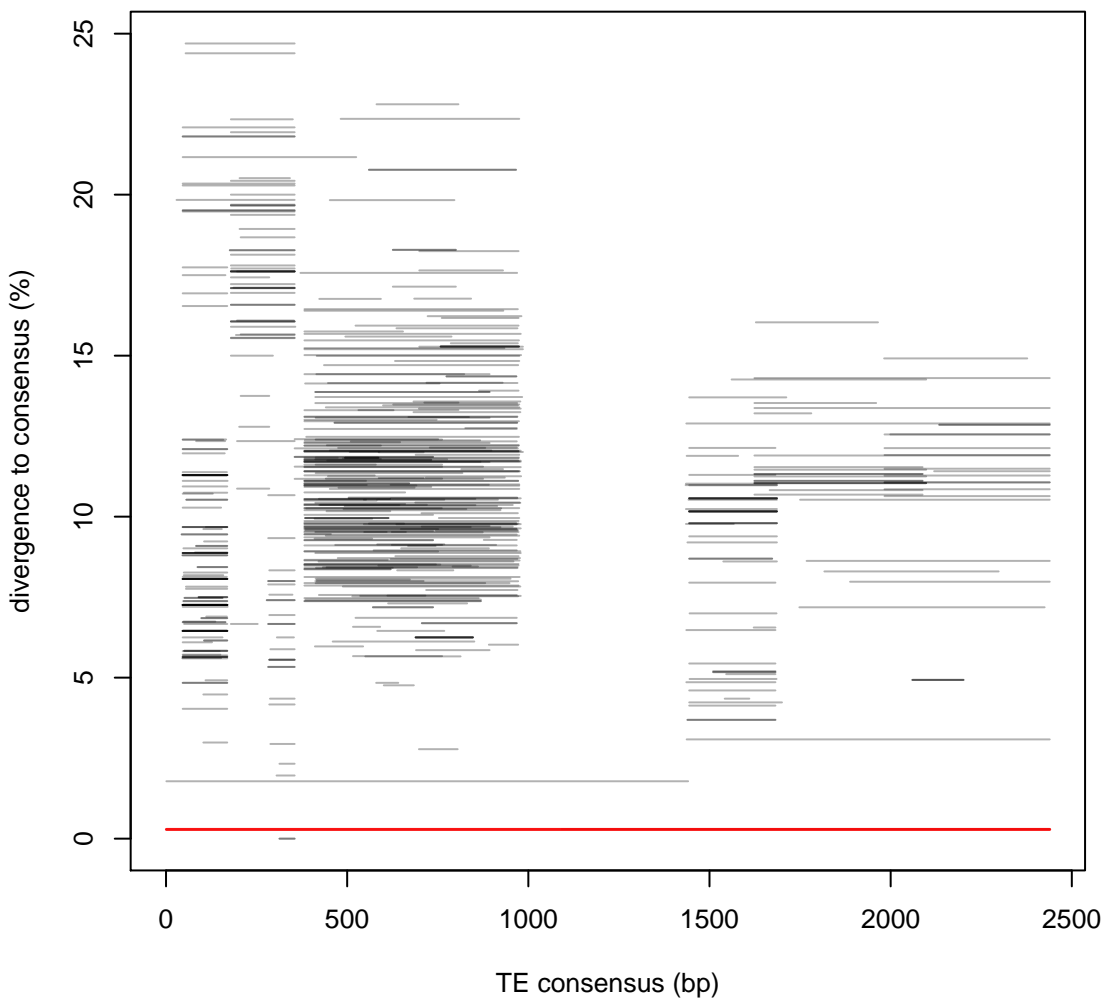
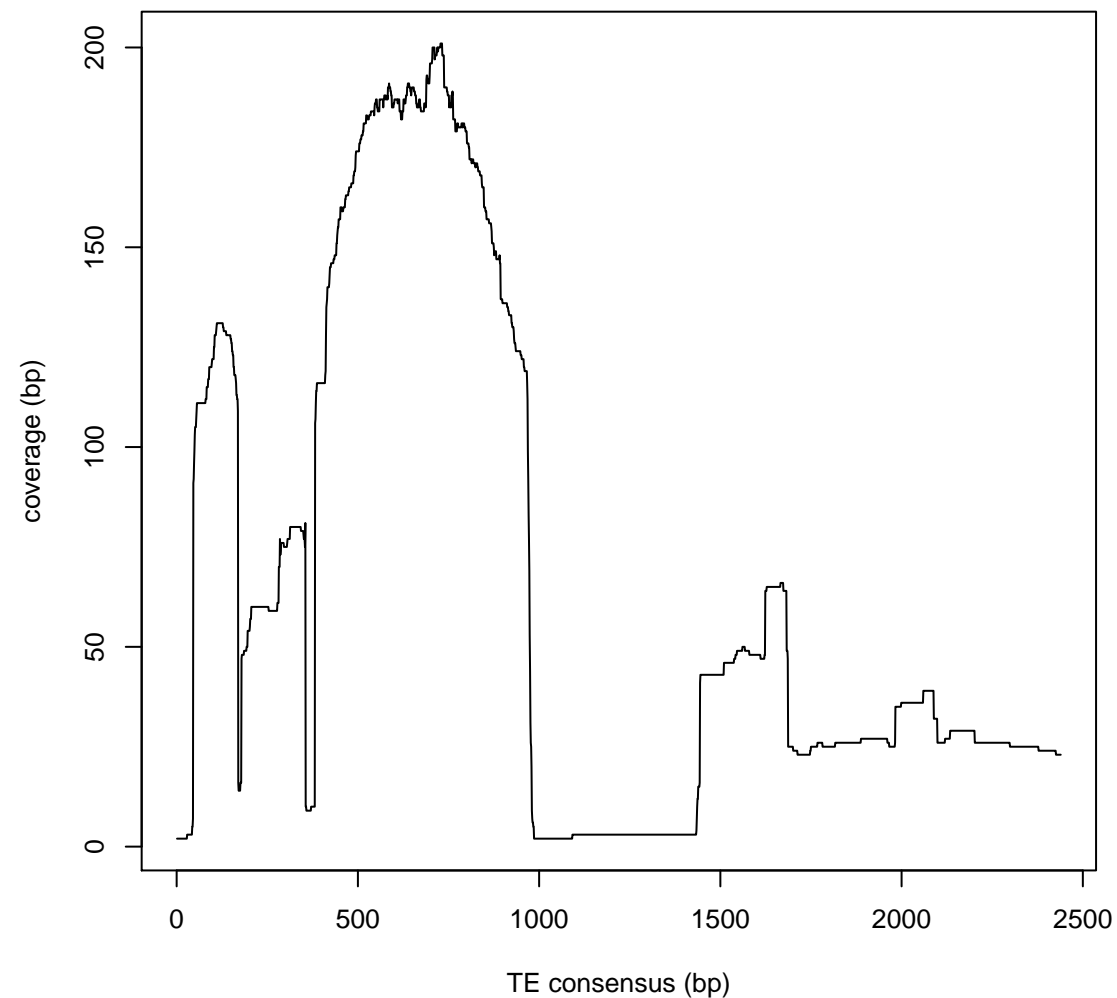


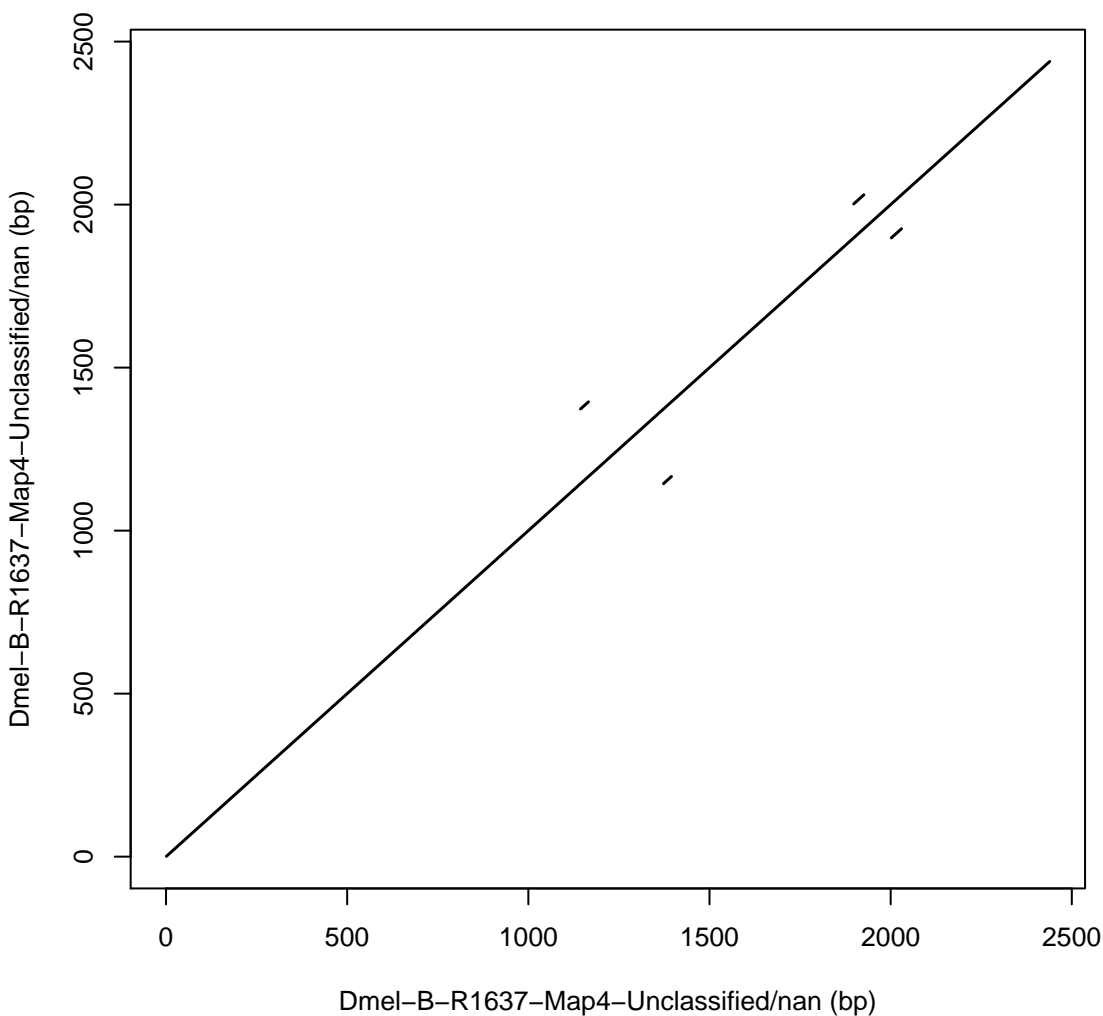
TE: Dmel-B-R1637-Map4-Unclassified/nan
consensus size: 2439bp; fragments: 559; full length: 1 (>=2195.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

