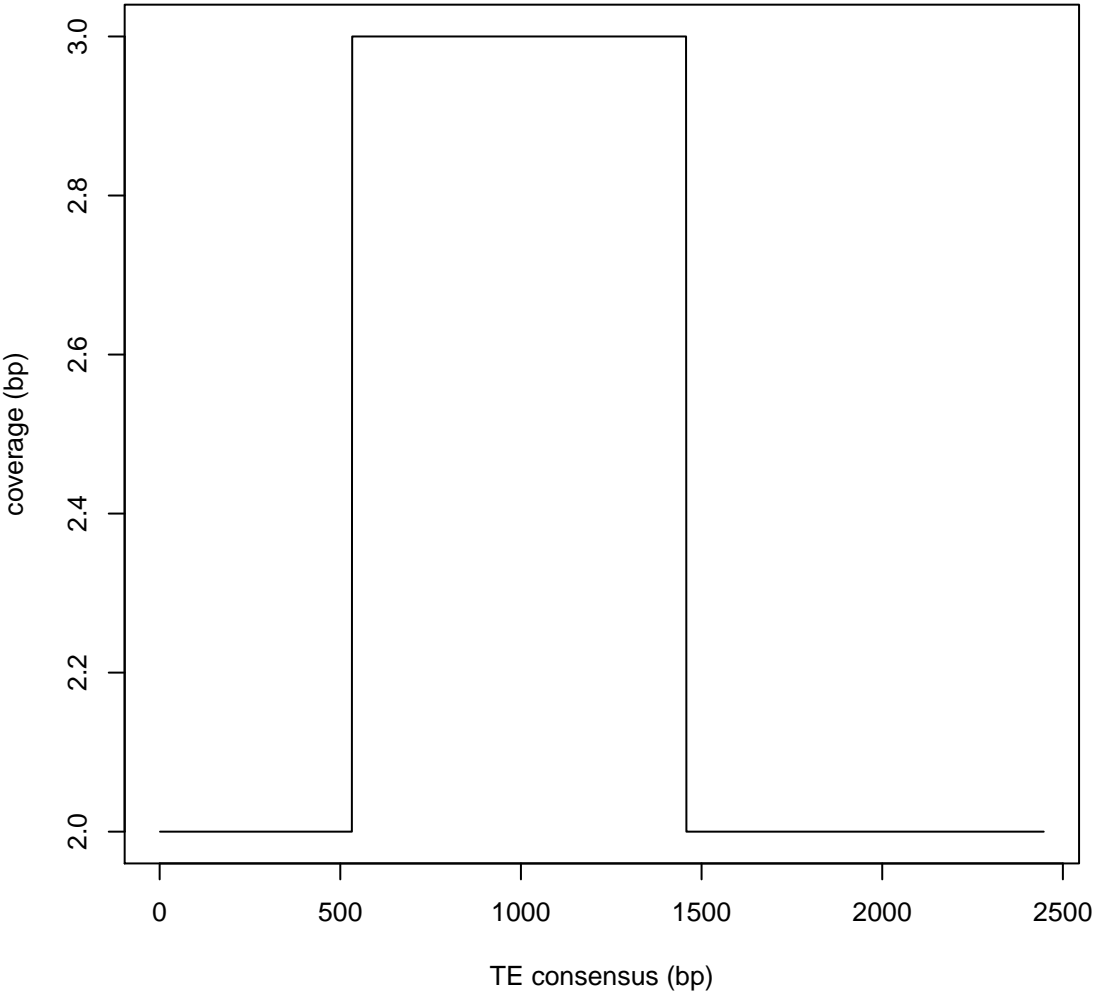


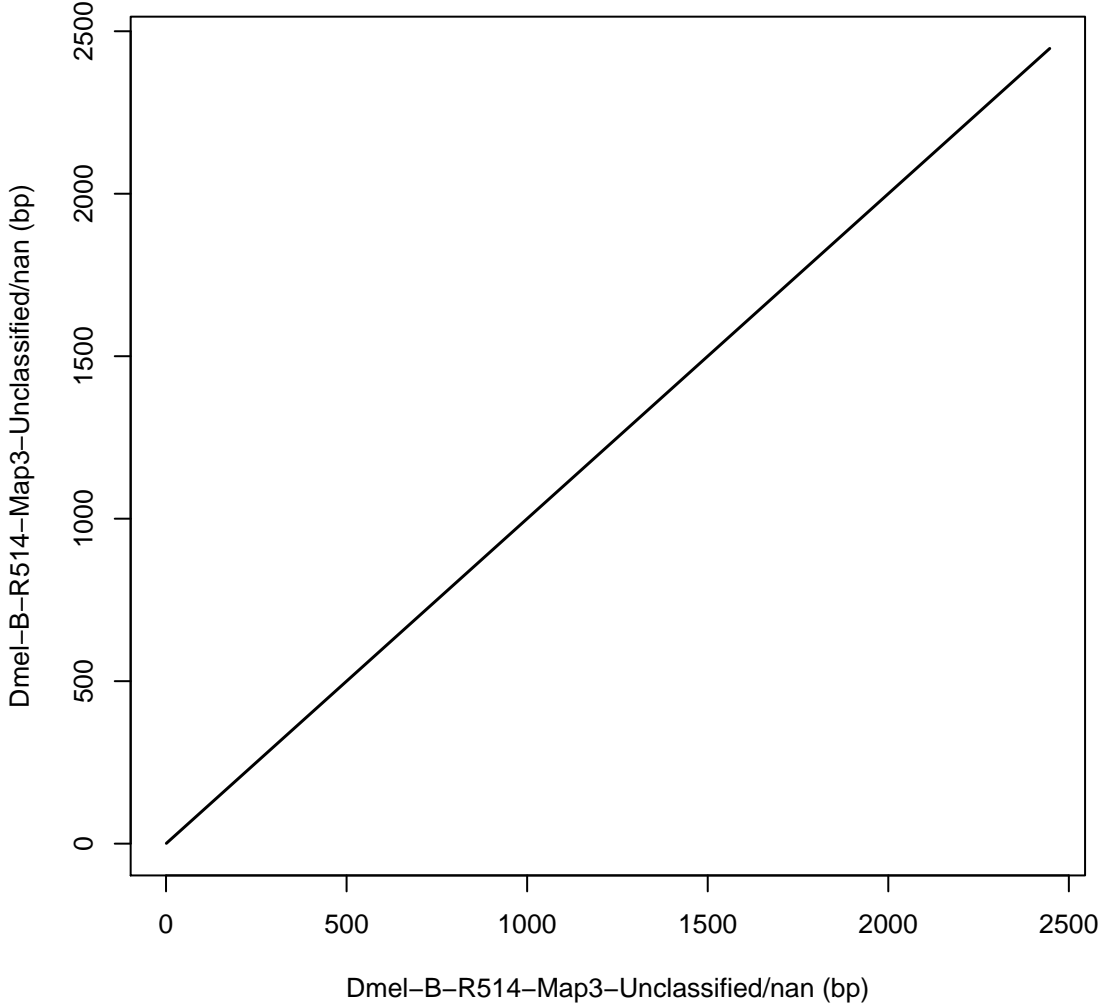
TE: Dmel-B-R514-Map3-Unclassified/nan
consensus size: 2447bp; fragments: 3; full length: 1 (>=2202.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

