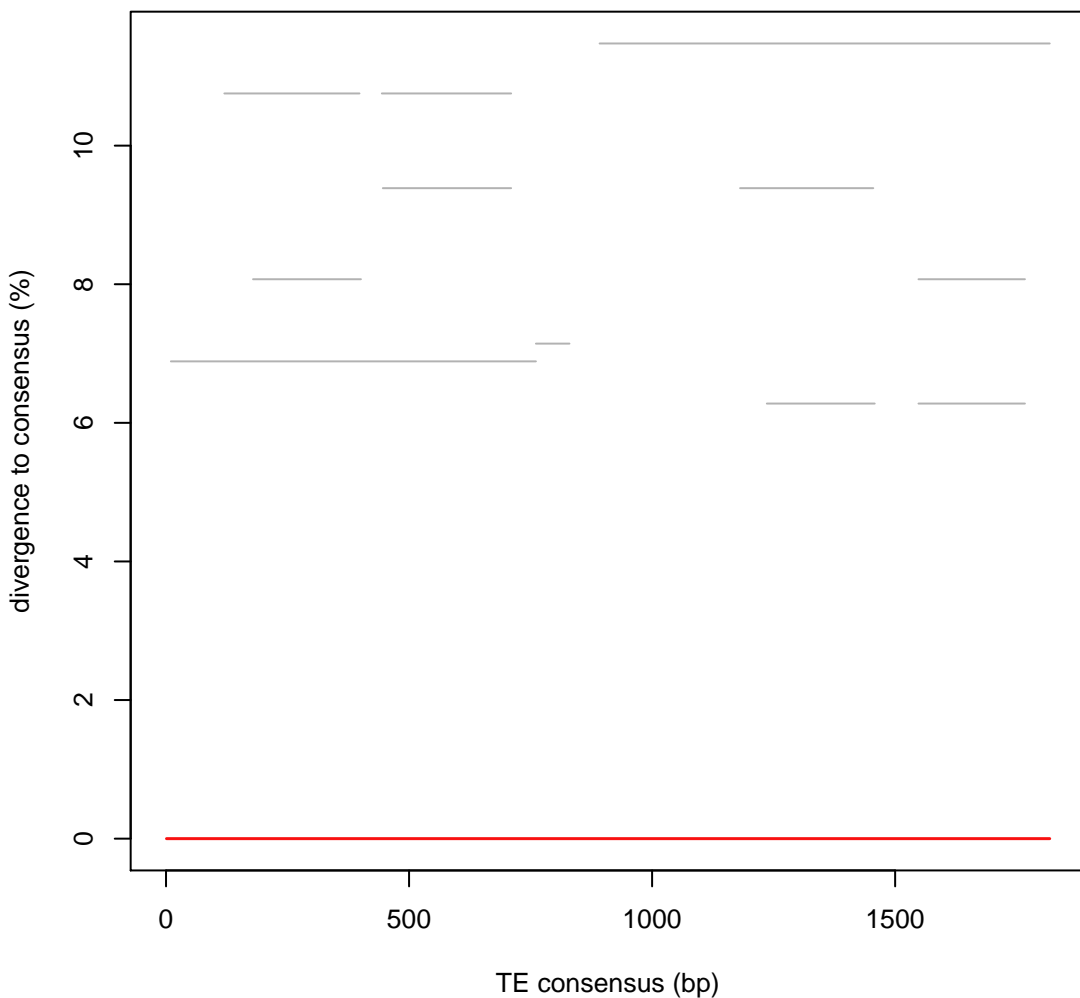
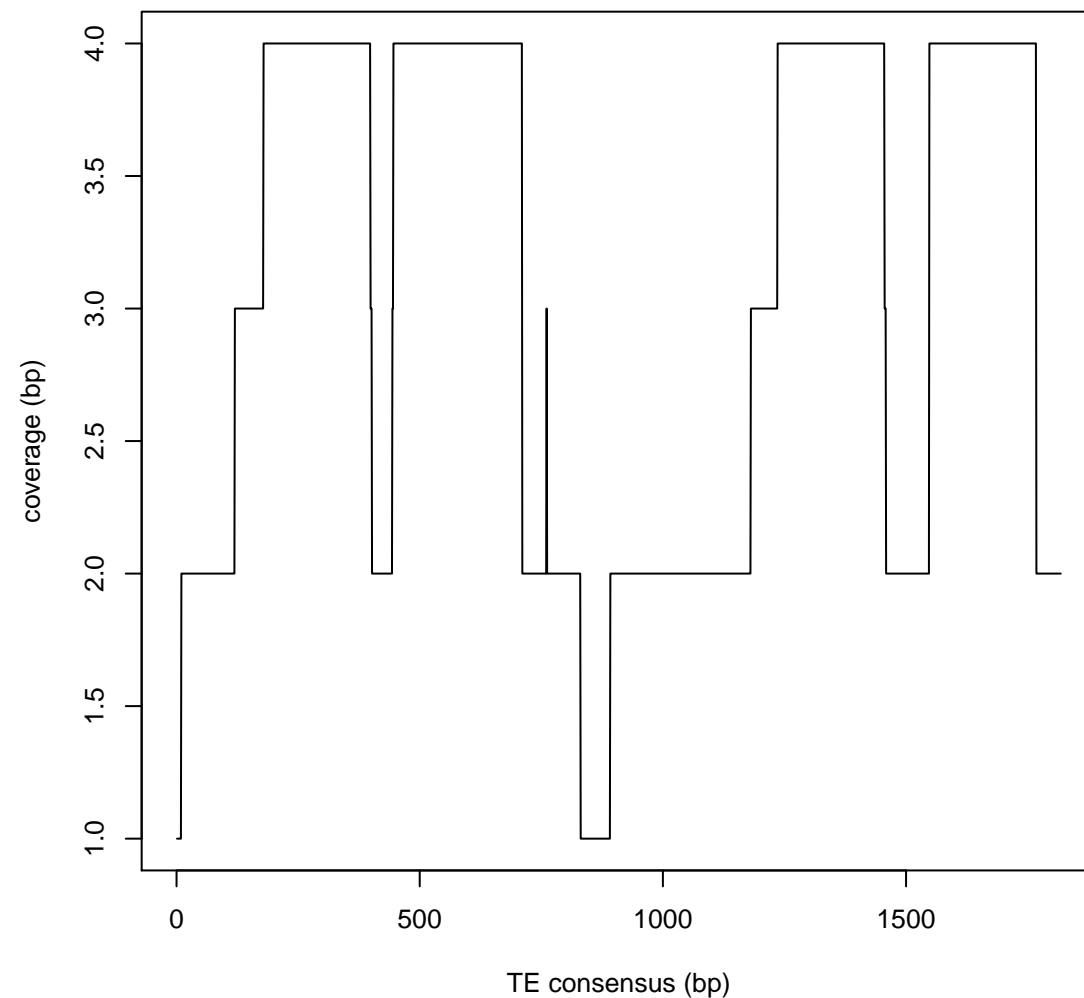


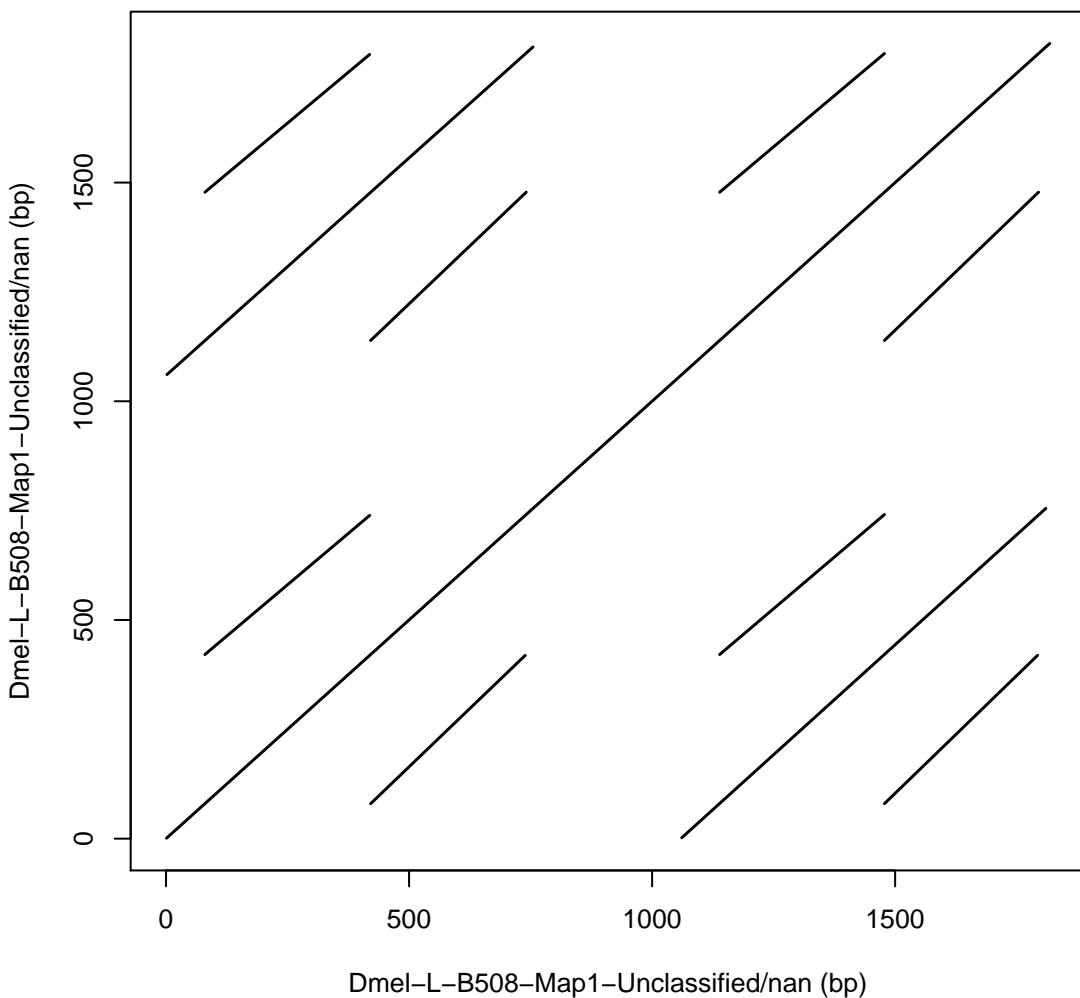
TE: Dmel-L-B508-Map1-Unclassified/nan
consensus size: 1818bp; fragments: 12; full length: 1 (≥ 1636.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

