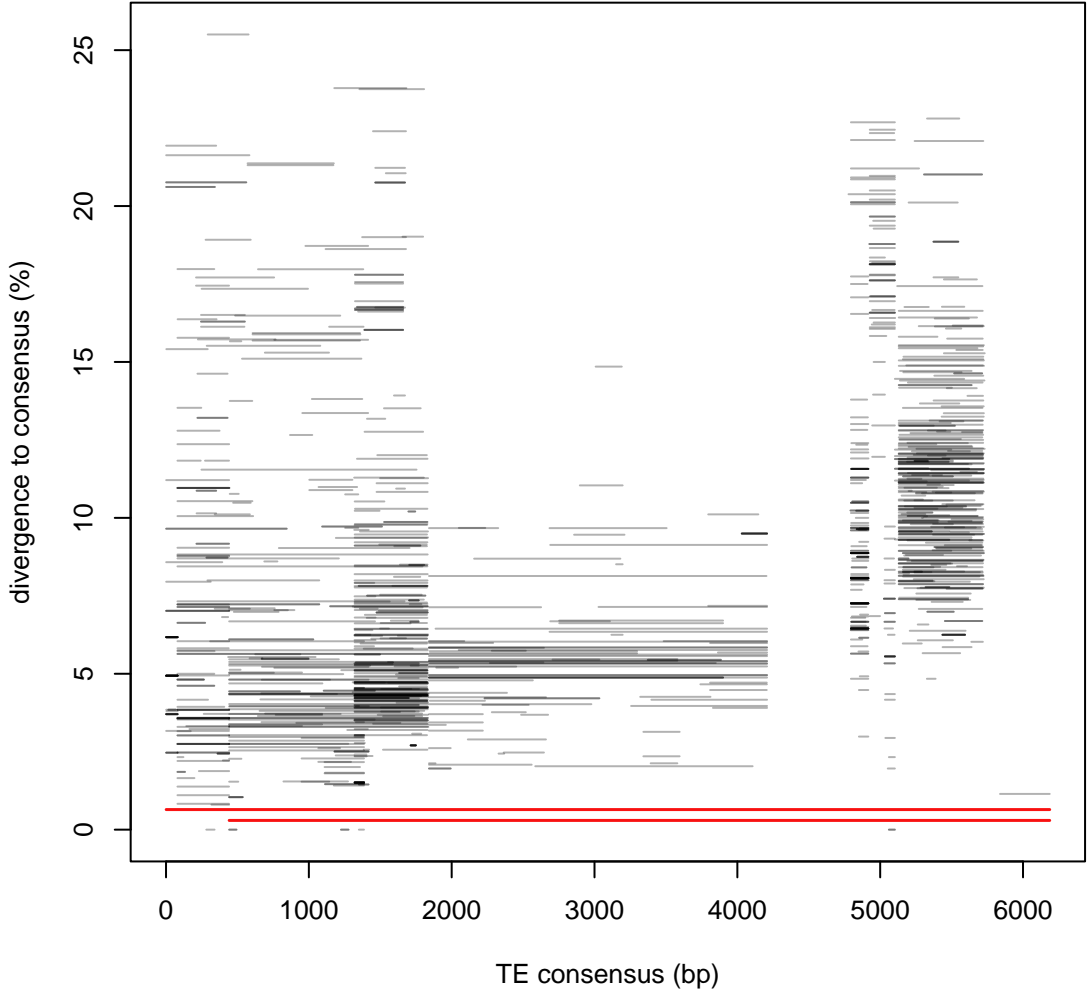
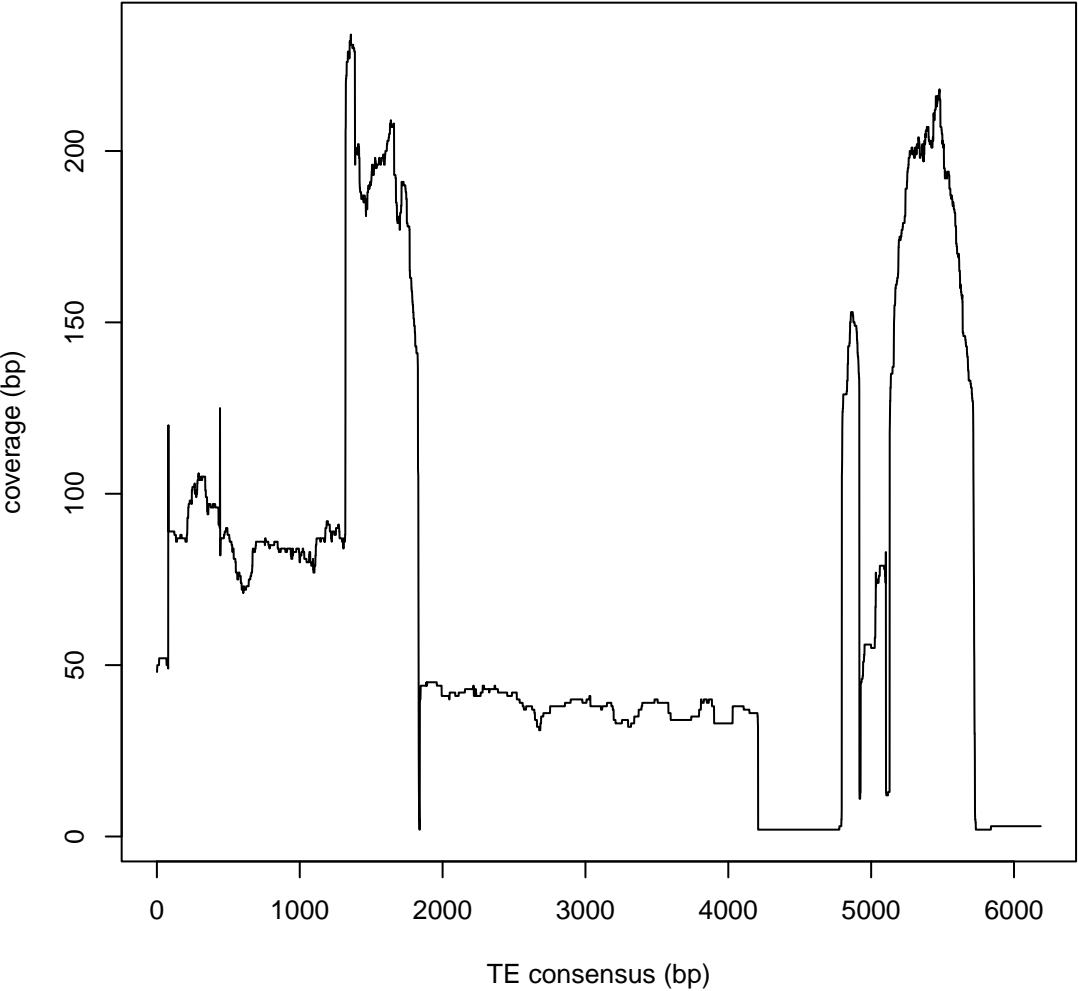


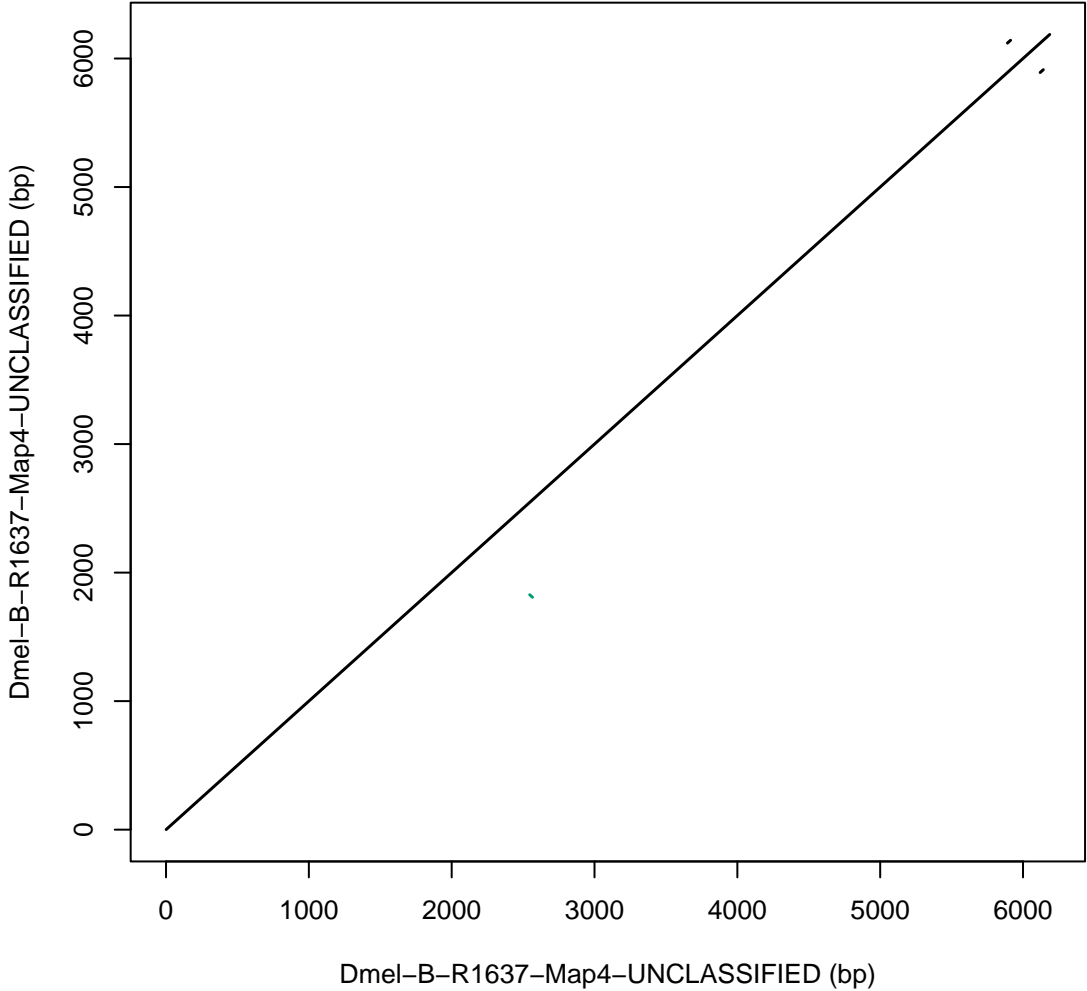
TE: Dmel-B-R1637-Map4-UNCLASSIFIED
consensus size: 6187bp; fragments: 1126; full length: 2 (>=5568.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

