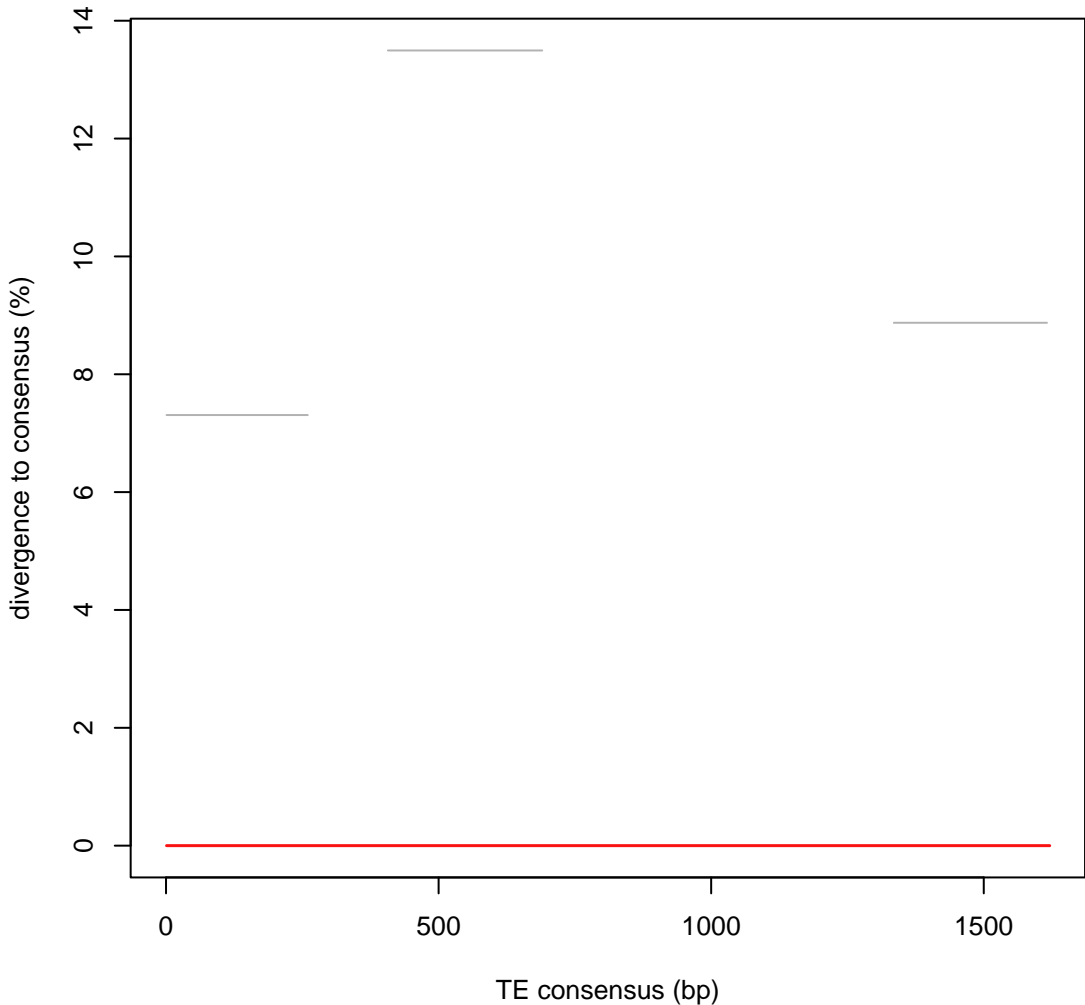
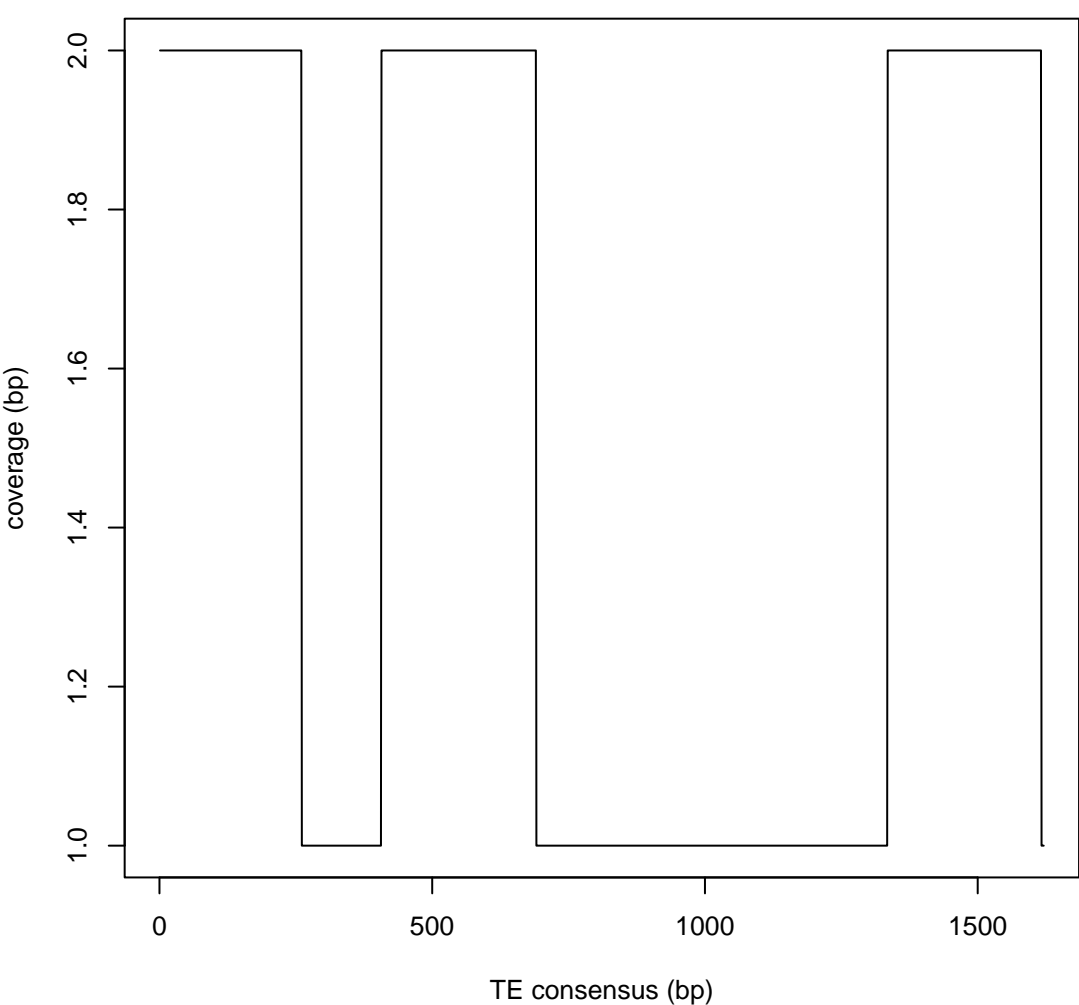


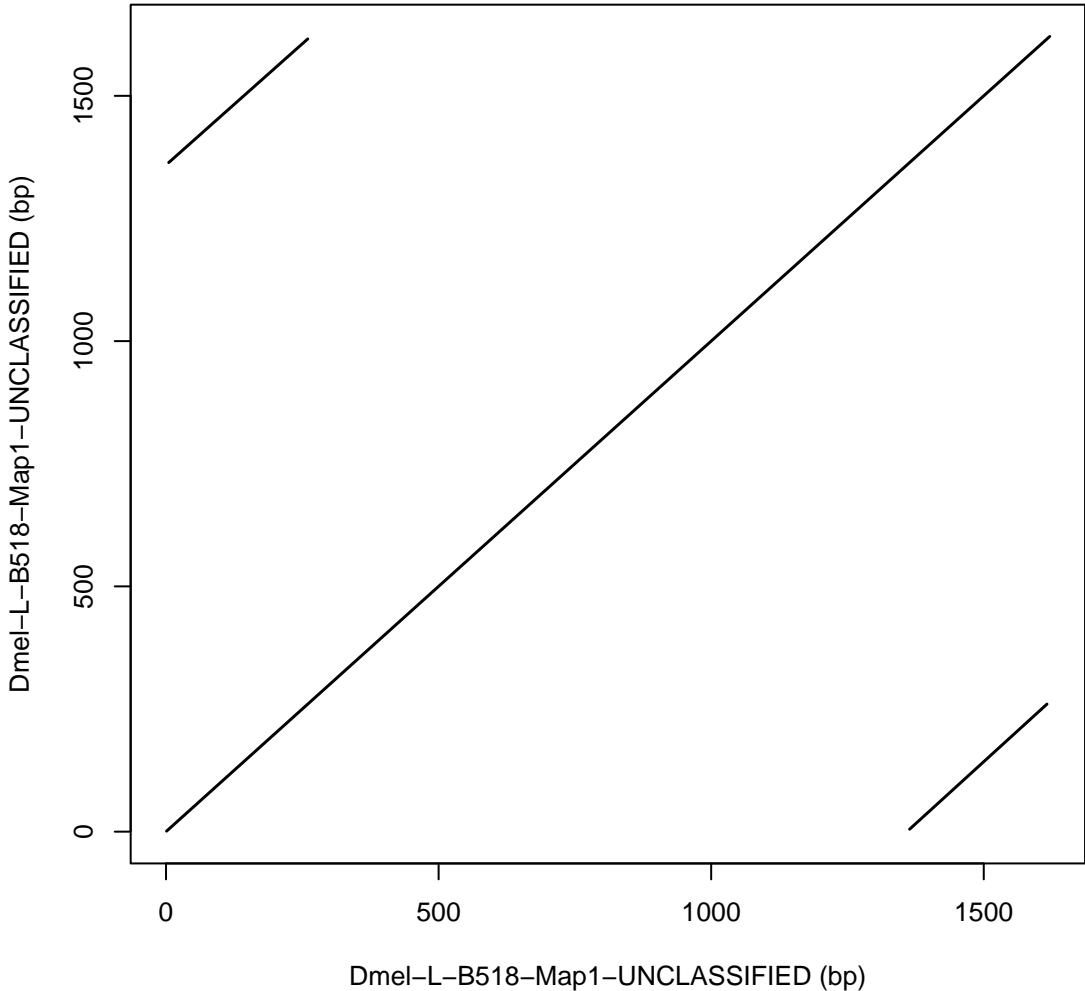
TE: Dmel-L-B518-Map1-UNCLASSIFIED  
consensus size: 1621bp; fragments: 4; full length: 1 (>=1458.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

