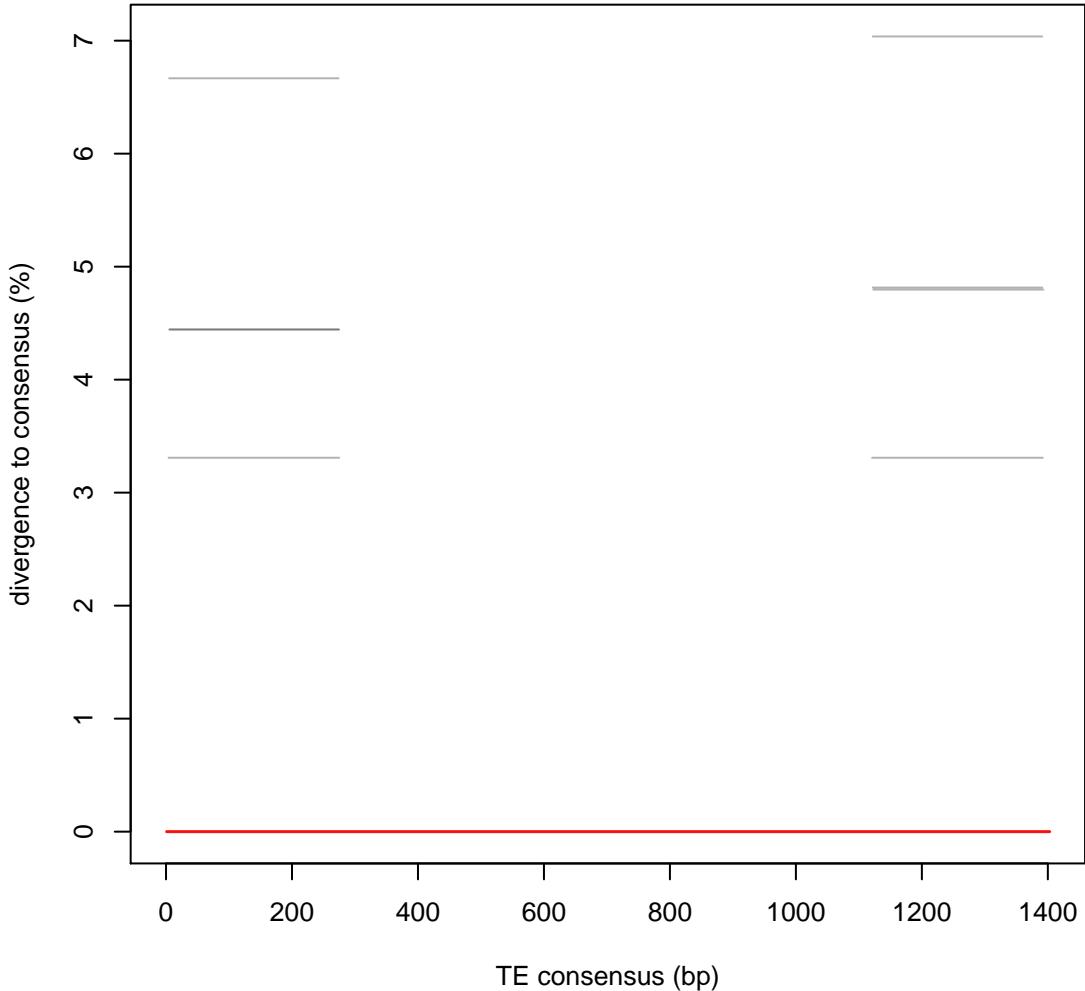
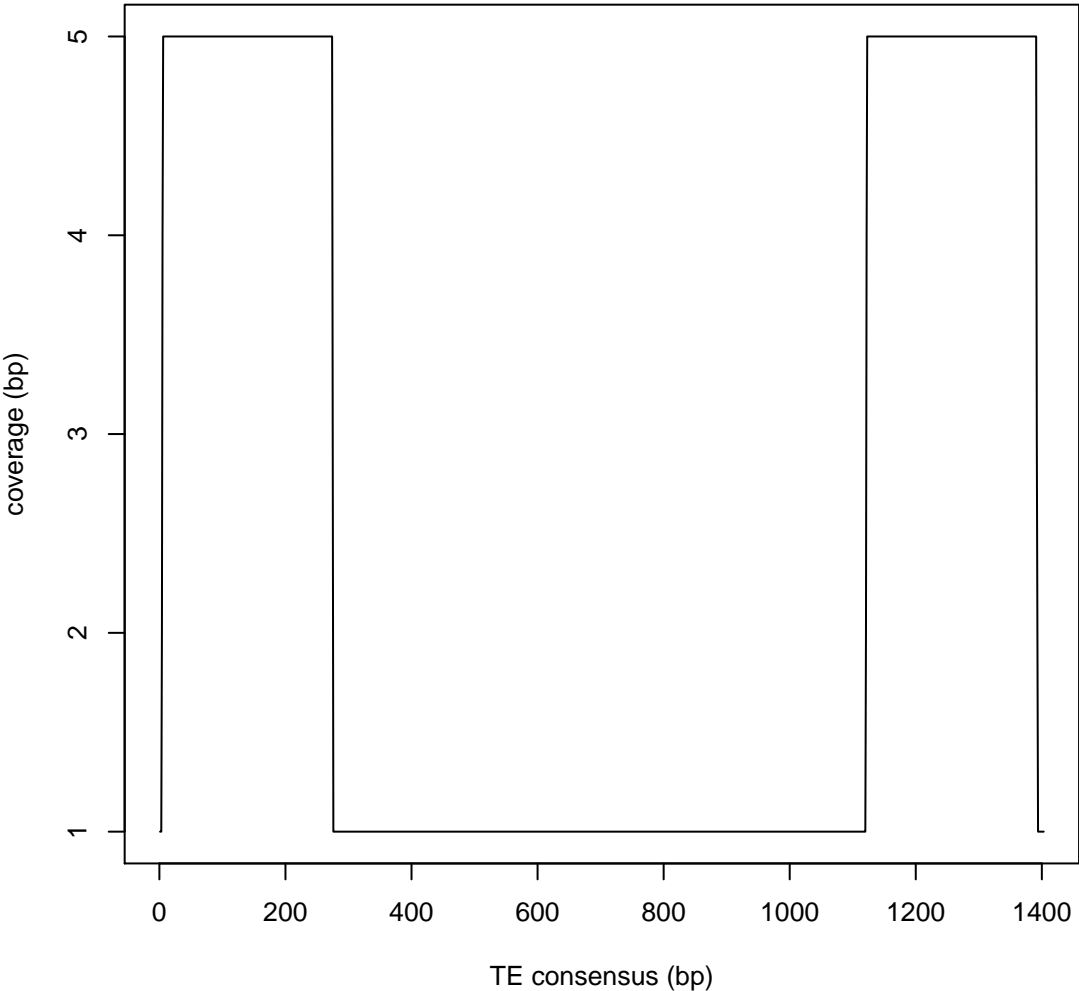


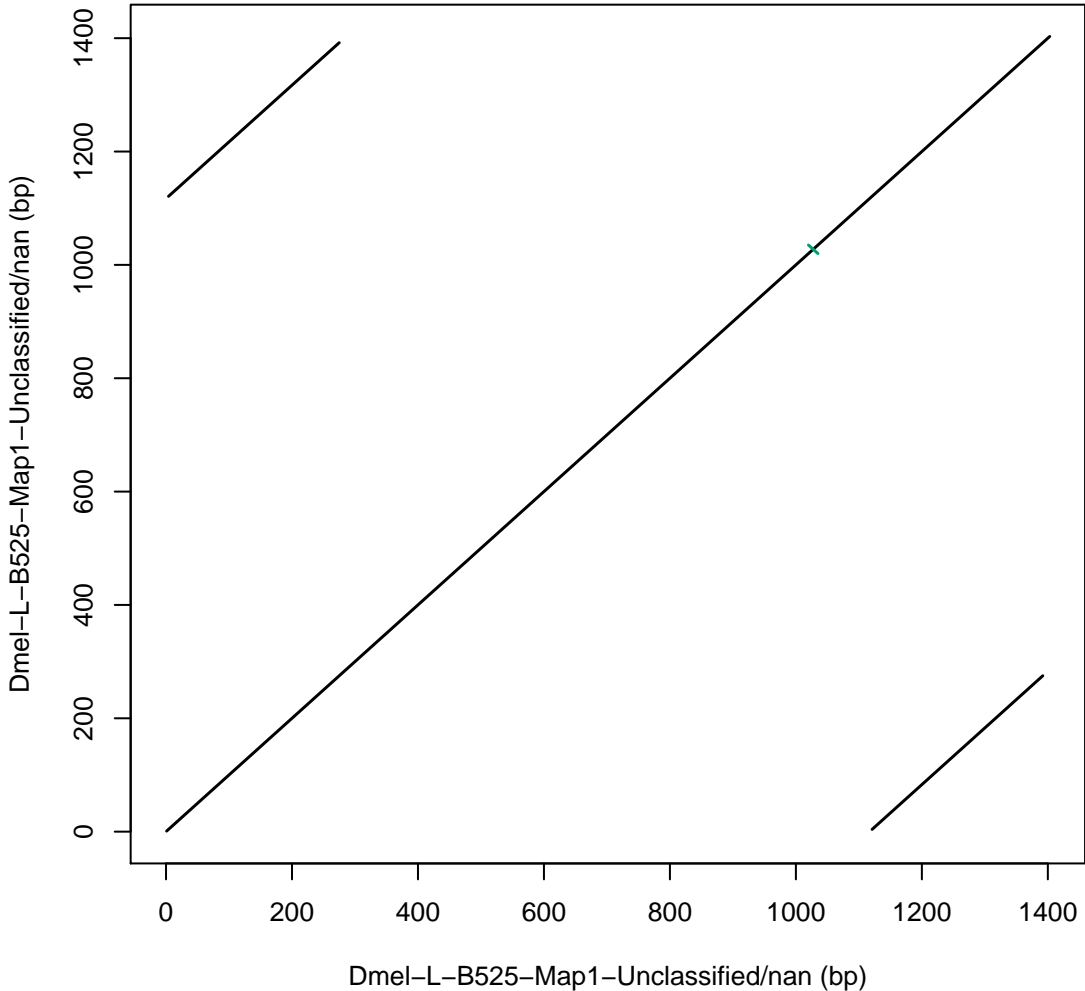
TE: Dmel-L-B525-Map1-Unclassified/nan
consensus size: 1403bp; fragments: 9; full length: 1 (>=1262.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

