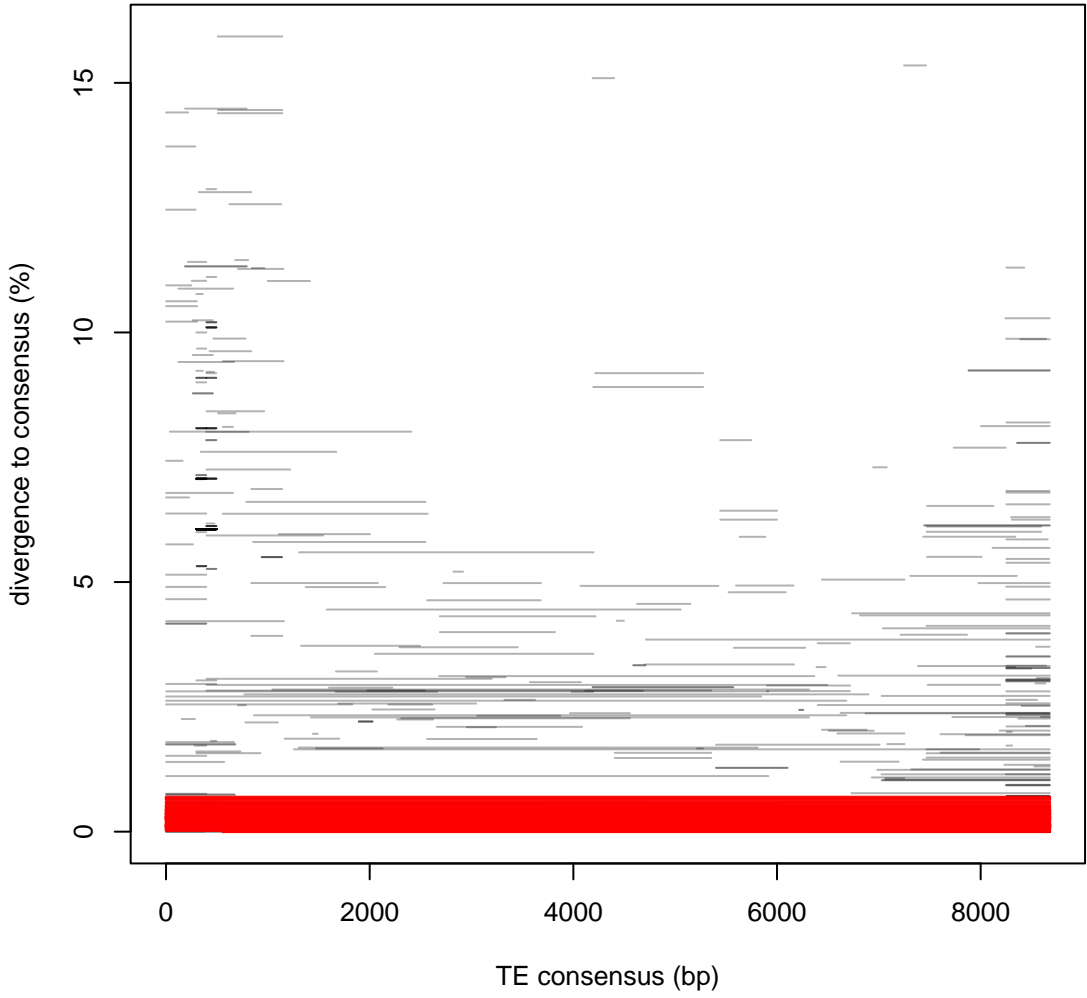
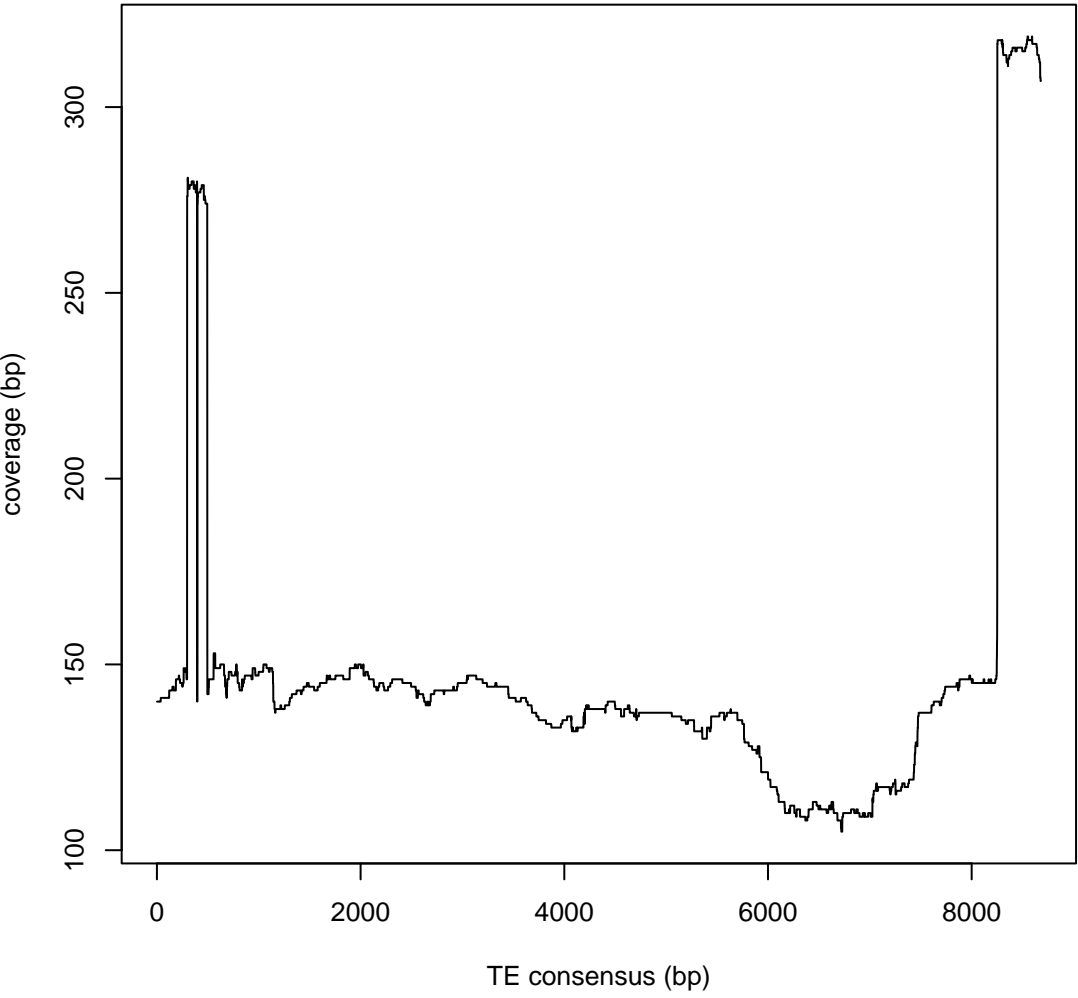


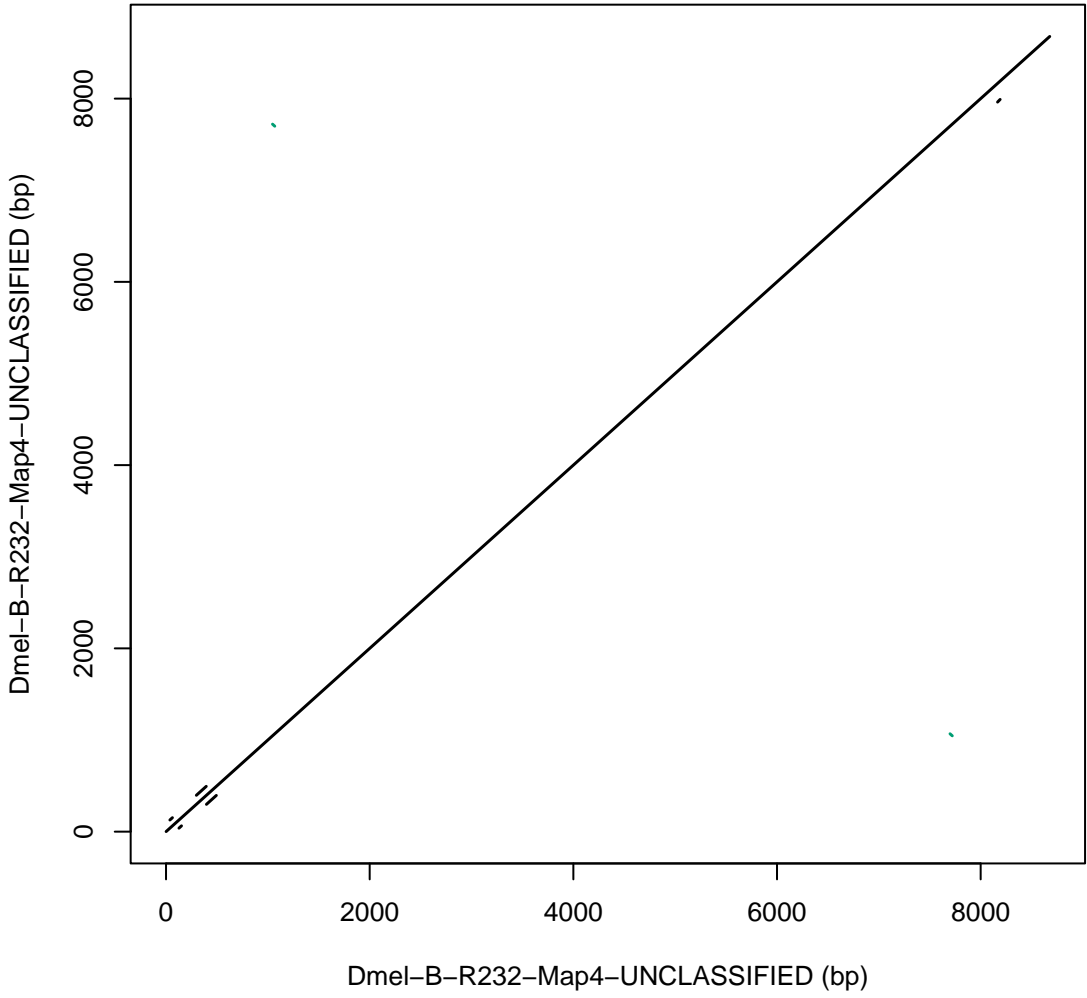
TE: Dmel-B-R232-Map4-UNCLASSIFIED
consensus size: 8678bp; fragments: 827; full length: 72 (>=7810.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

