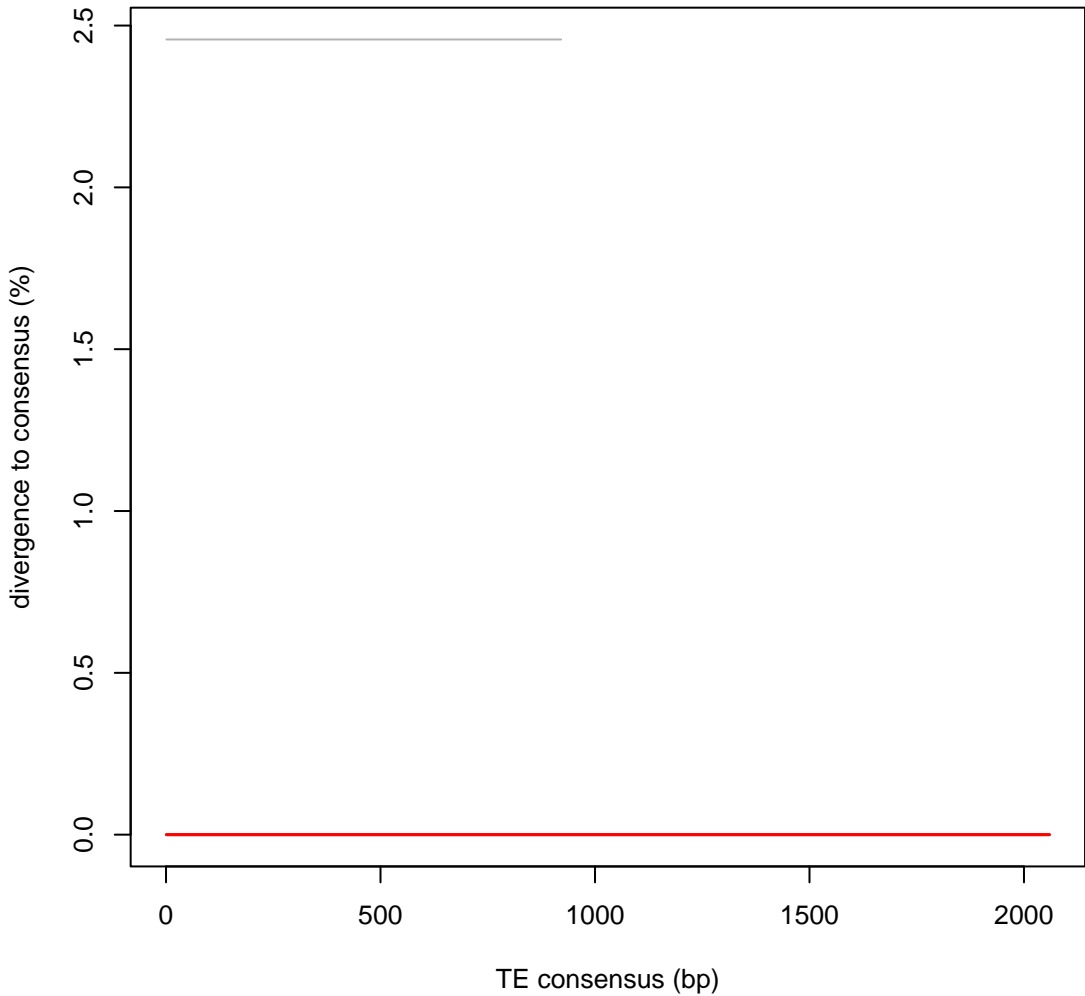
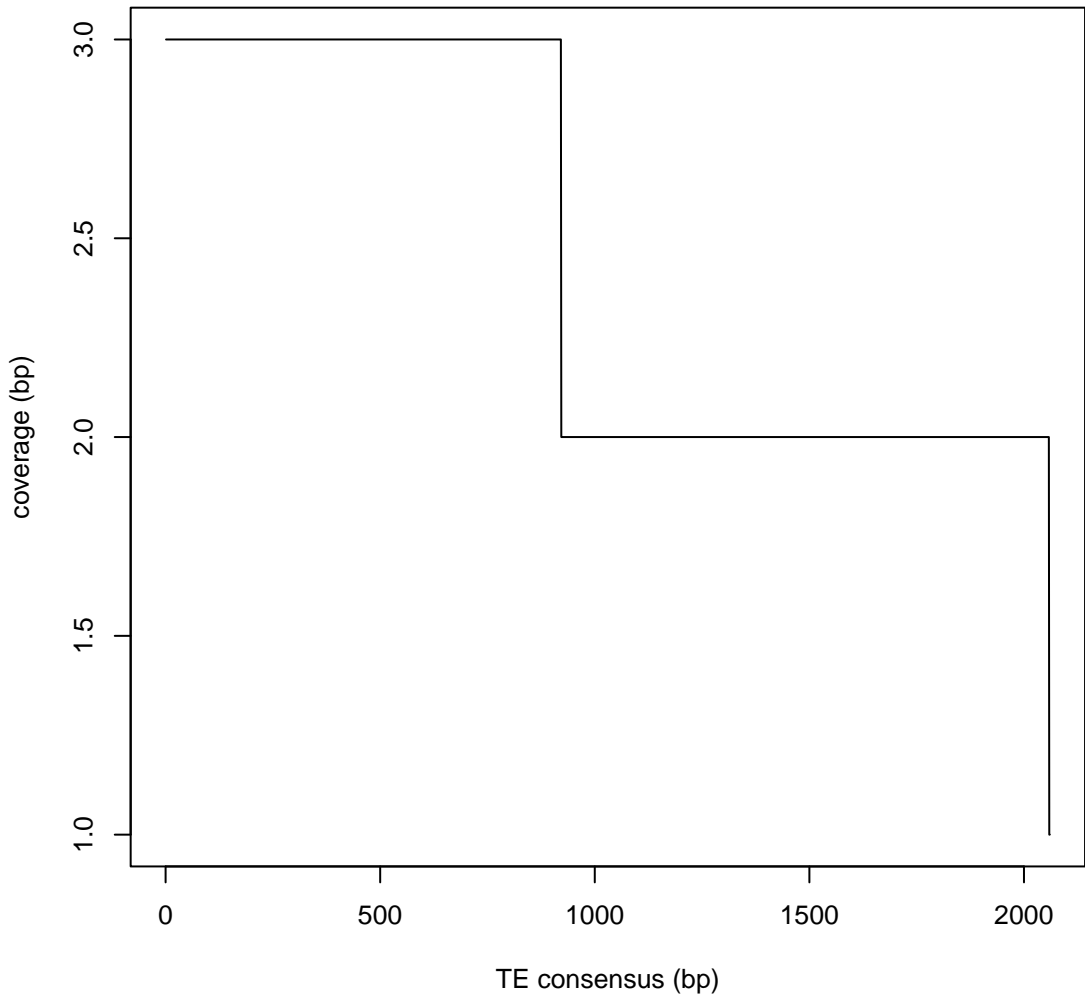


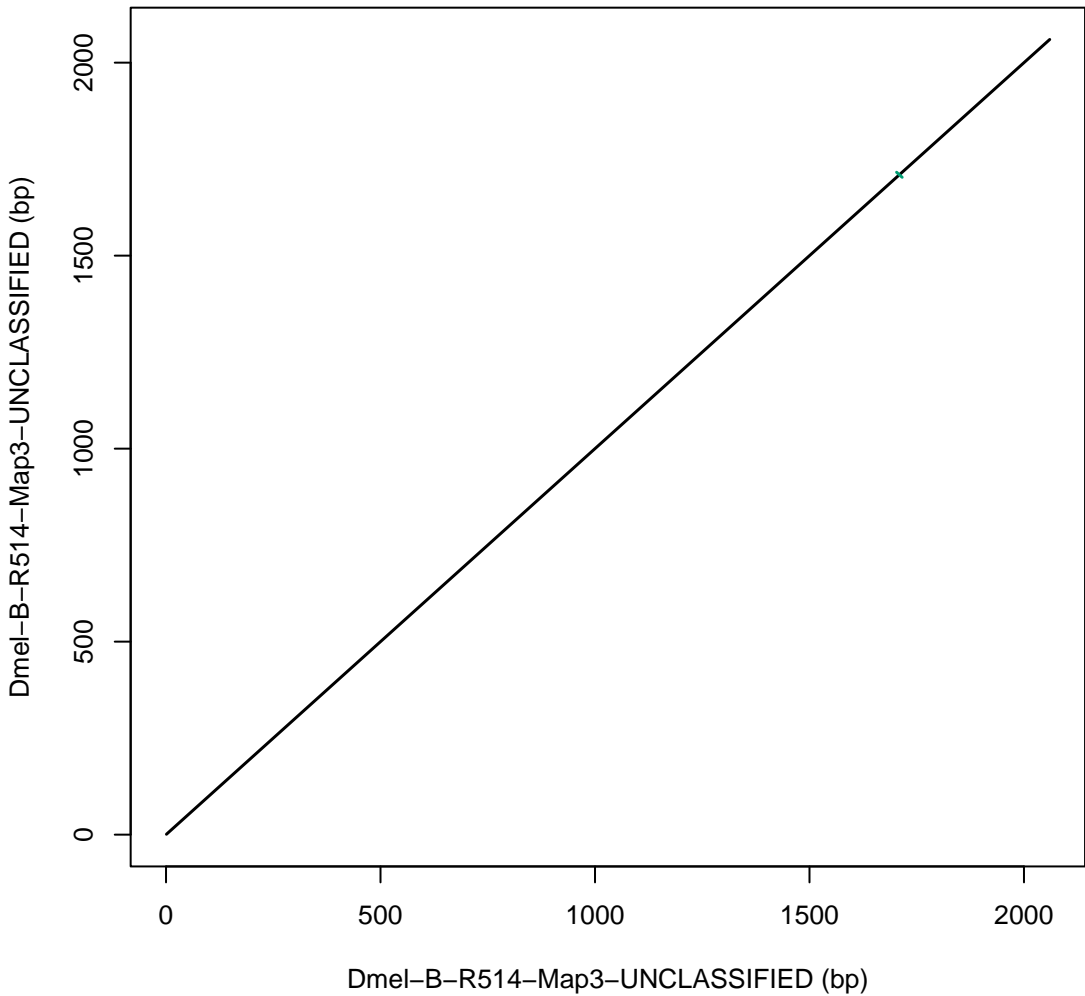
TE: Dmel-B-R514-Map3-UNCLASSIFIED  
consensus size: 2060bp; fragments: 3; full length: 2 (>=1854bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

