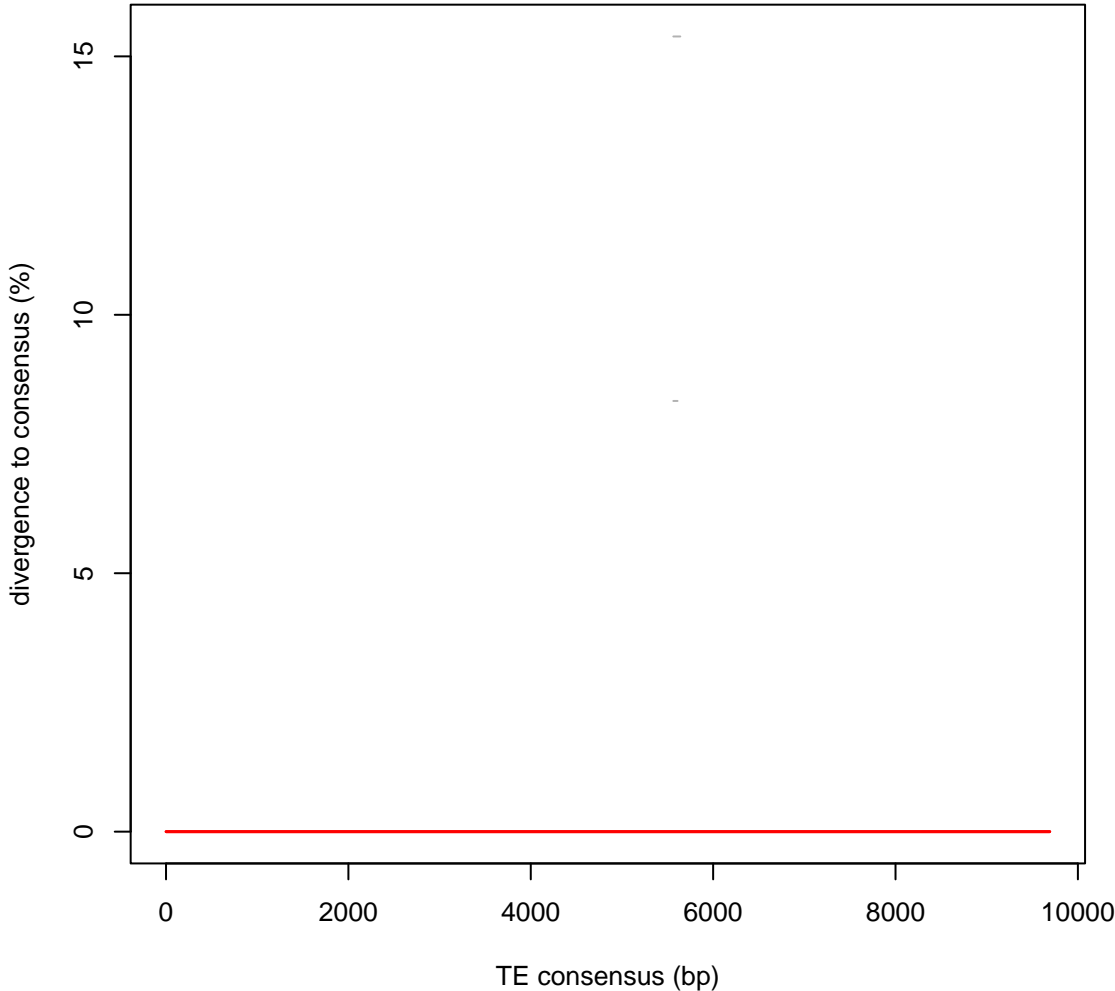
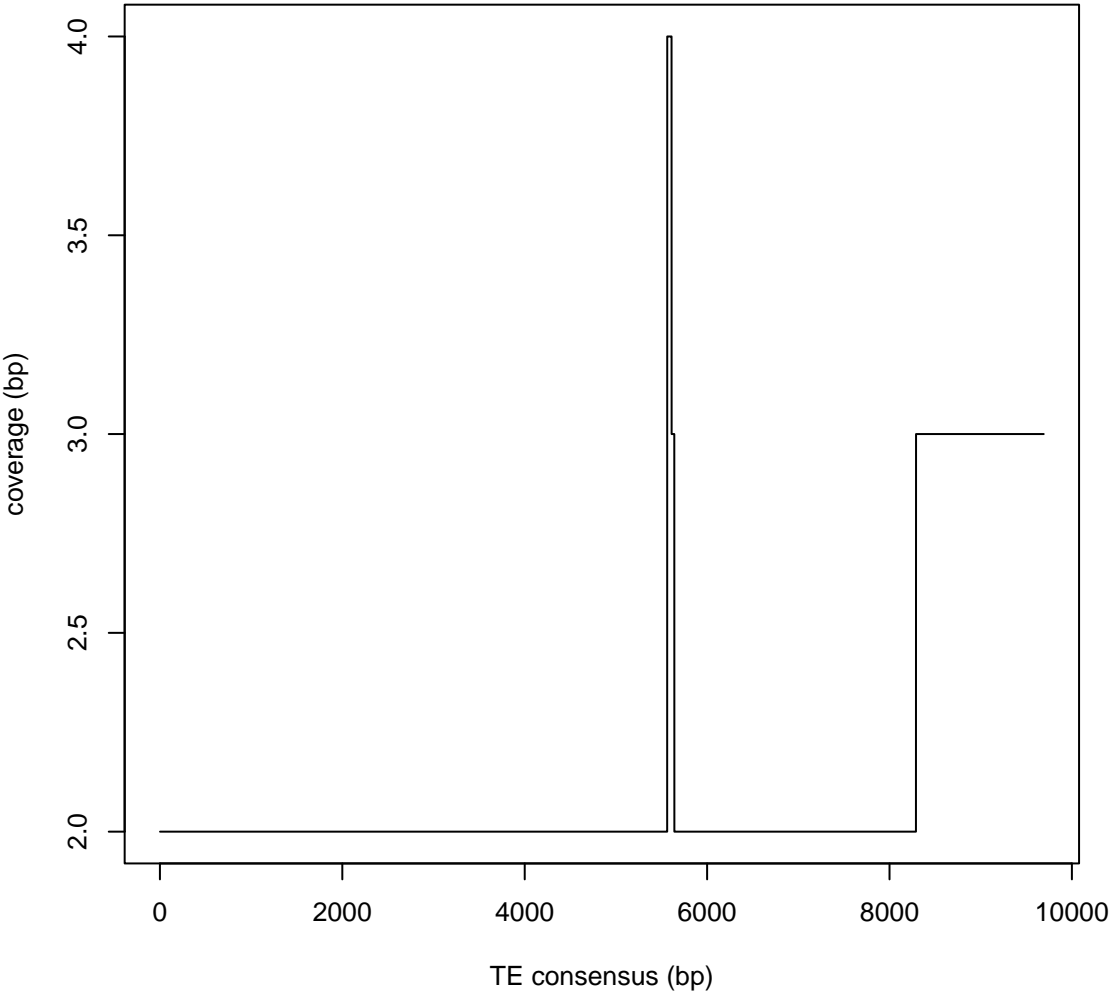


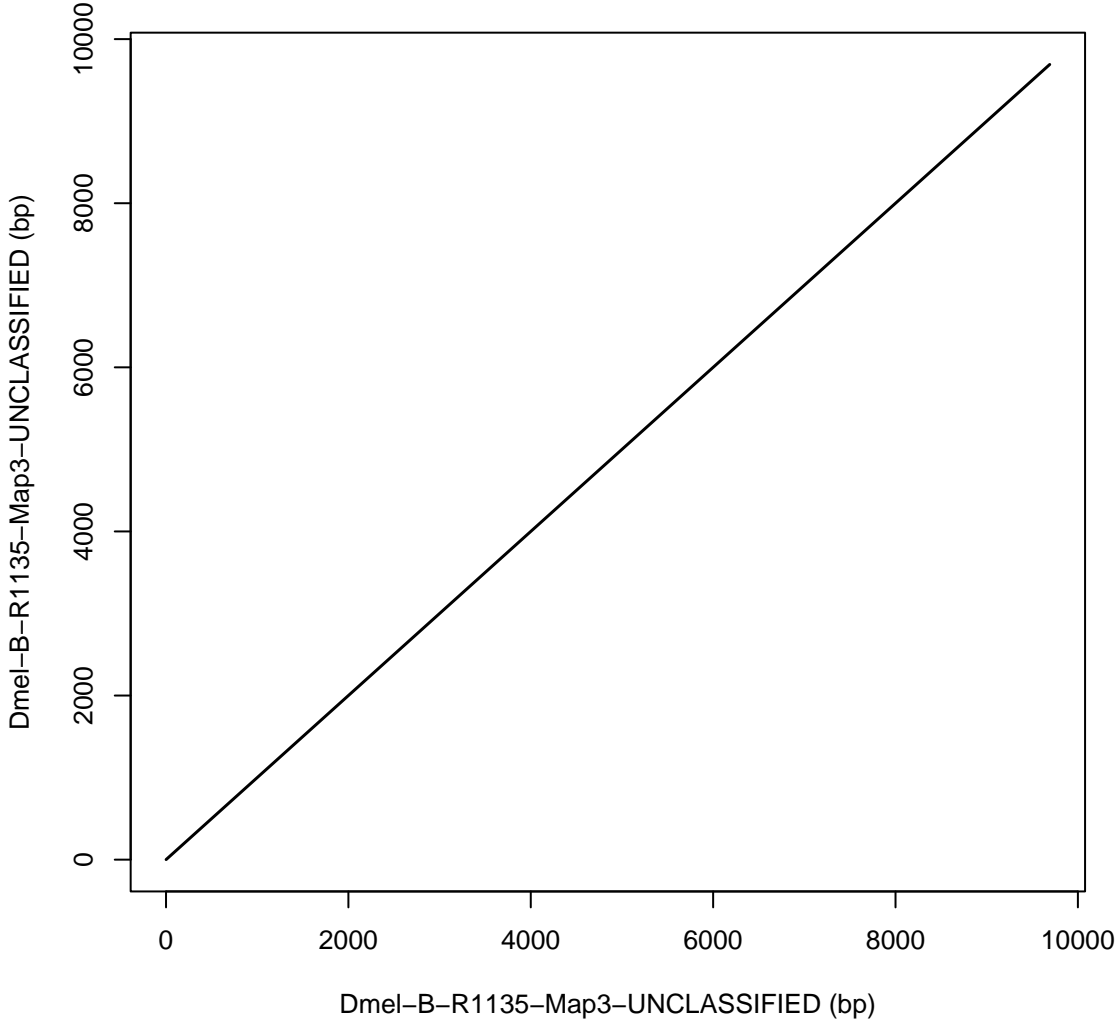
TE: Dmel-B-R1135-Map3-UNCLASSIFIED
consensus size: 9691bp; fragments: 5; full length: 2 (>=8721.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

