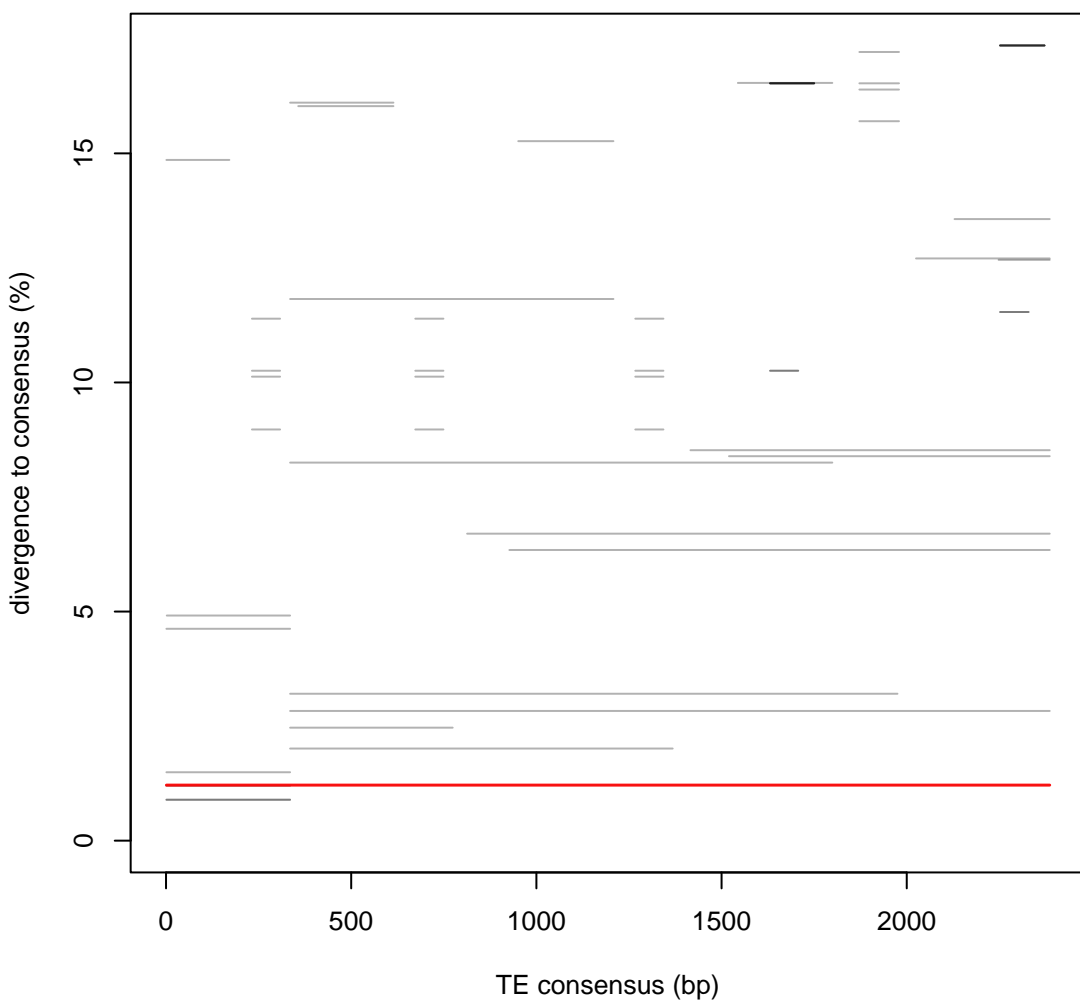
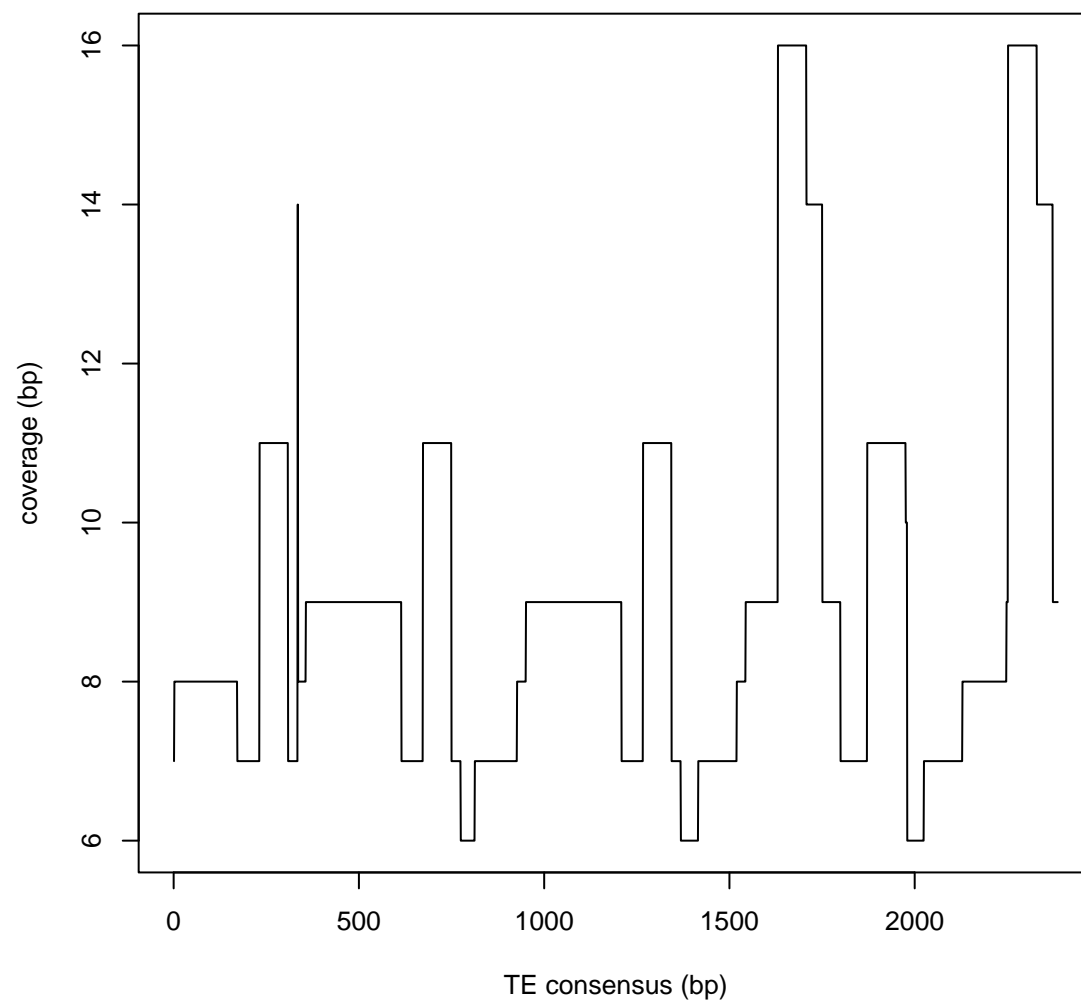


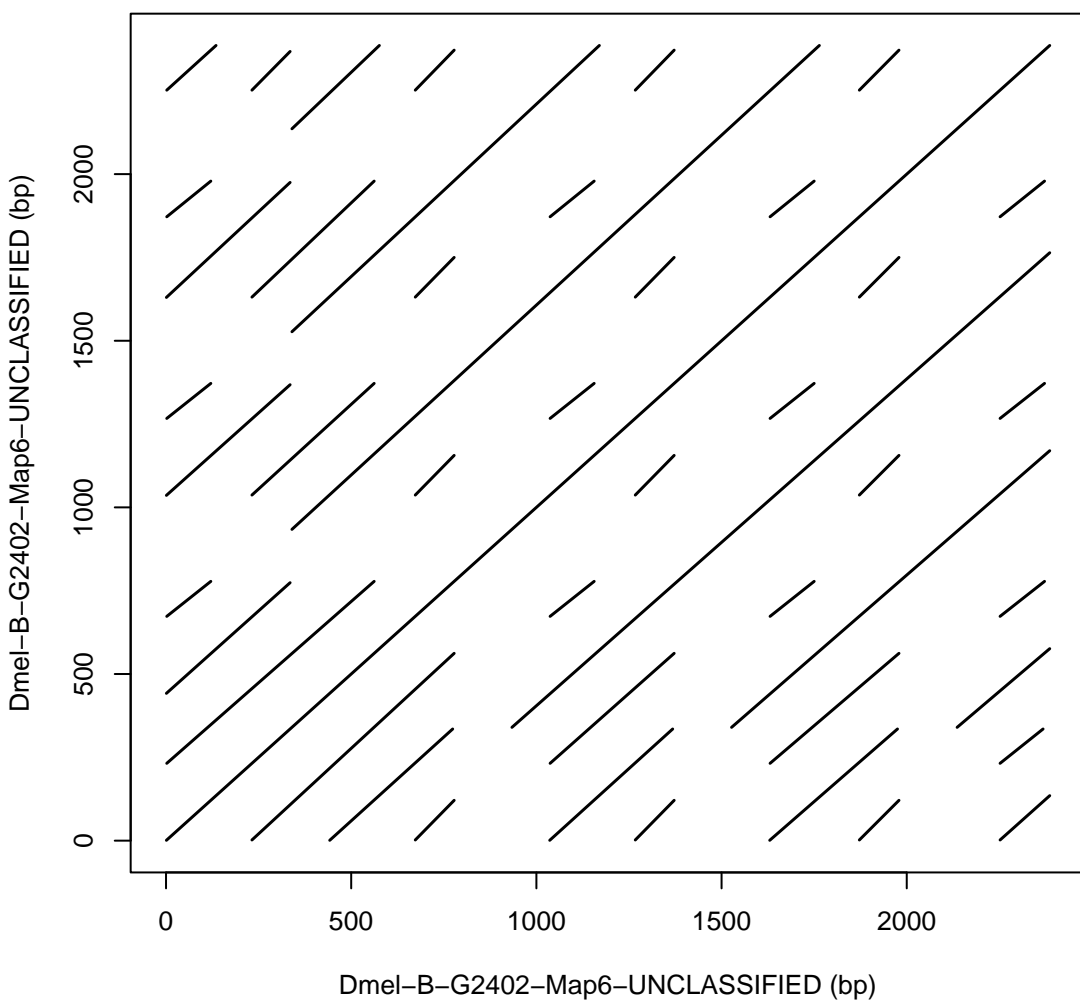
TE: Dmel-B-G2402-Map6-UNCLASSIFIED
consensus size: 2386bp; fragments: 55; full length: 1 (≥ 2147.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

