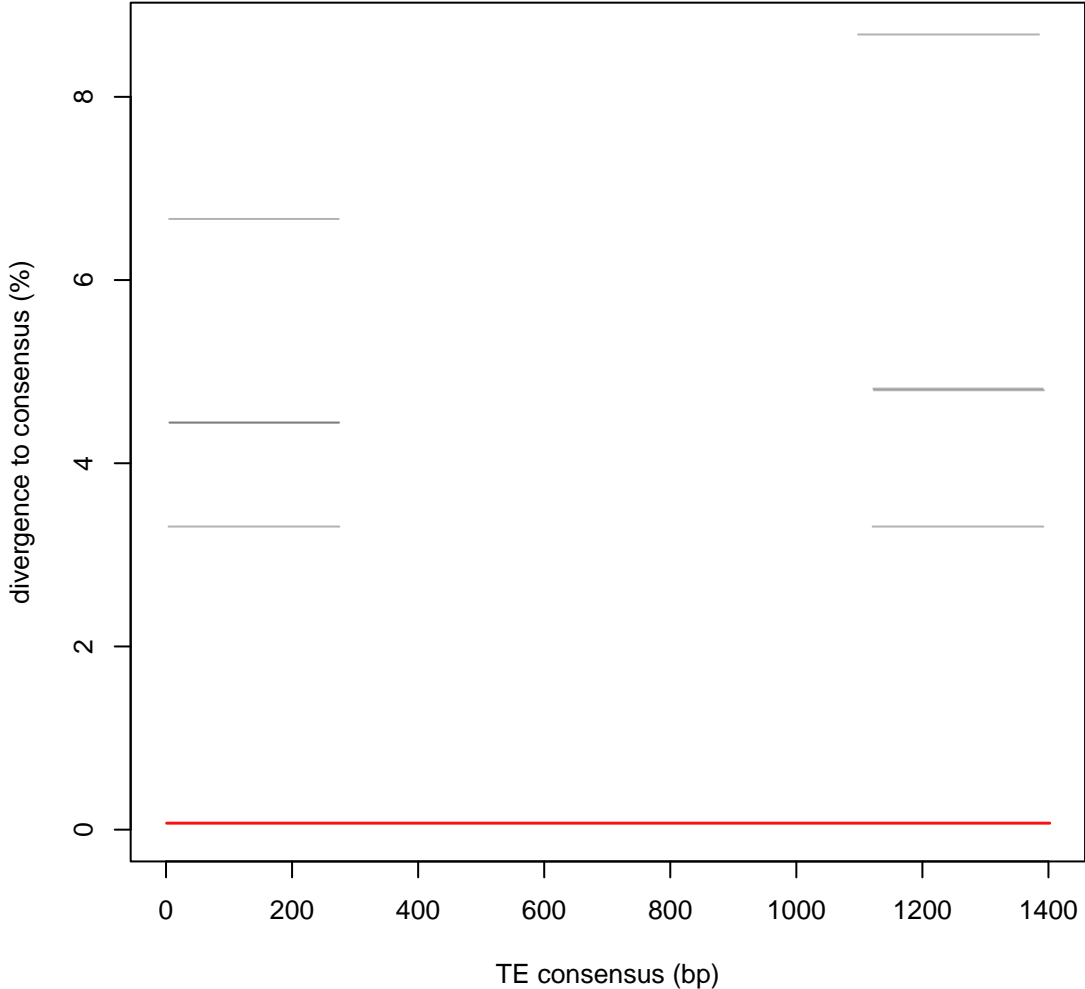
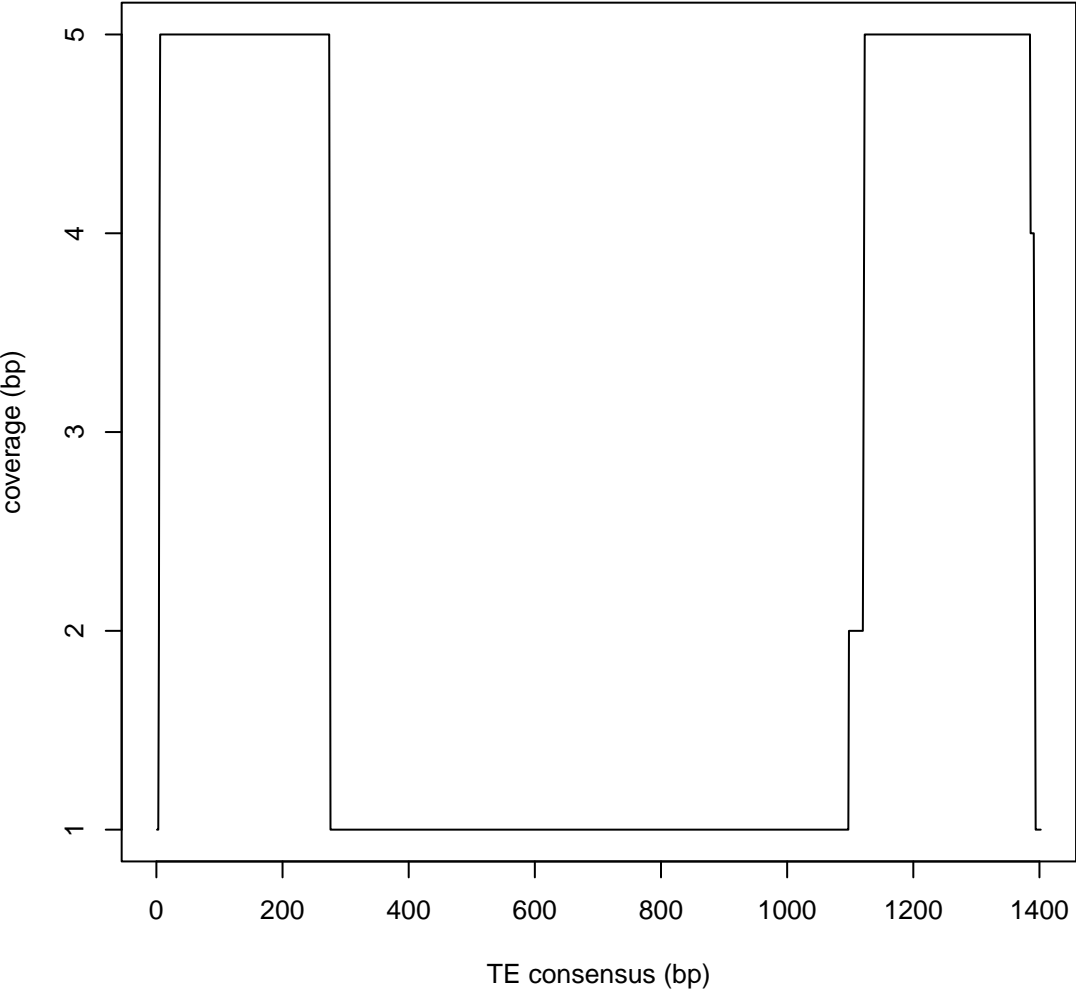


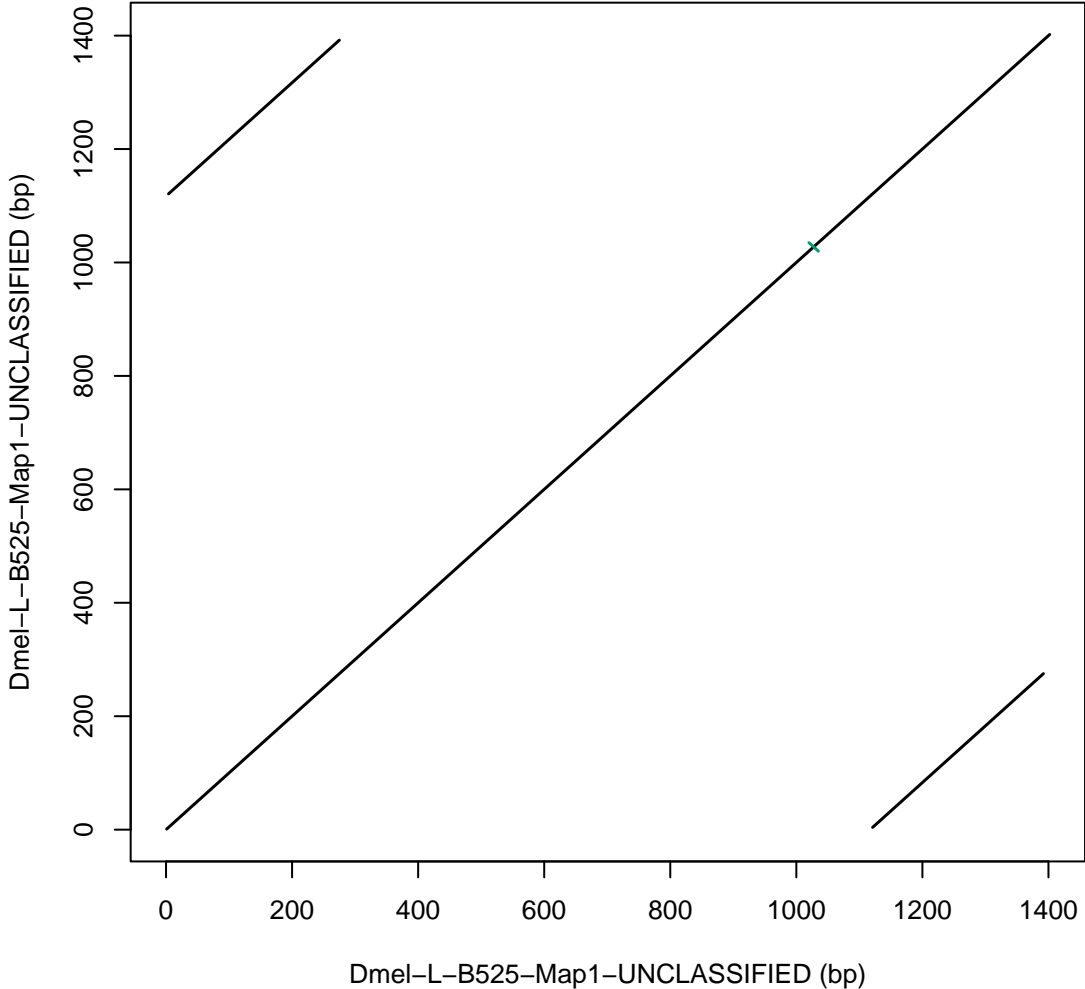
TE: Dmel-L-B525-Map1-UNCLASSIFIED  
consensus size: 1402bp; fragments: 9; full length: 1 (>=1261.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

