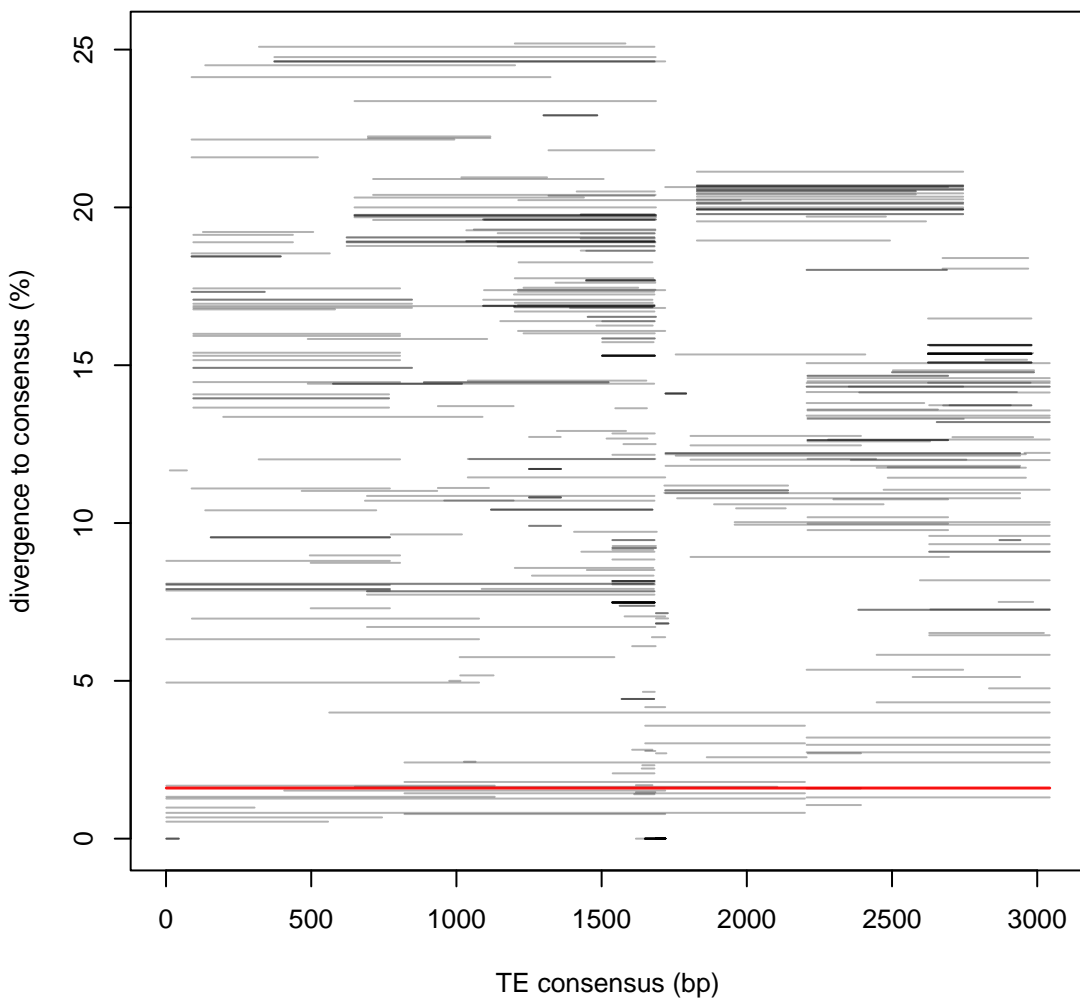
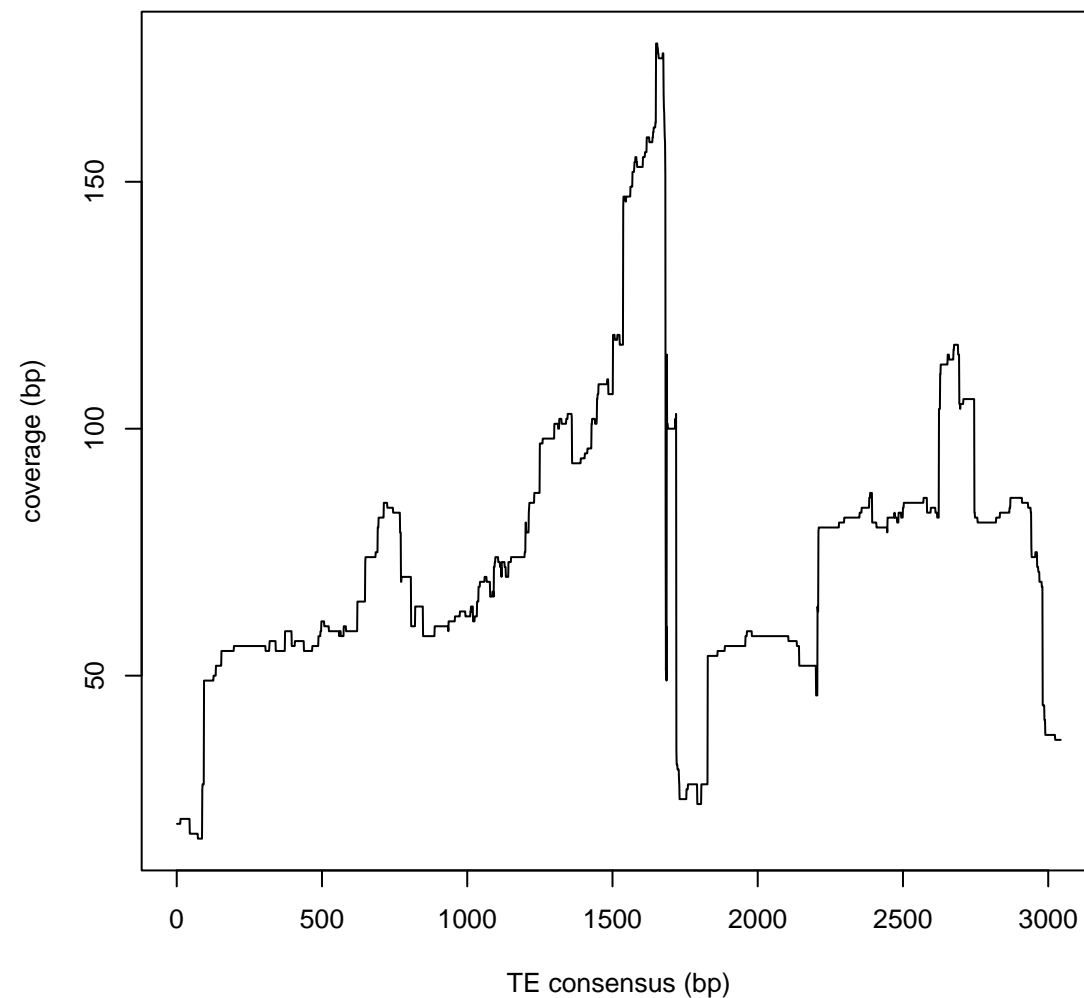


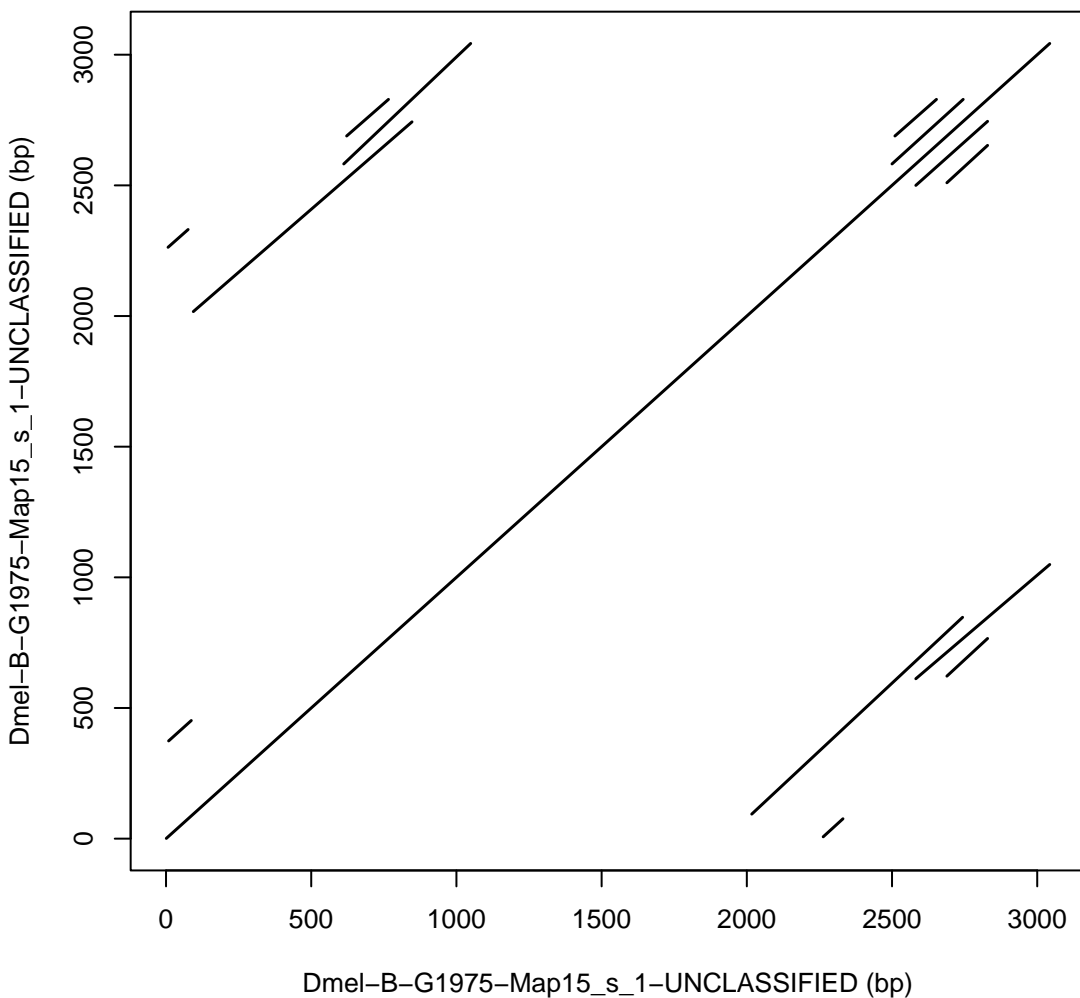
TE: Dmel-B-G1975-Map15_s_1-UNCLASSIFIED
consensus size: 3043bp; fragments: 502; full length: 1 (>=2738.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

