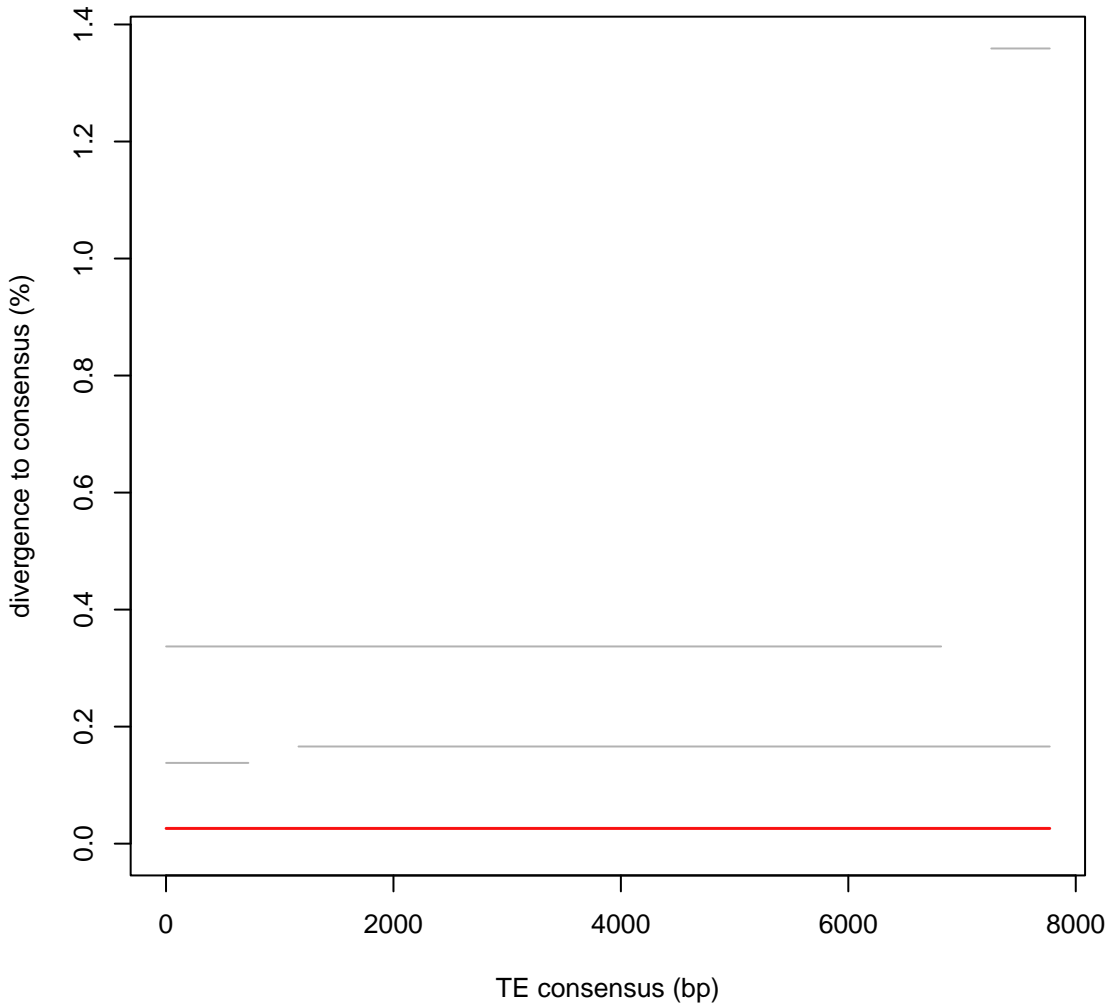
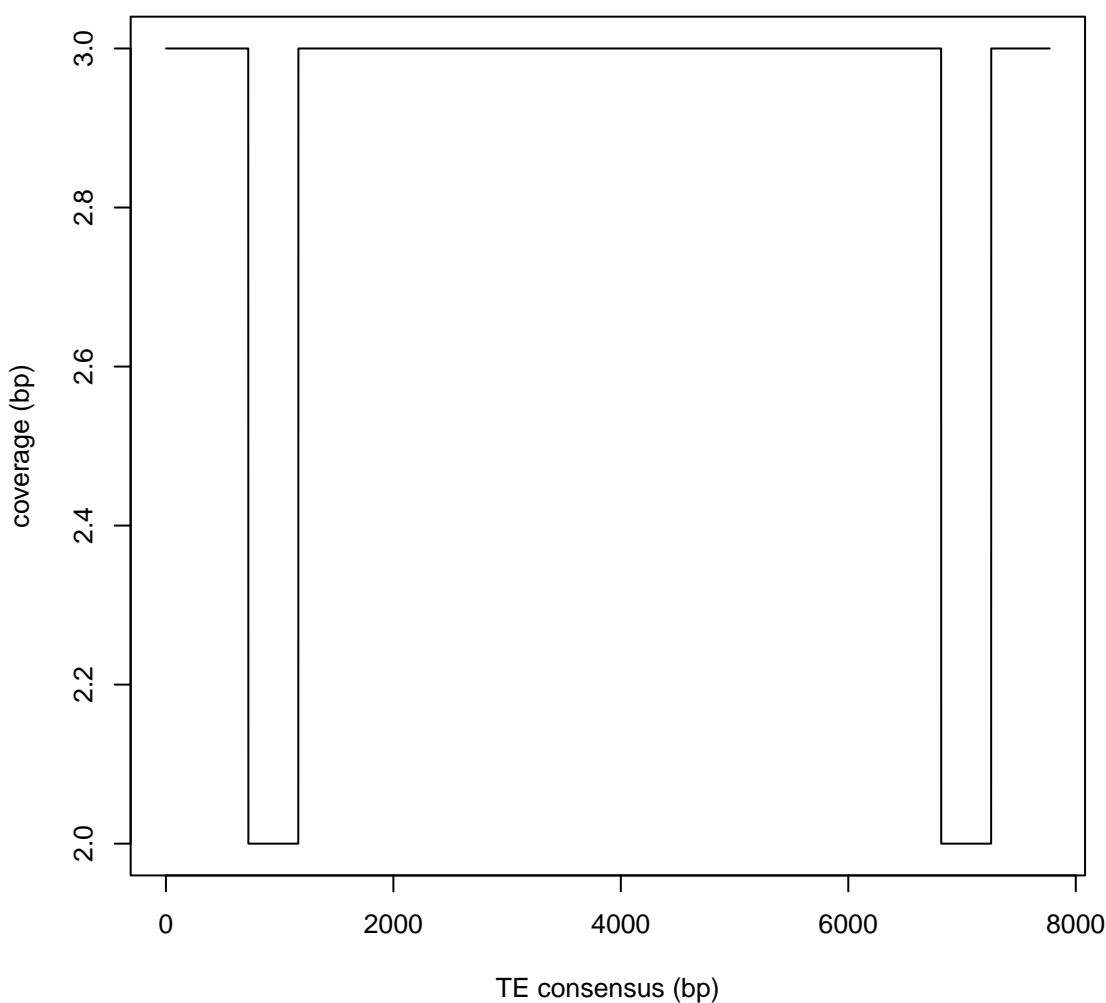


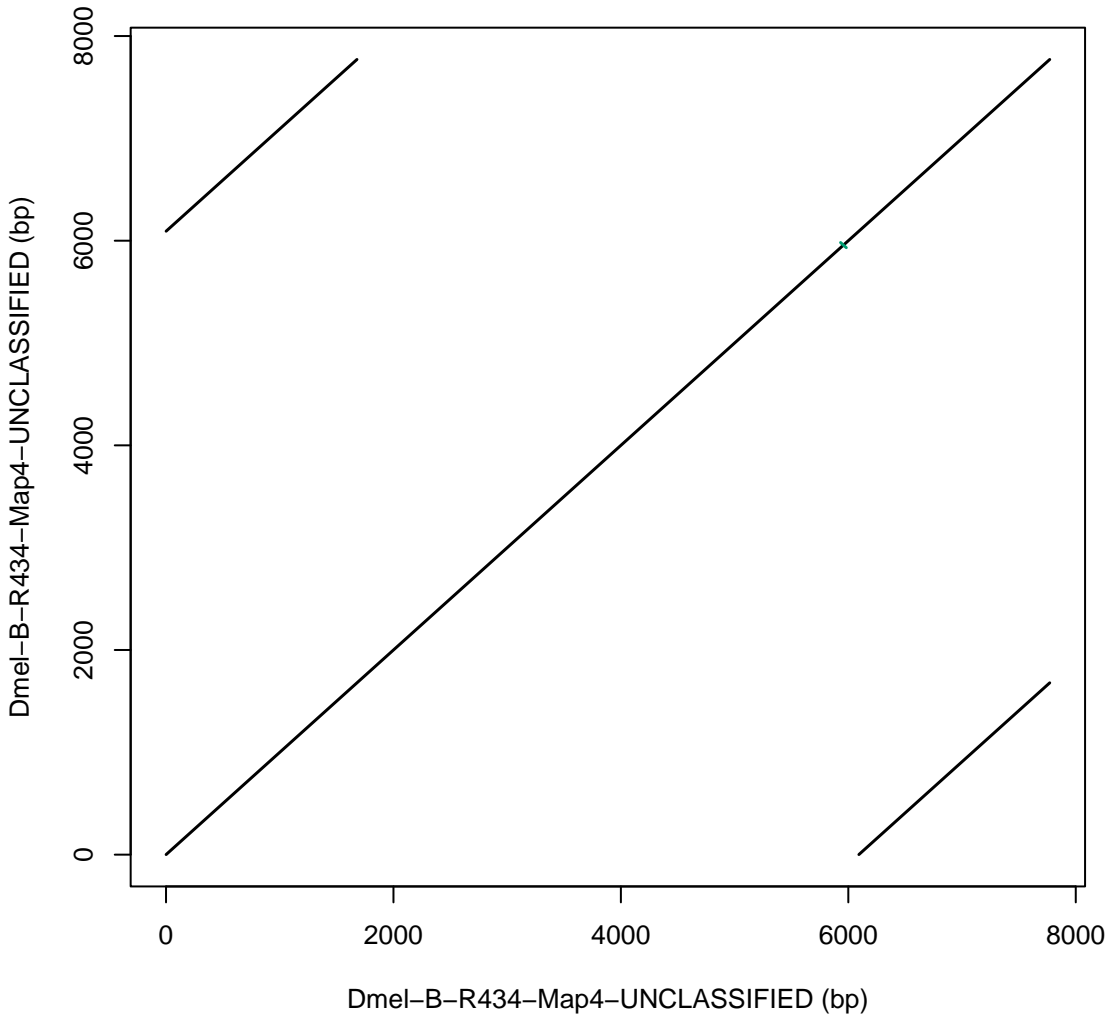
TE: Dmel-B-R434-Map4-UNCLASSIFIED  
consensus size: 7771bp; fragments: 5; full length: 1 (>=6993.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

