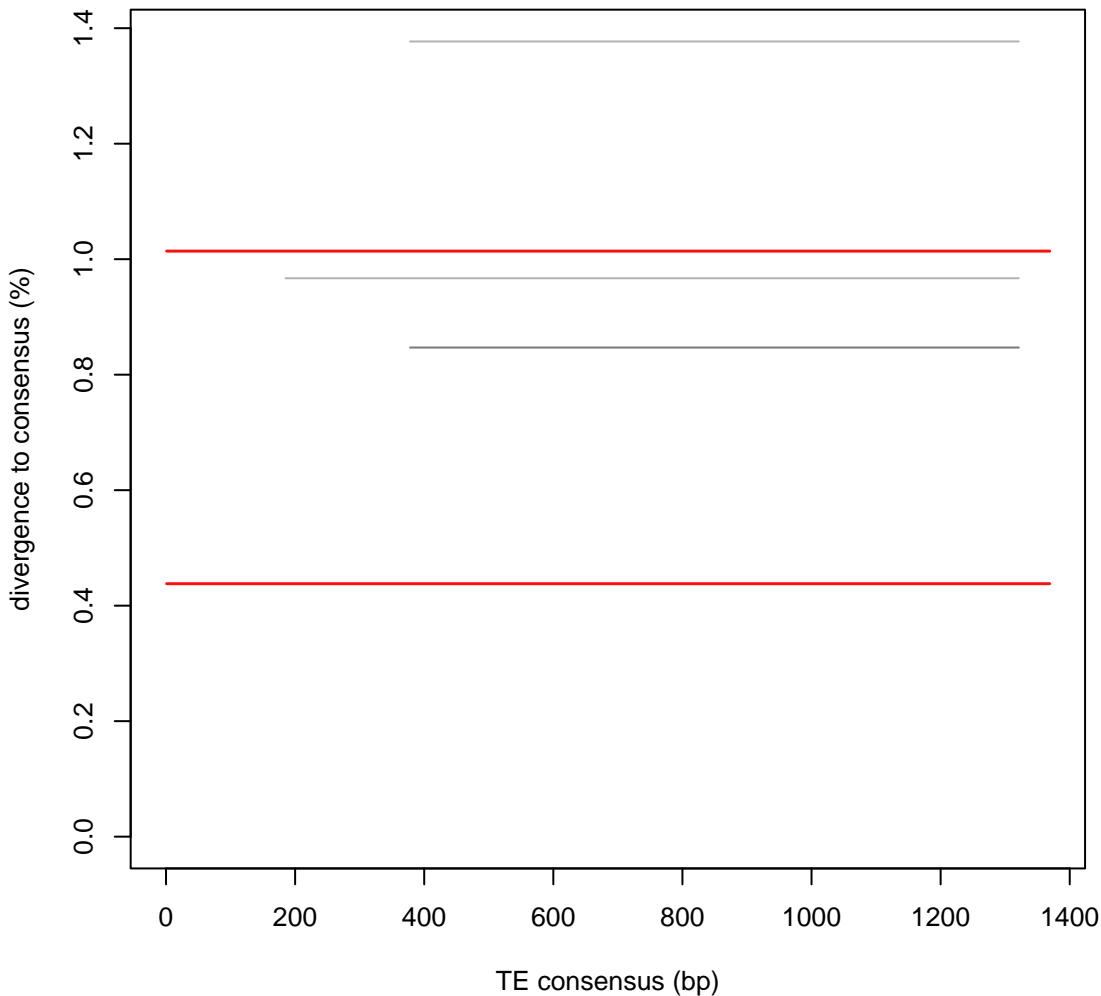
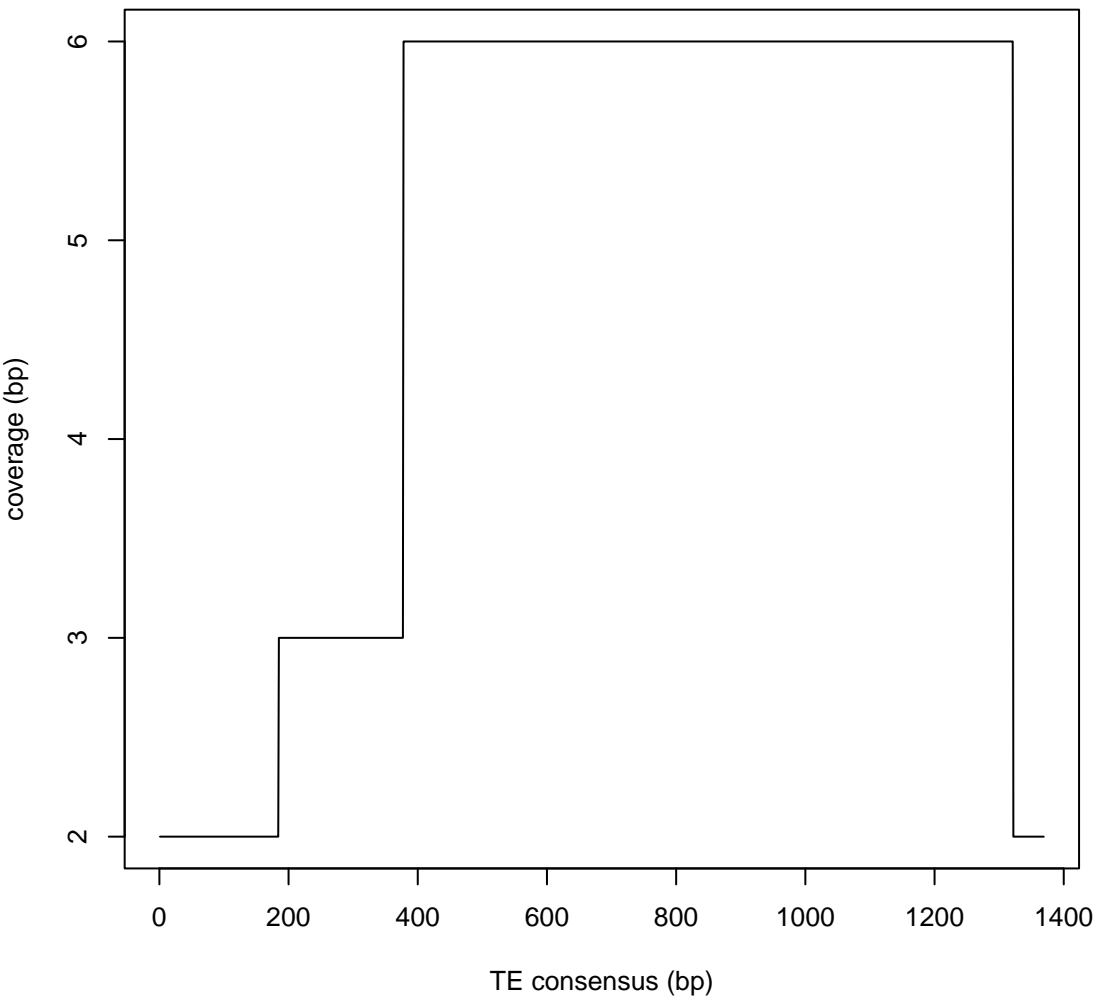


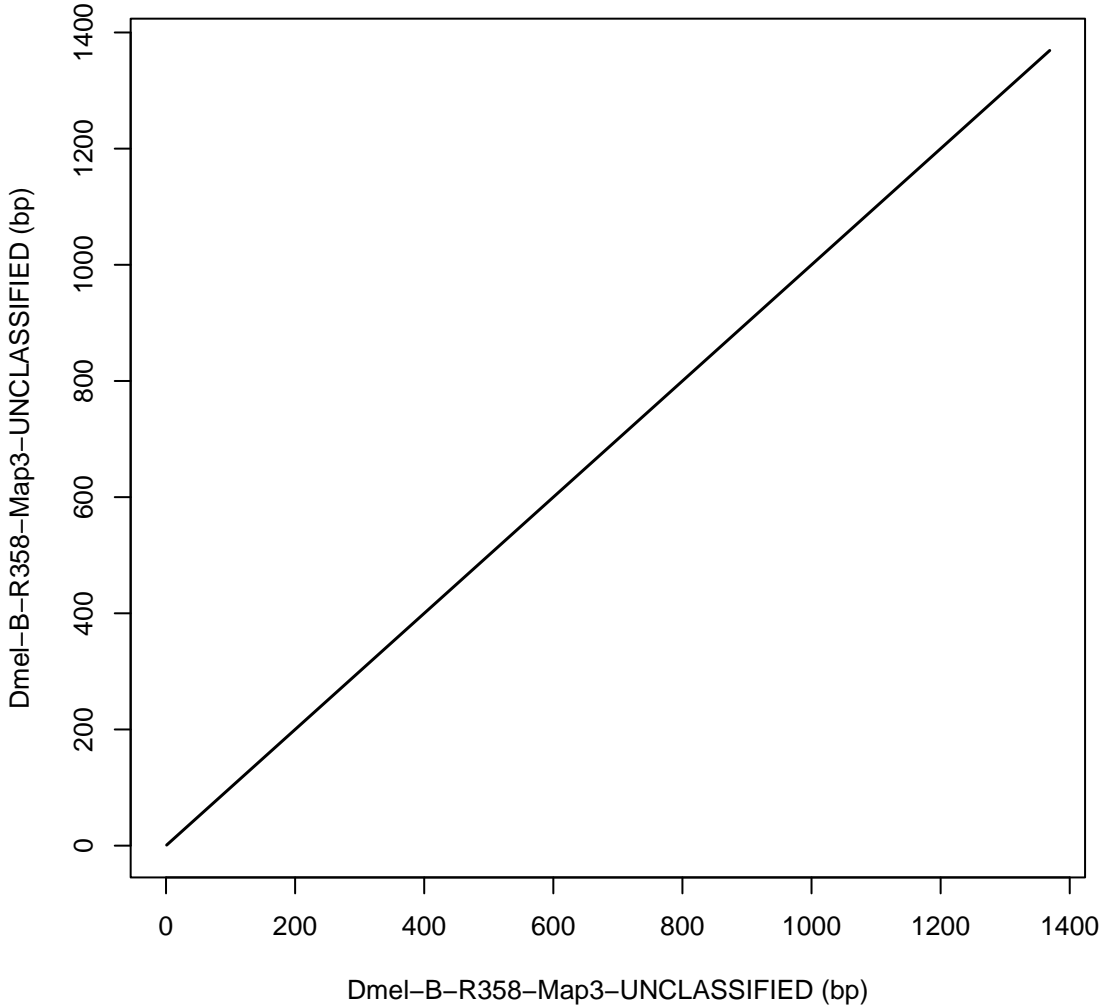
TE: Dmel-B-R358-Map3-UNCLASSIFIED  
consensus size: 1369bp; fragments: 6; full length: 2 (>=1232.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

