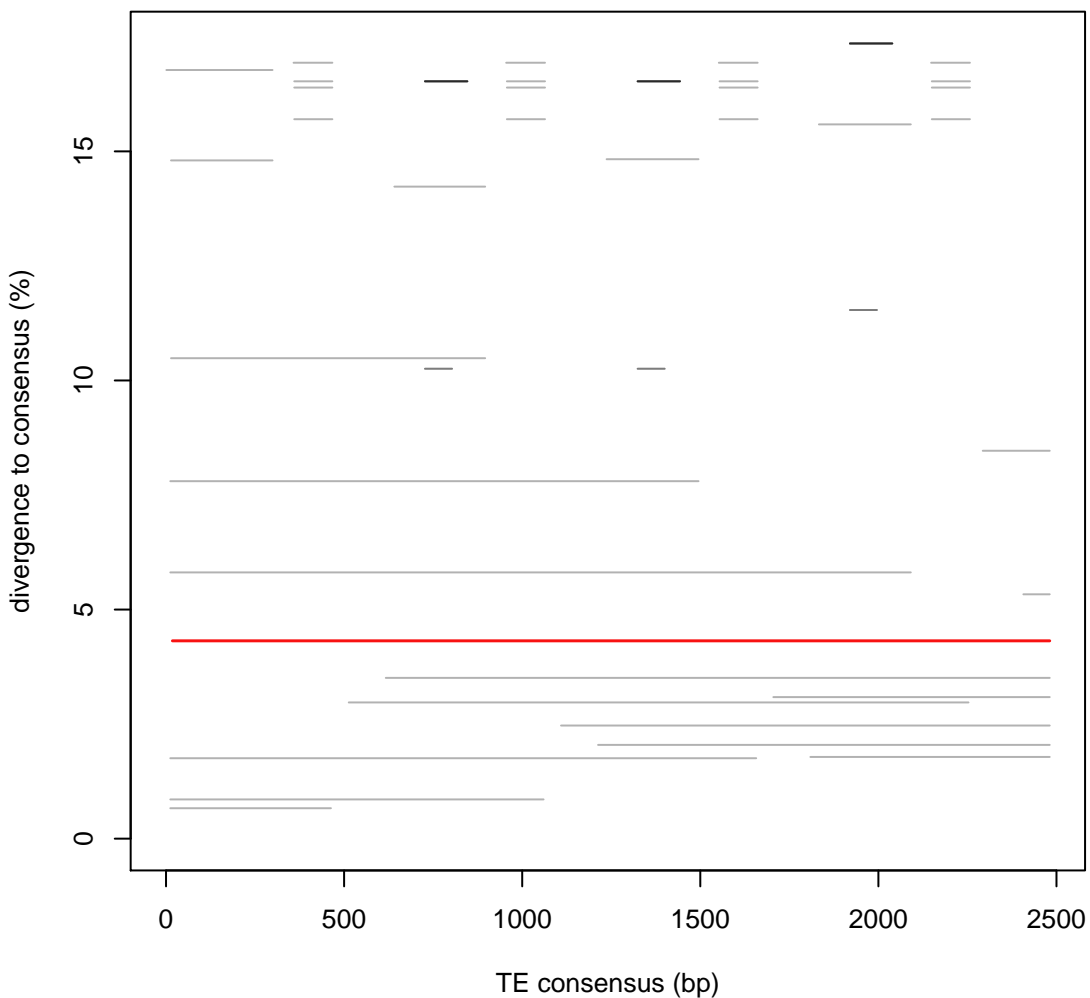
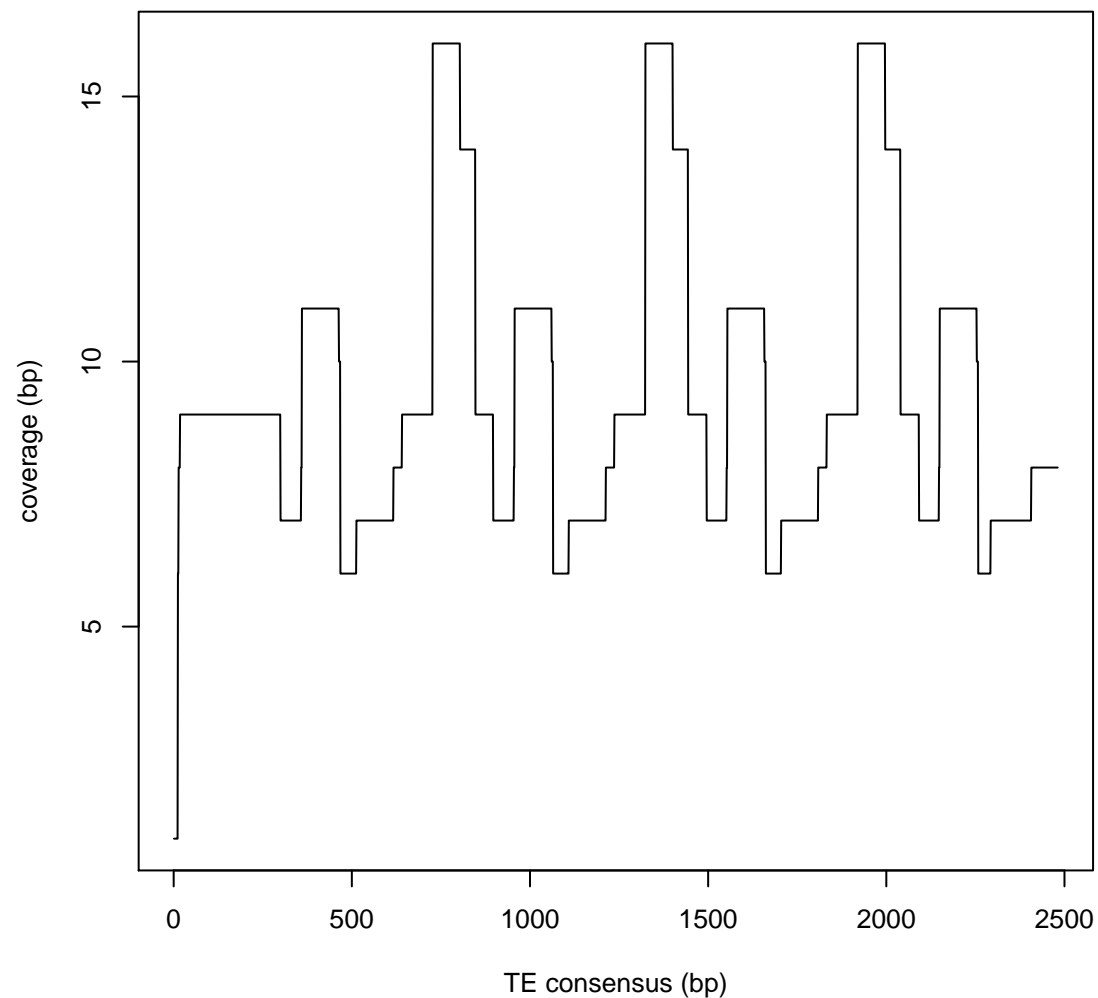


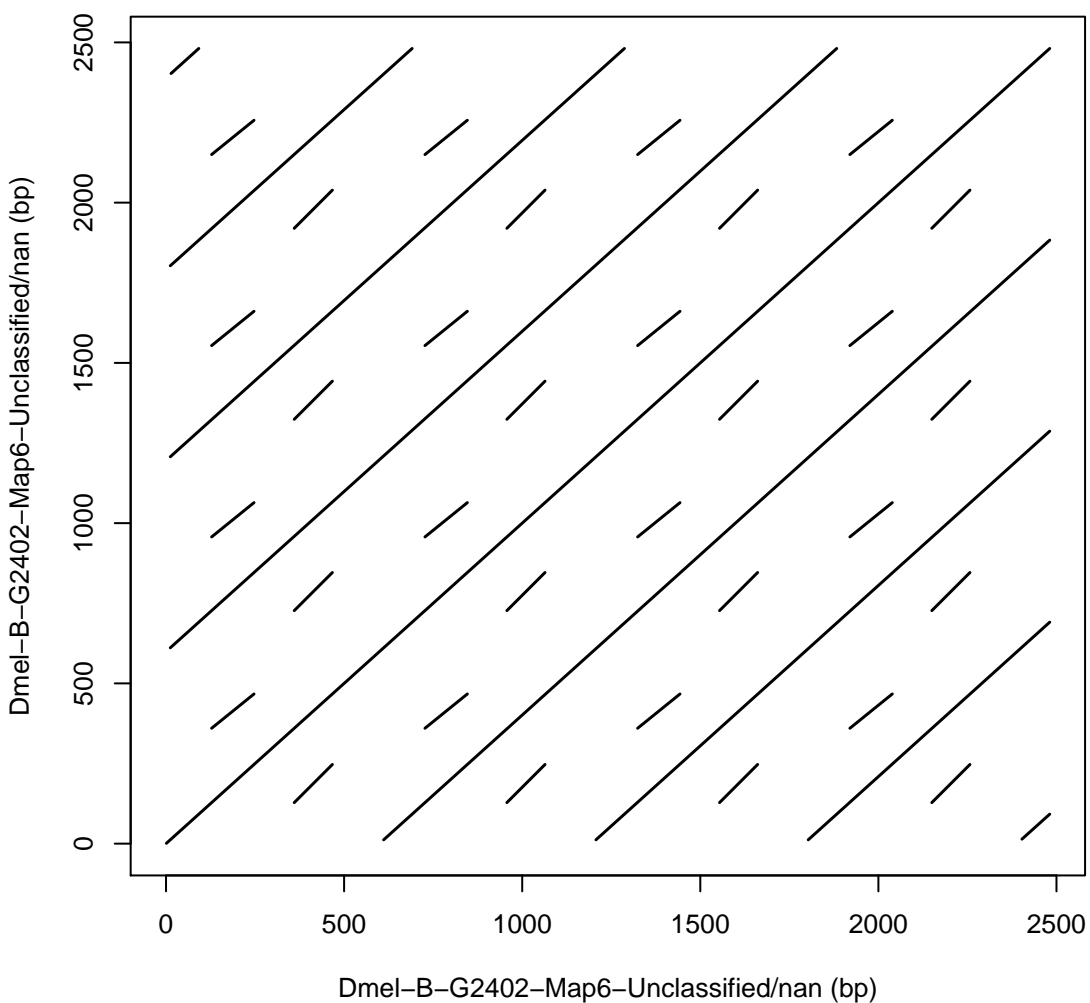
TE: Dmel-B-G2402-Map6-Unclassified/nan
consensus size: 2481bp; fragments: 57; full length: 1 (>=2232.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

