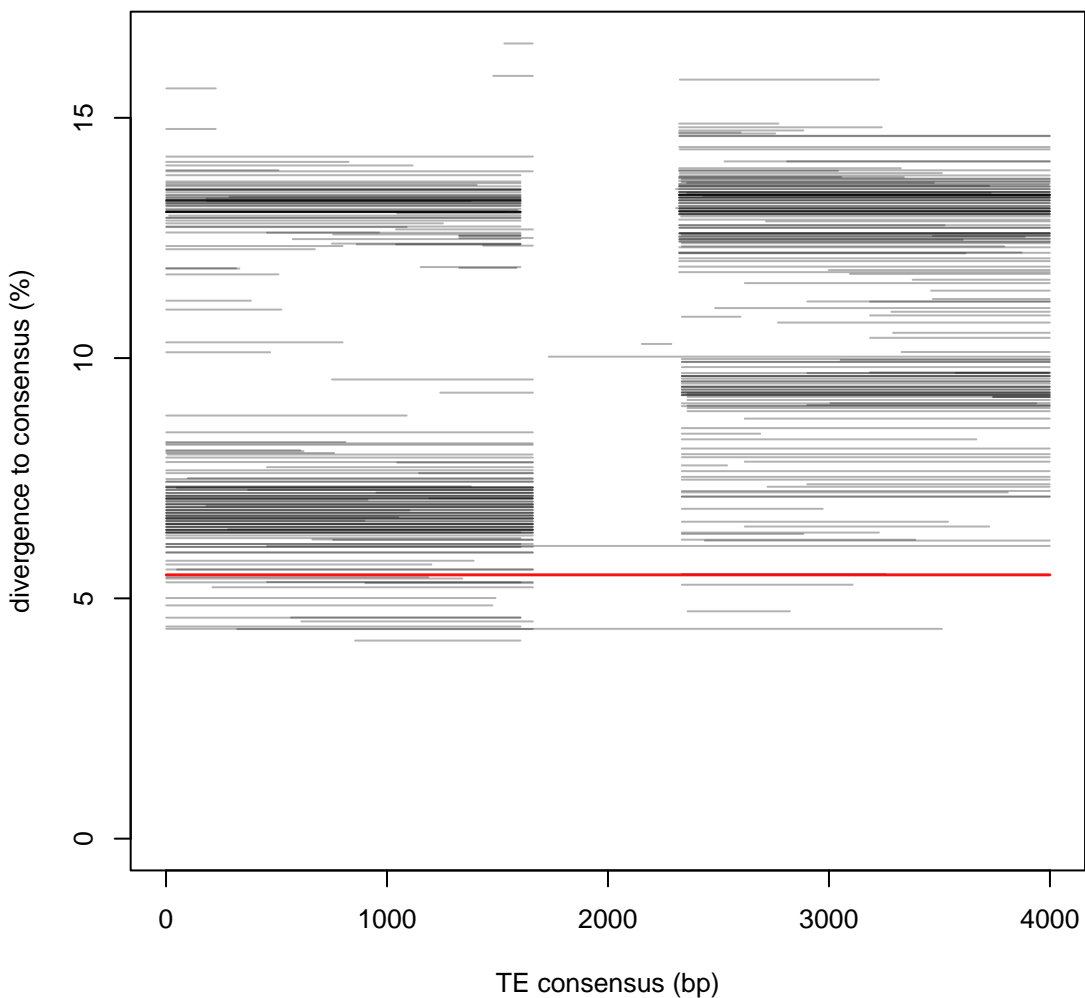
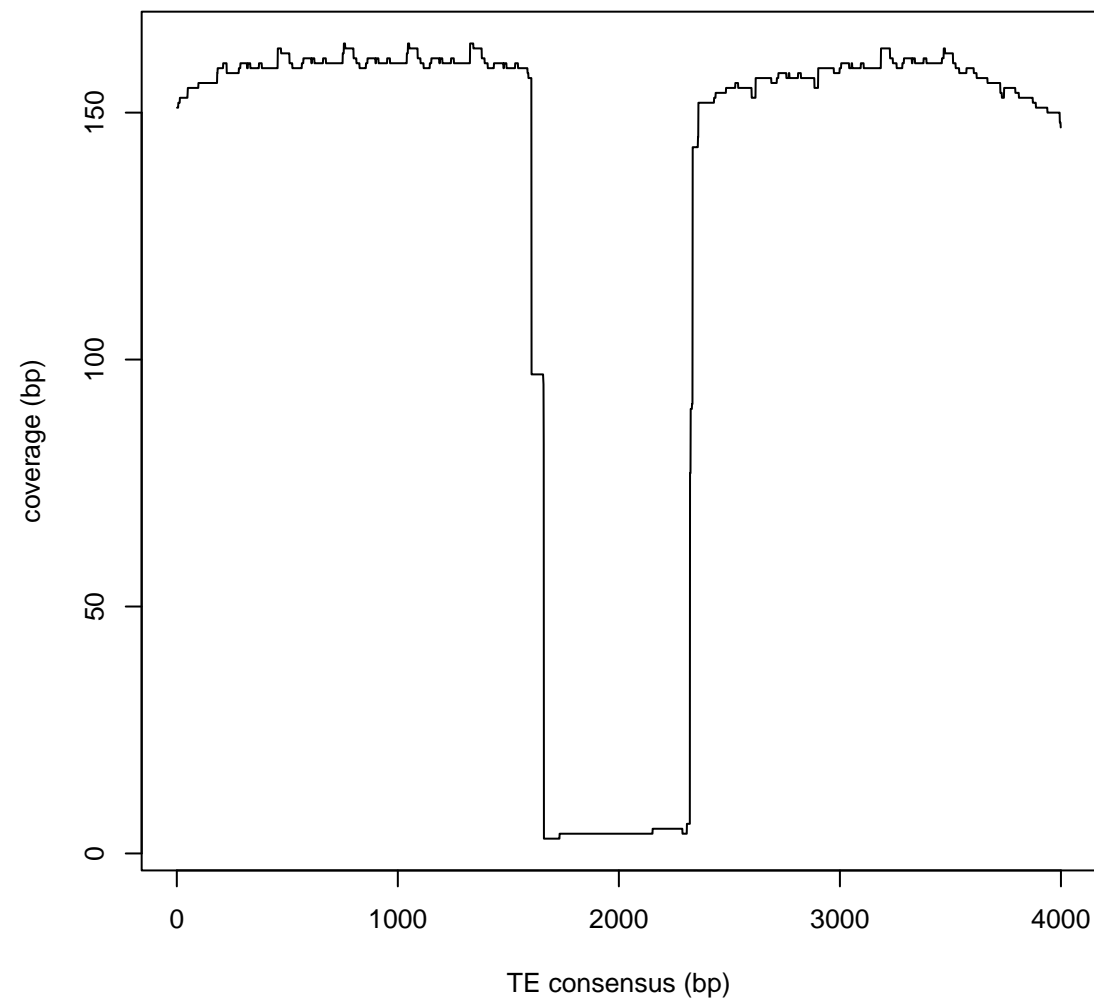


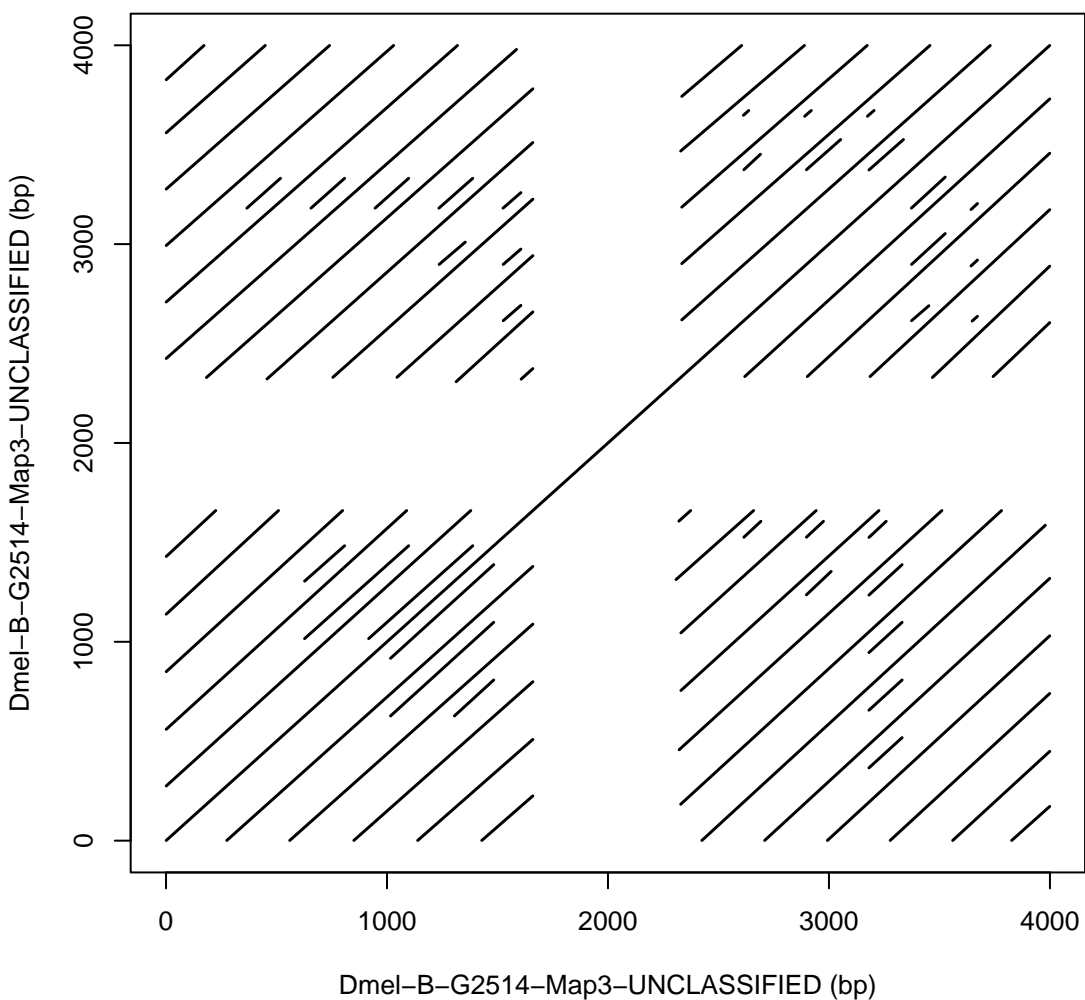
TE: Dmel-B-G2514-Map3-UNCLASSIFIED
consensus size: 3999bp; fragments: 379; full length: 1 (>=3599.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

