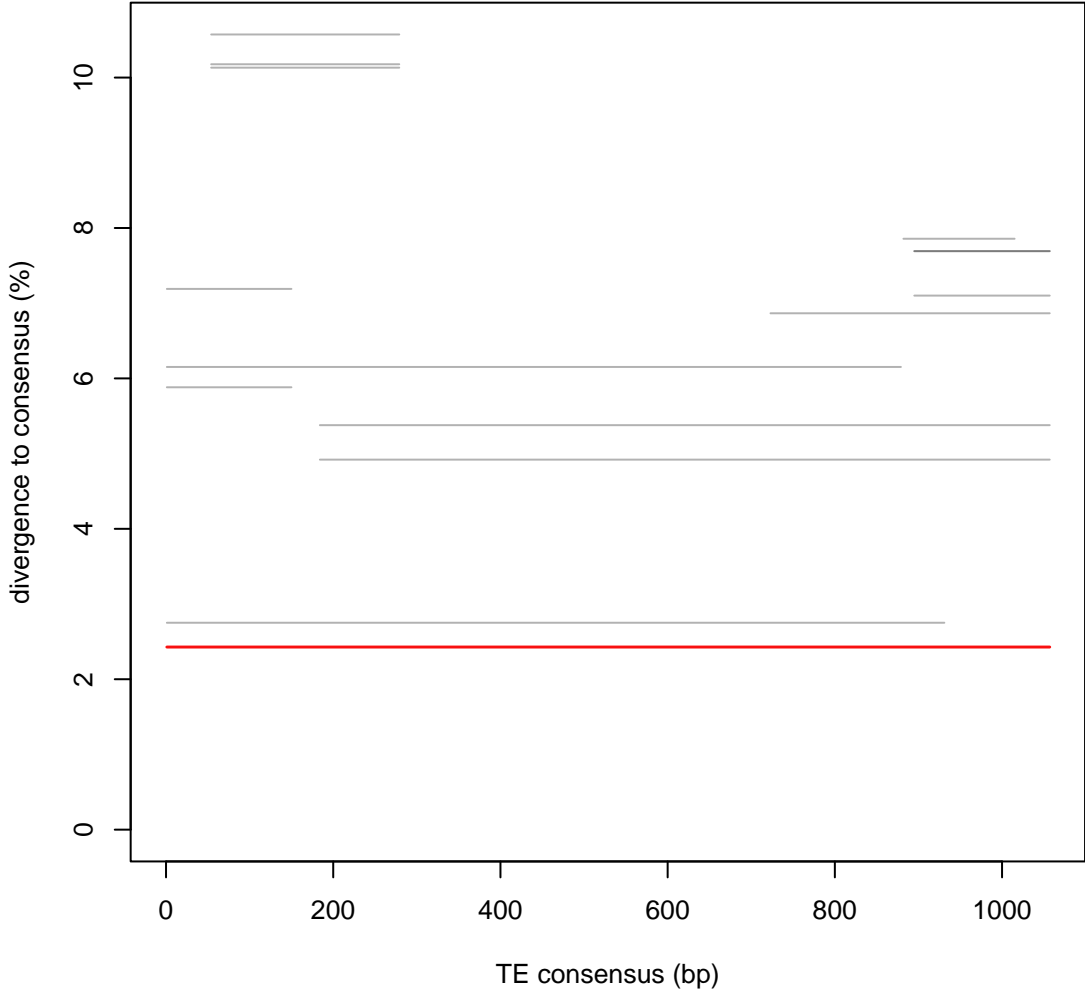
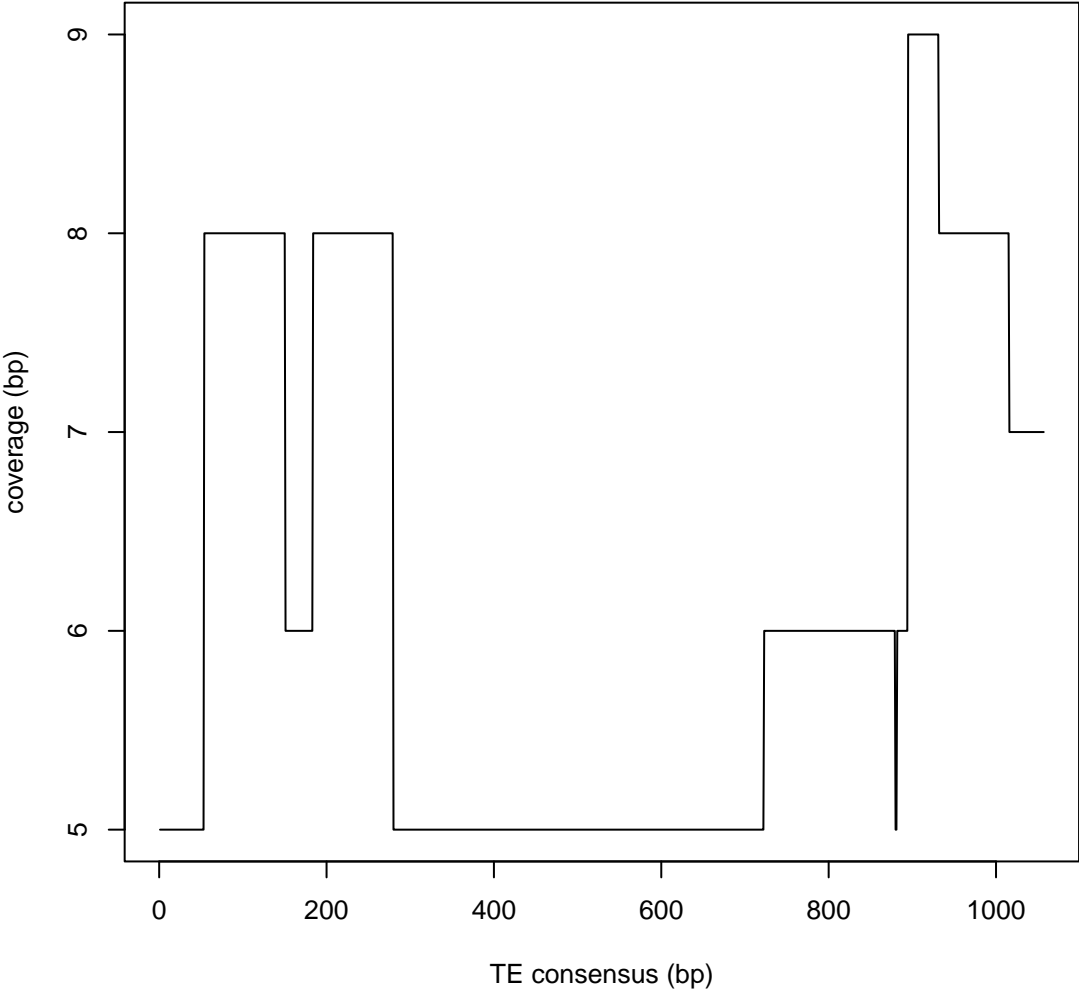


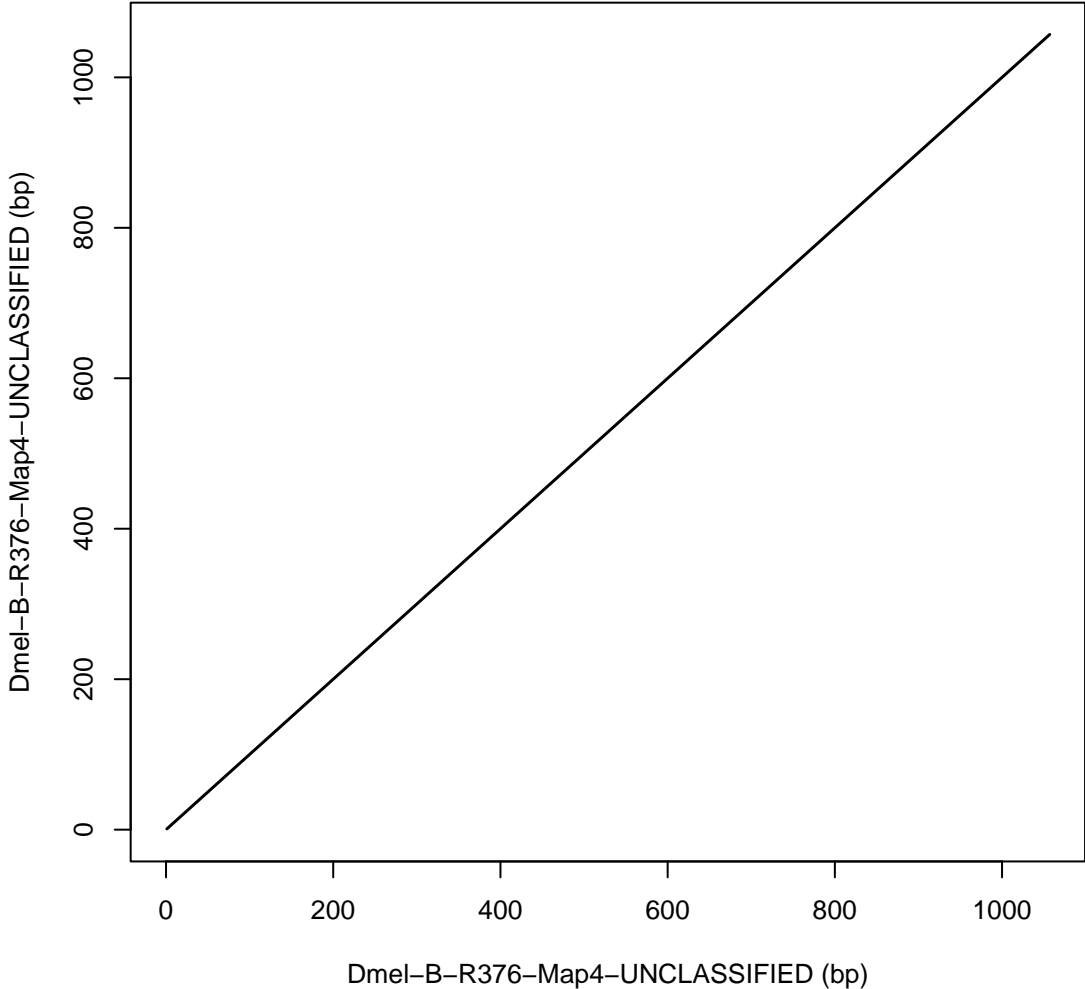
TE: Dmel-B-R376-Map4-UNCLASSIFIED  
consensus size: 1057bp; fragments: 15; full length: 1 (>=951.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

