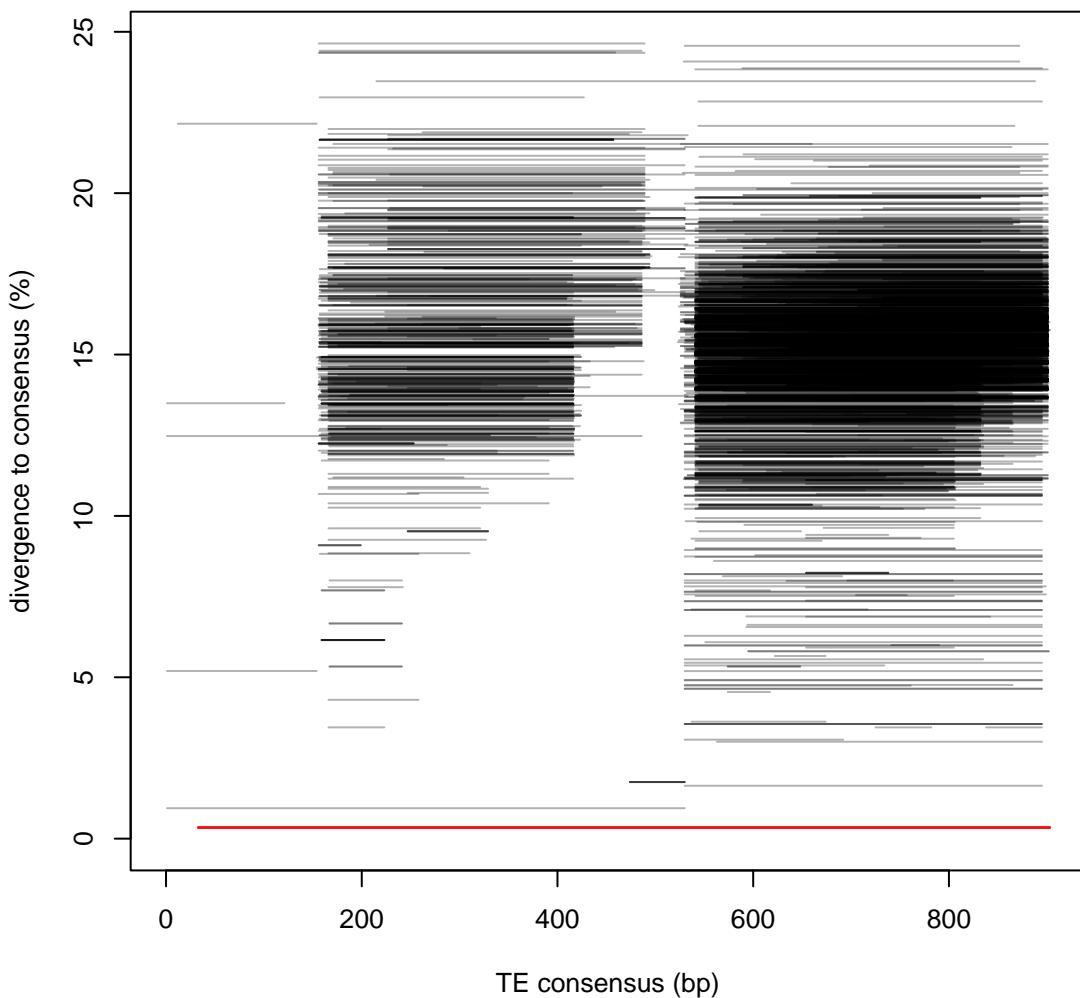
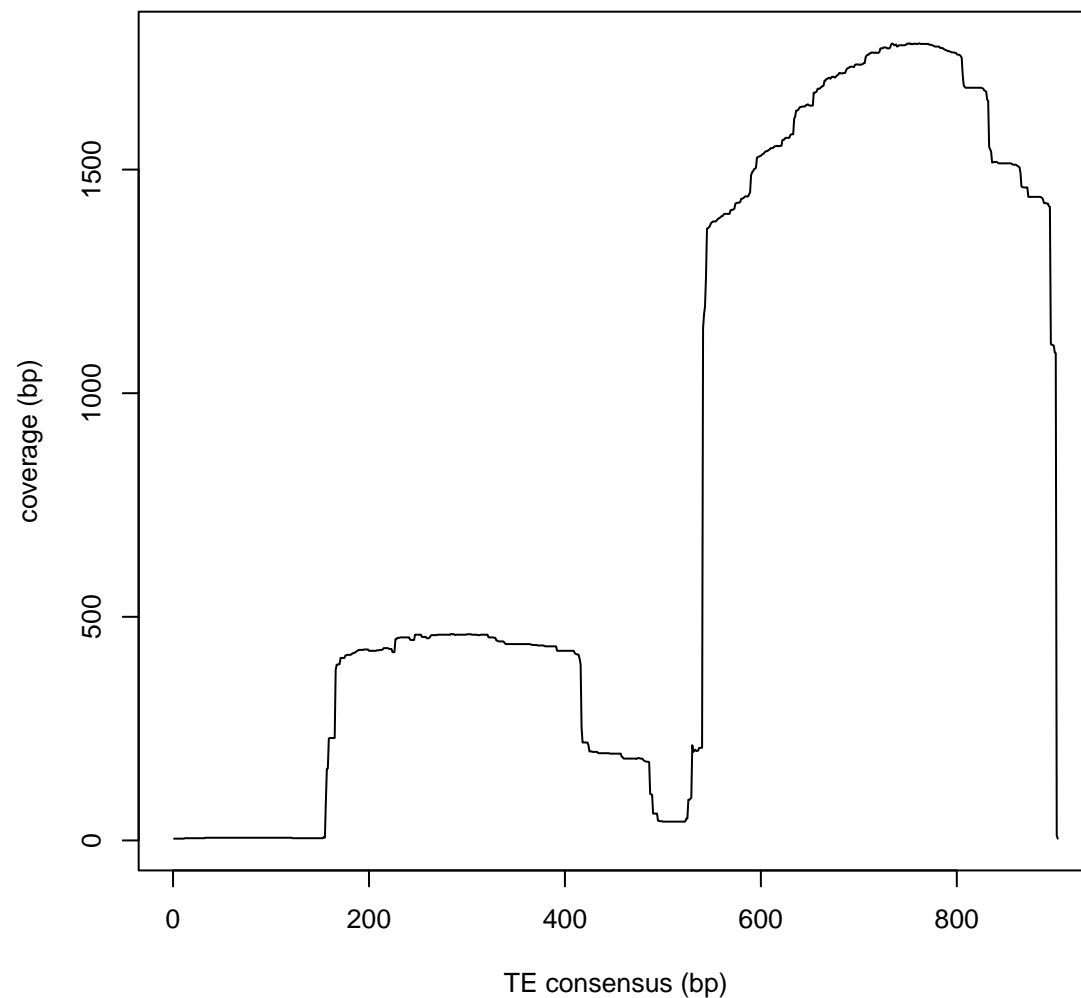


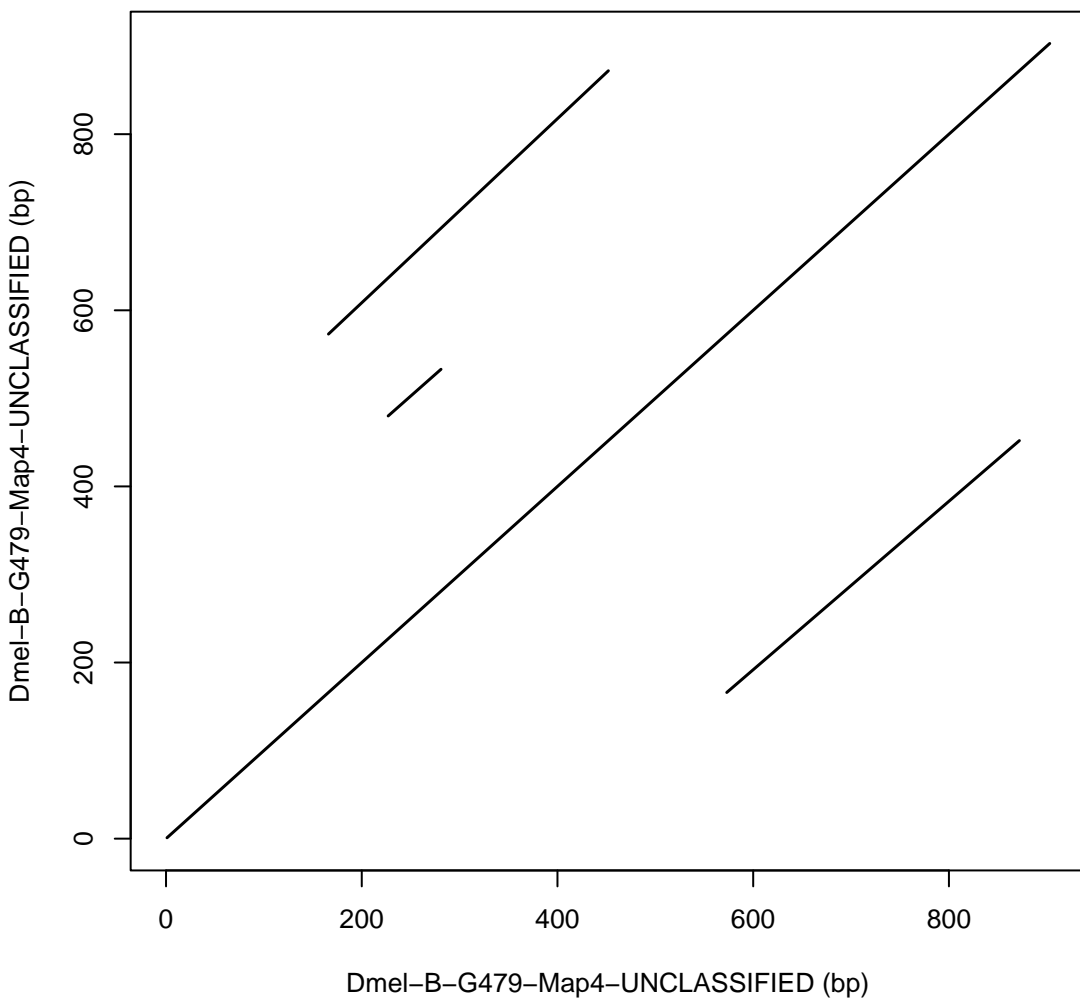
TE: Dmel-B-G479-Map4-UNCLASSIFIED  
consensus size: 903bp; fragments: 2323; full length: 1 (>=812.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

