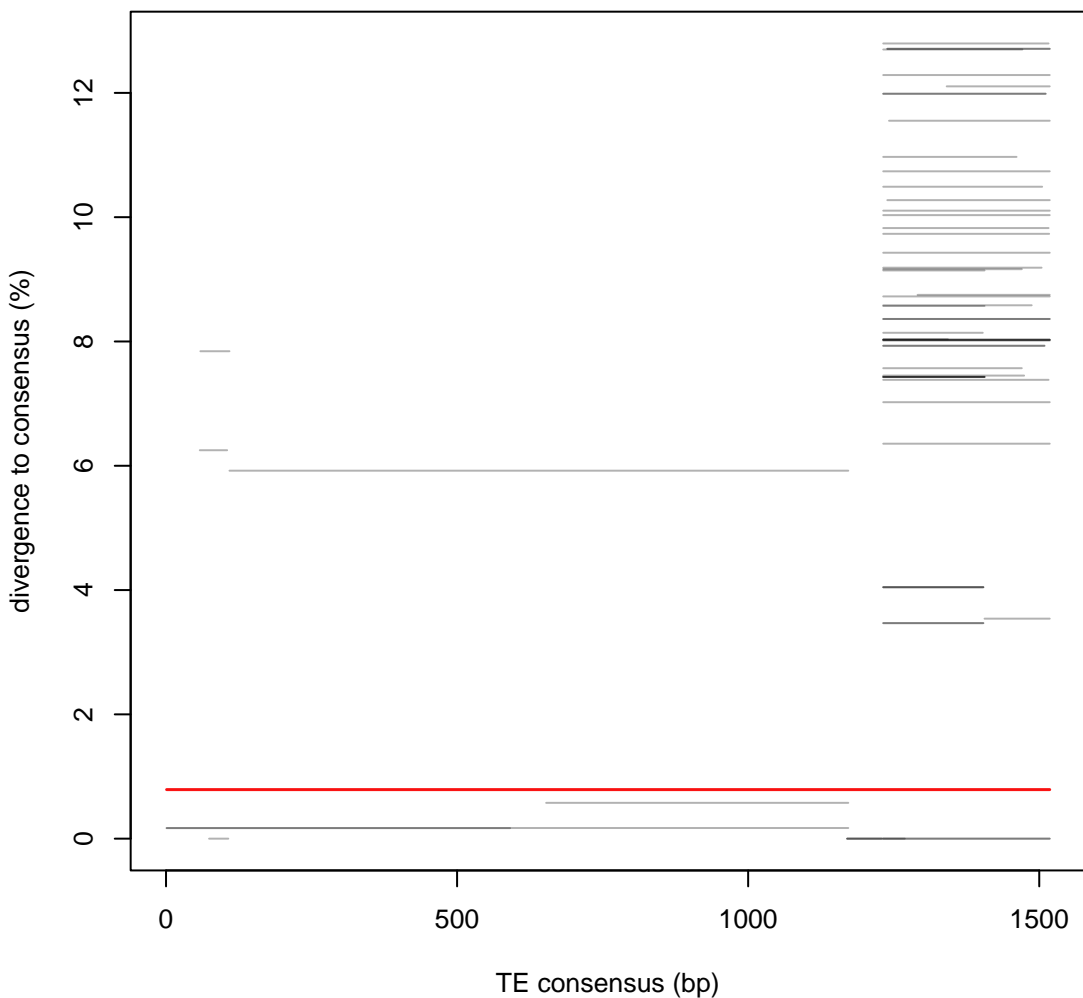
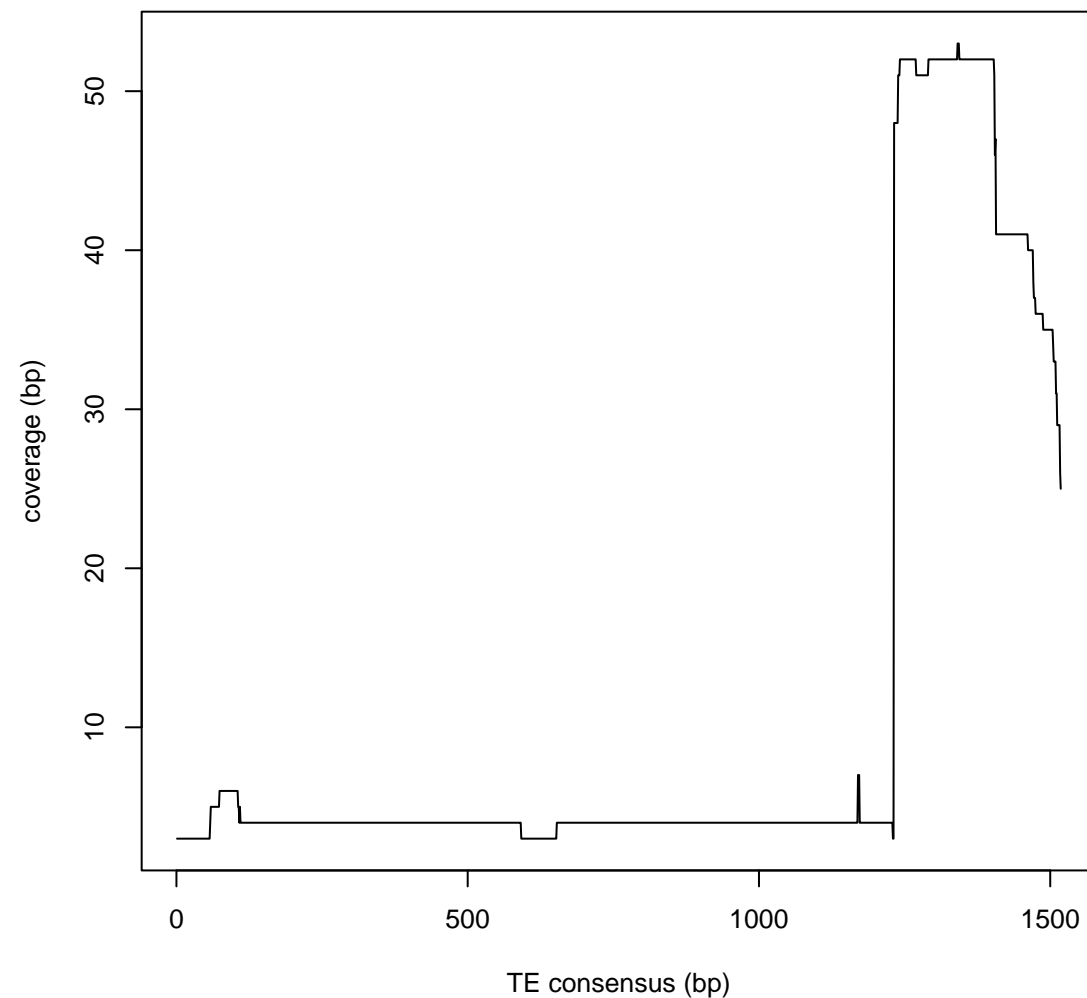


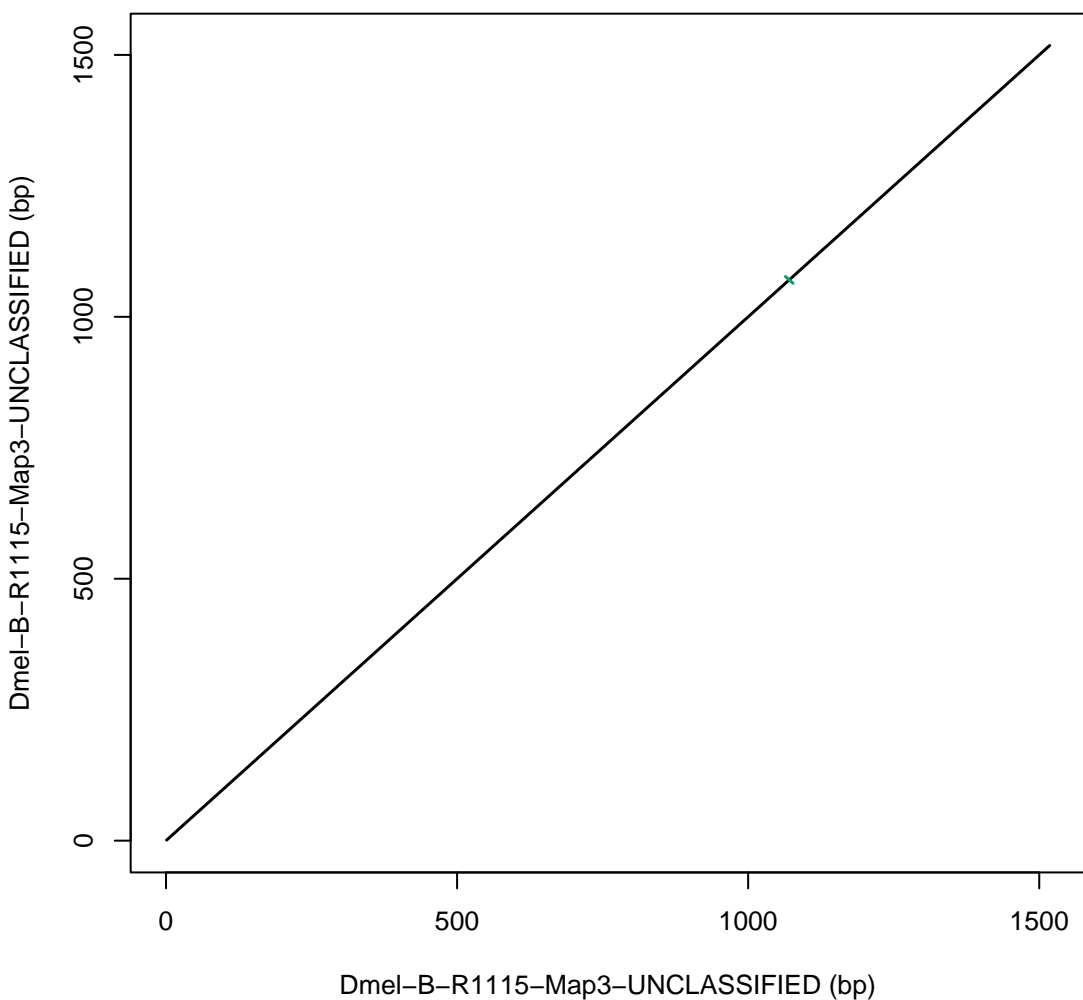
TE: Dmel-B-R1115-Map3-UNCLASSIFIED
consensus size: 1518bp; fragments: 63; full length: 1 (>=1366.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

