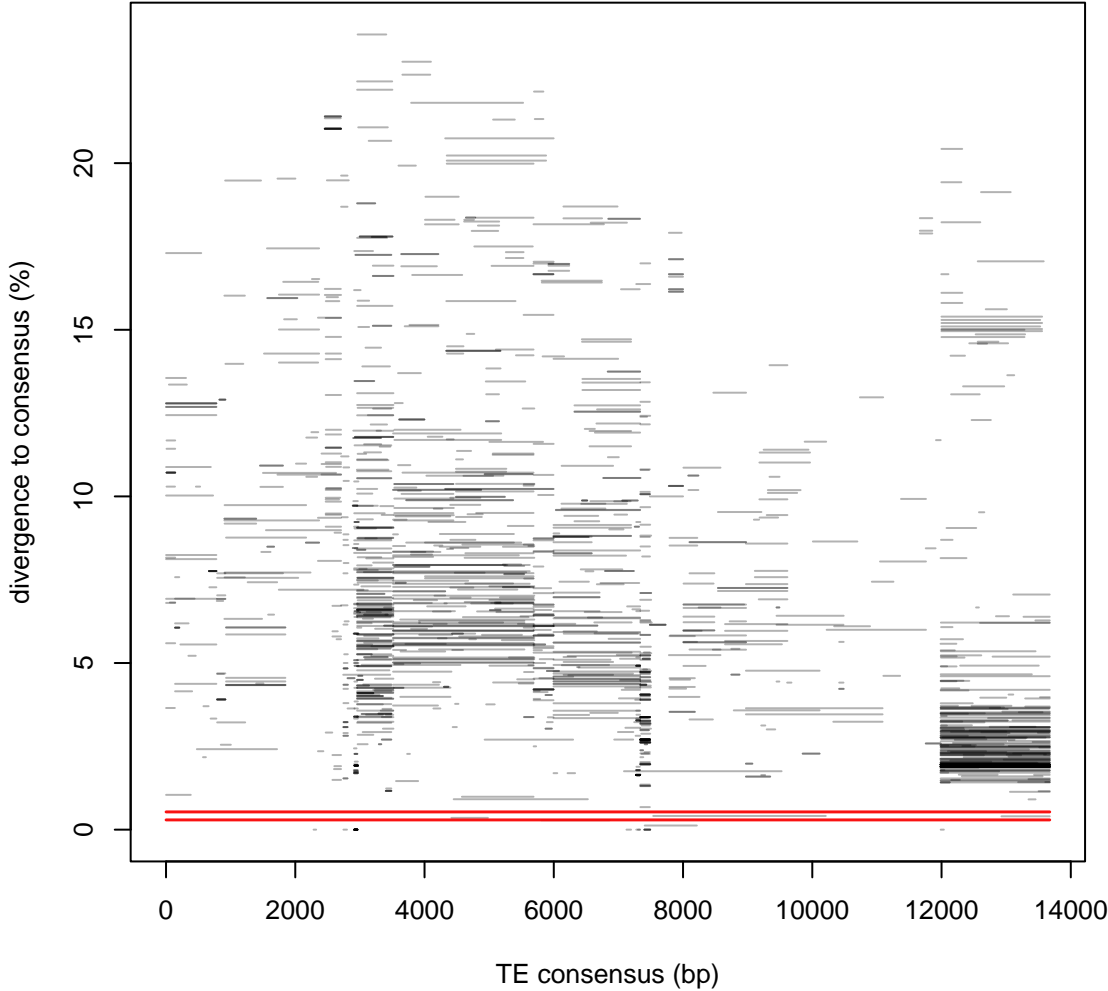
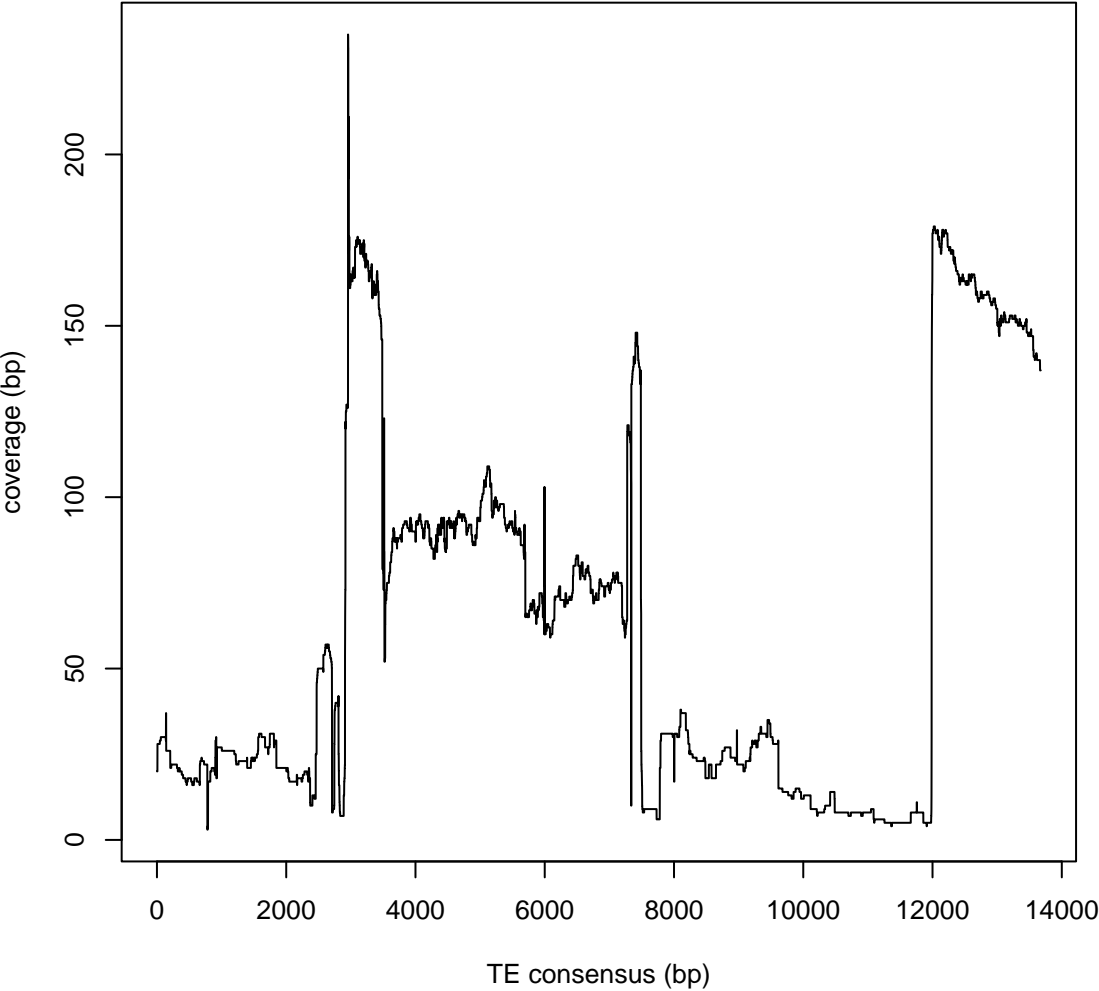


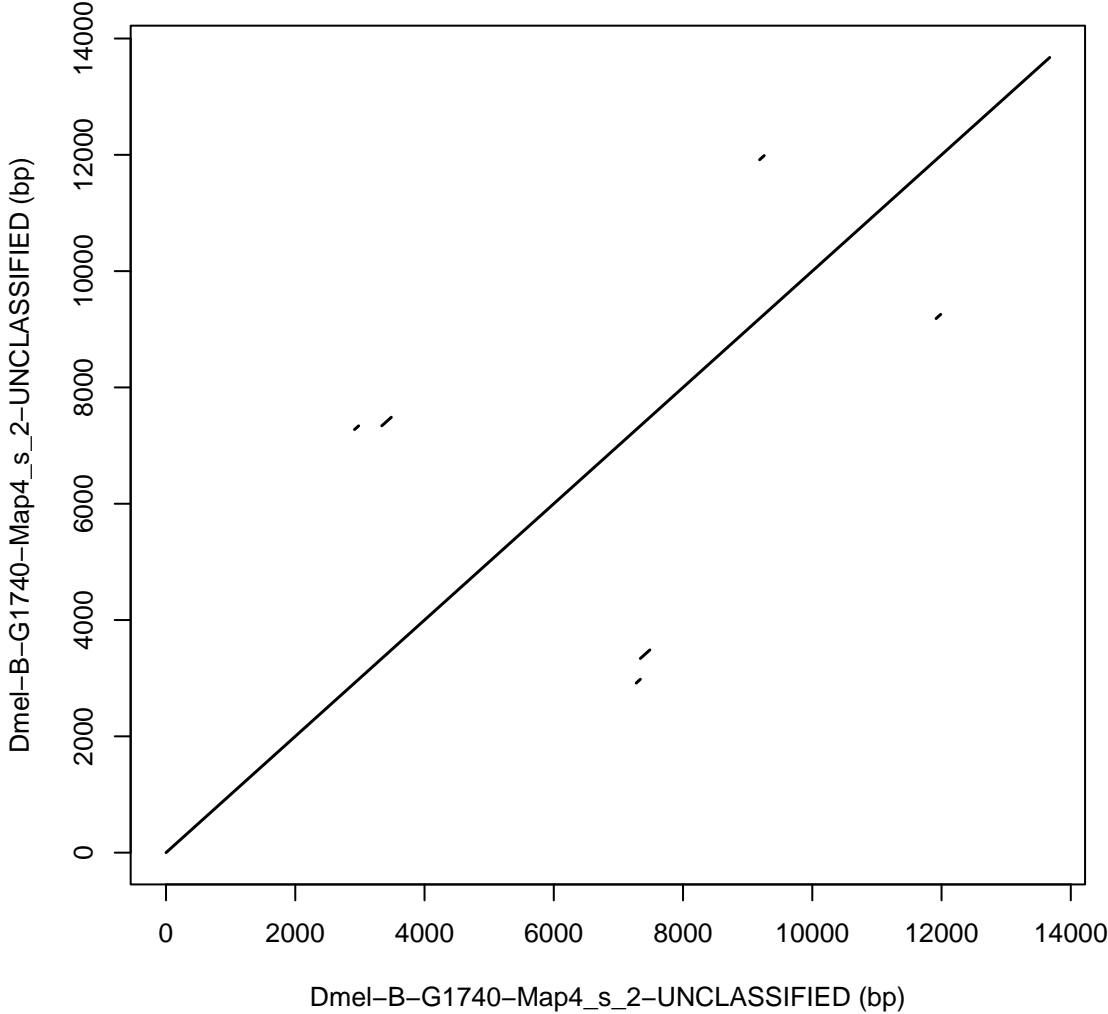
TE: Dmel-B-G1740-Map4_s_2-UNCLASSIFIED
consensus size: 13674bp; fragments: 1573; full length: 2 (>=12306.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

