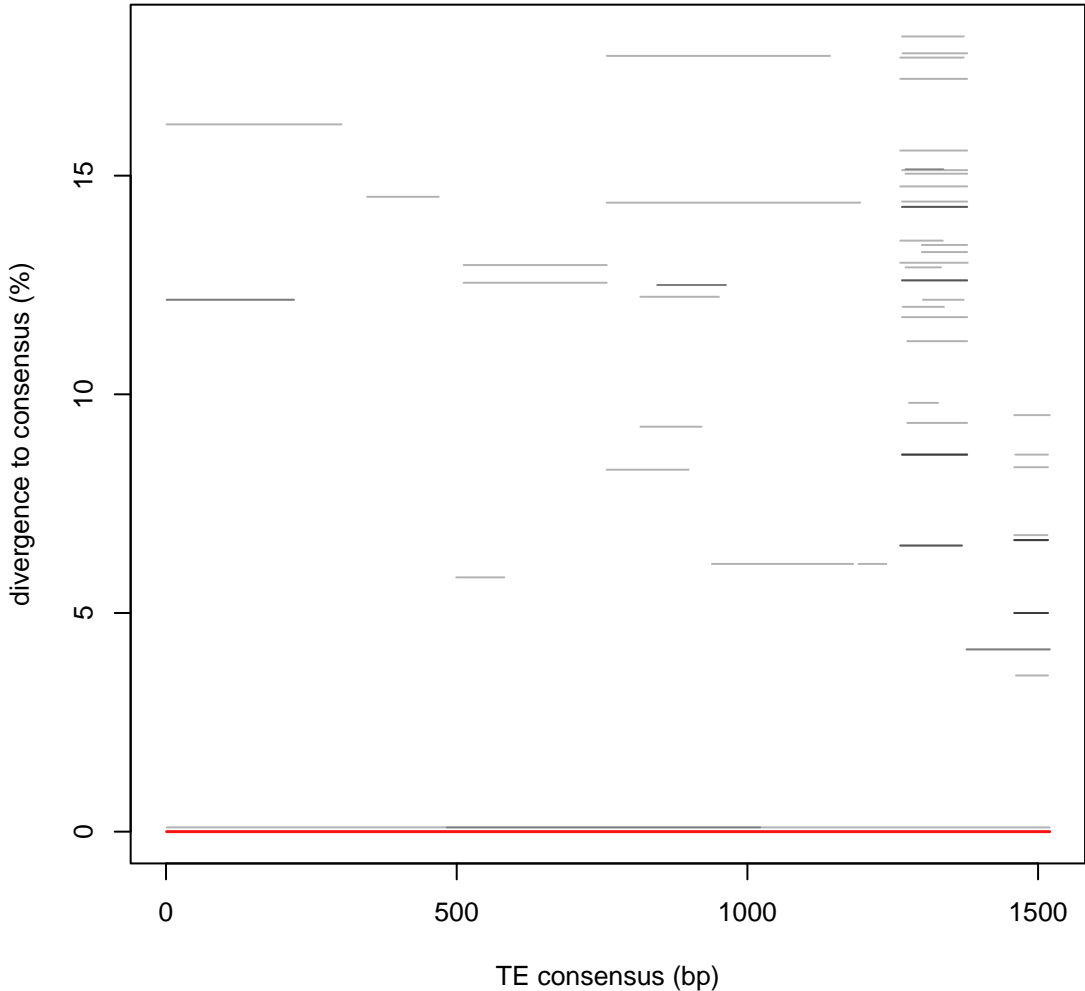
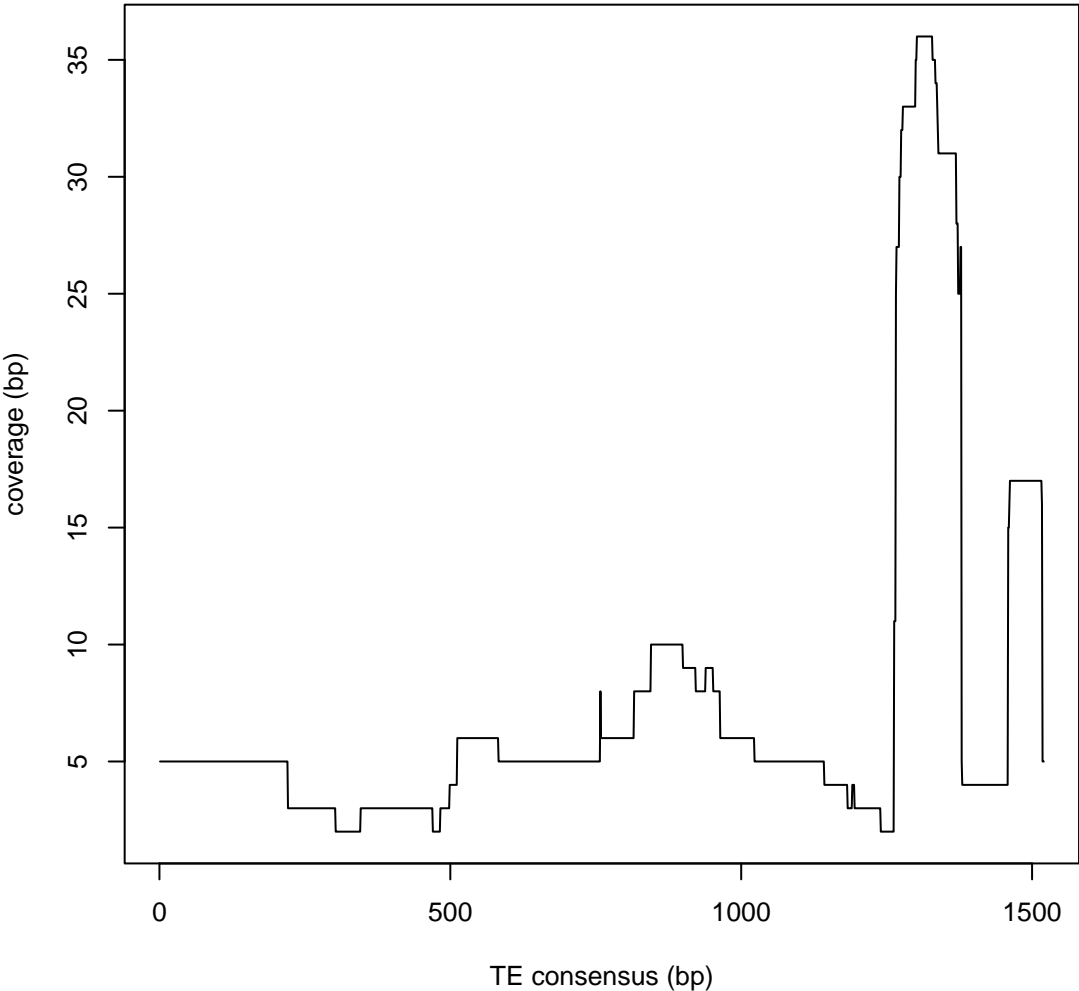


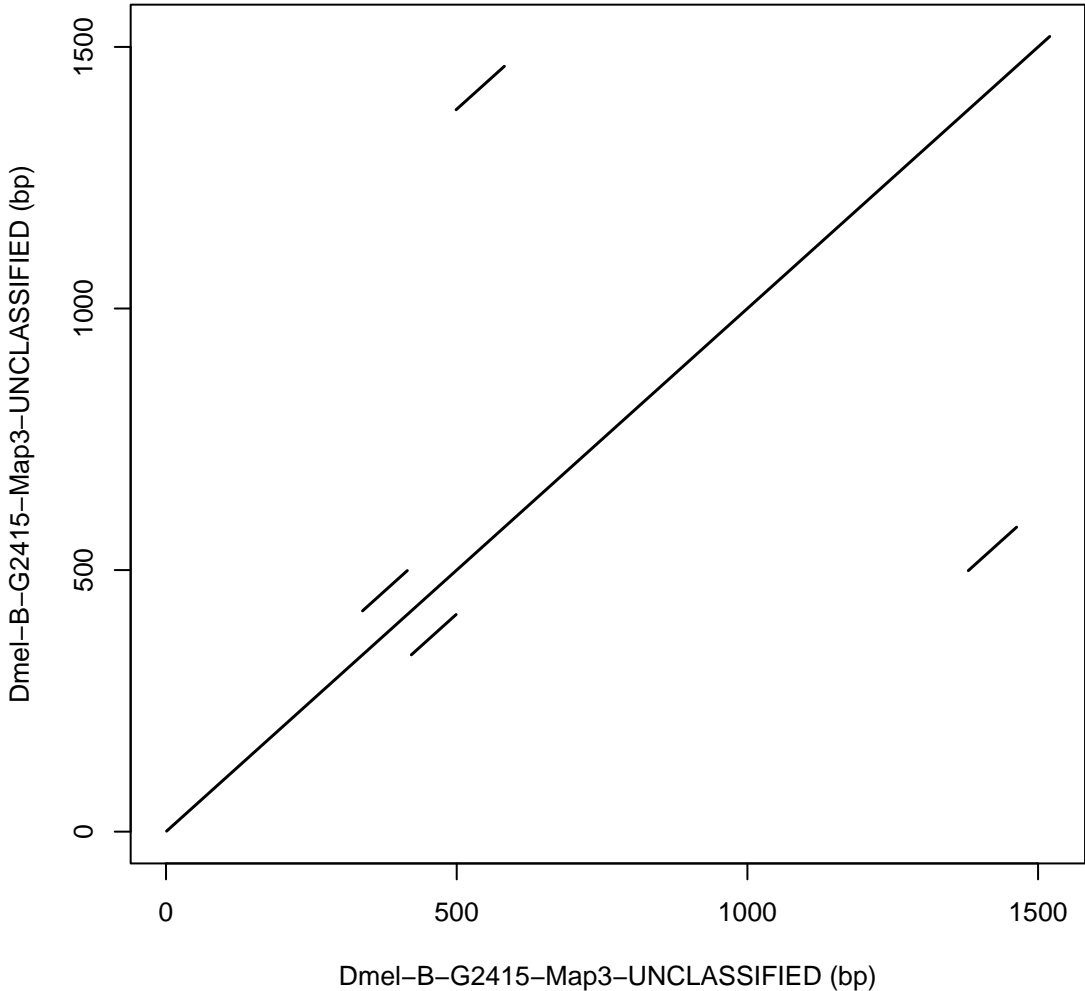
TE: Dmel-B-G2415-Map3-UNCLASSIFIED
consensus size: 1520bp; fragments: 68; full length: 1 (>=1368bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

