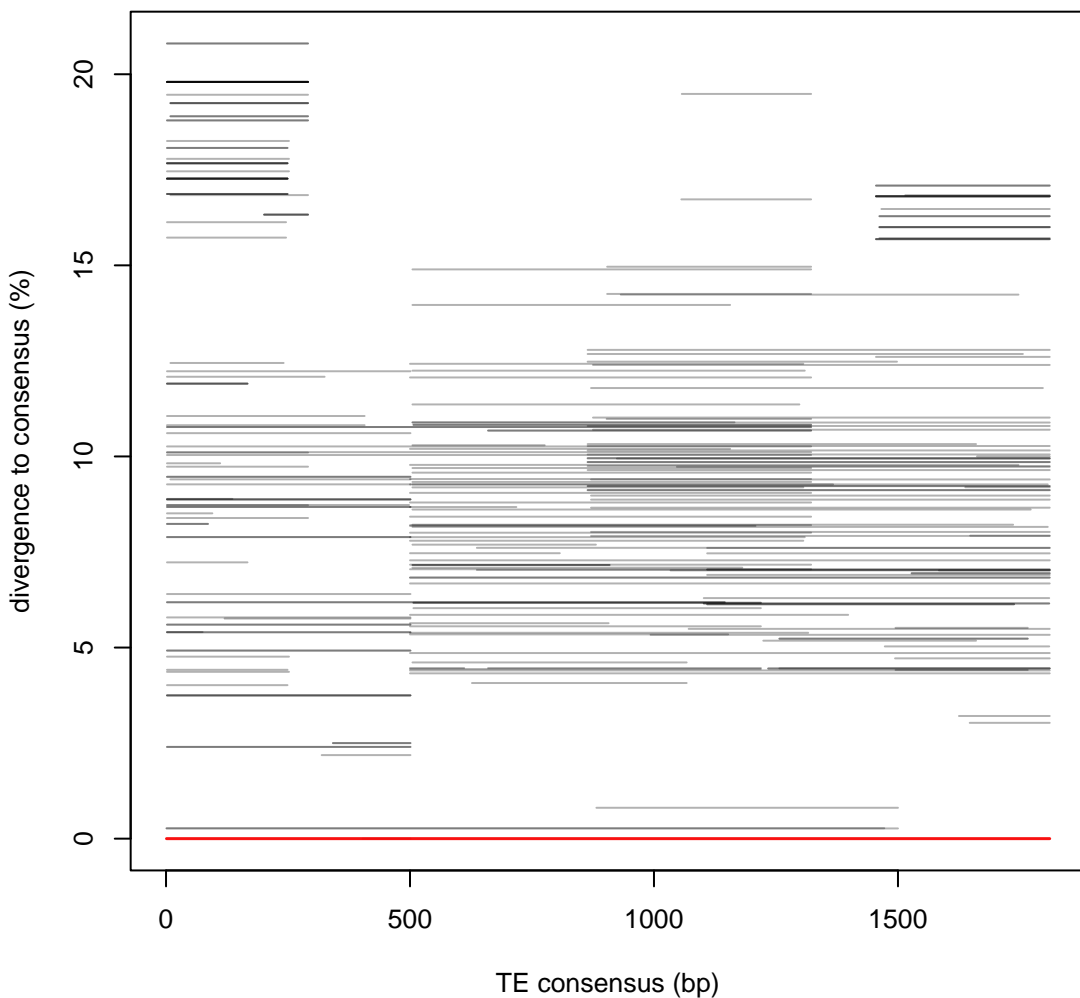
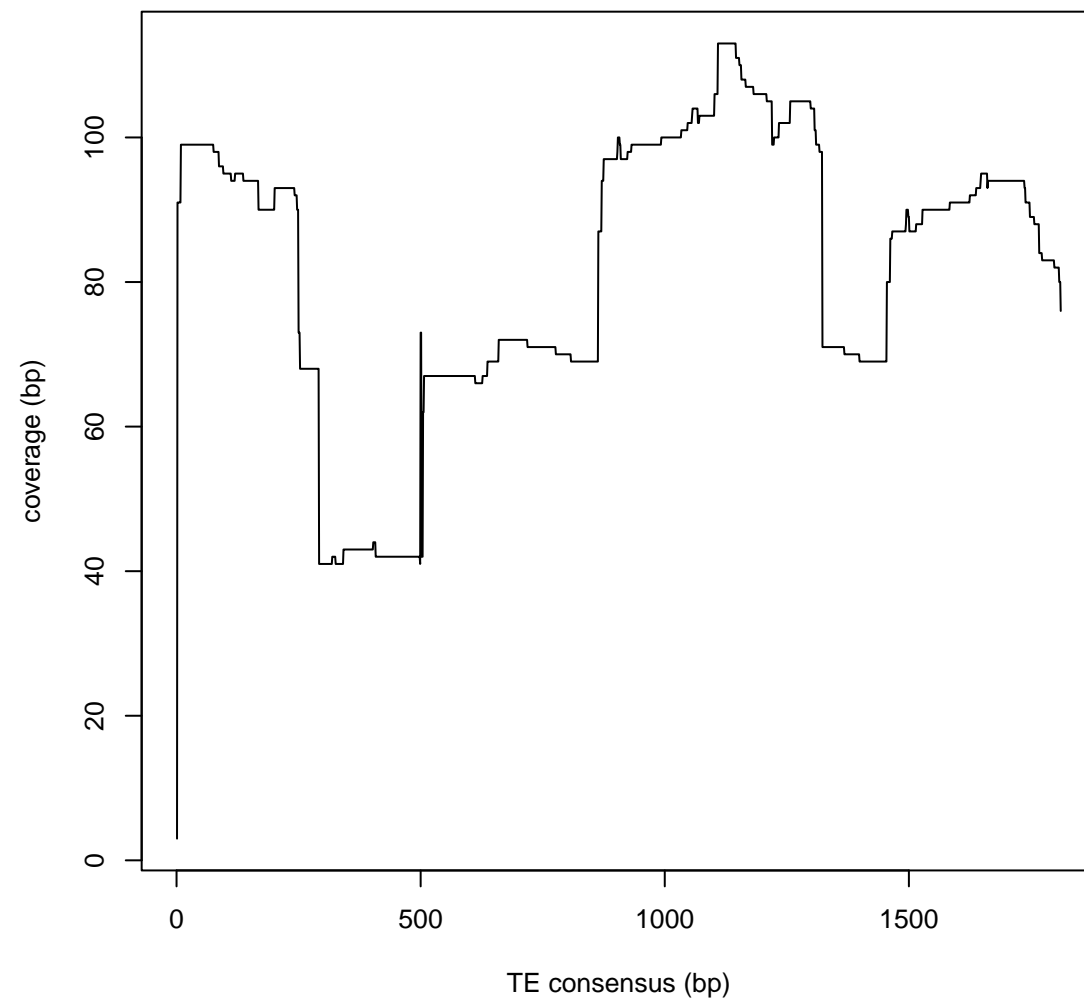


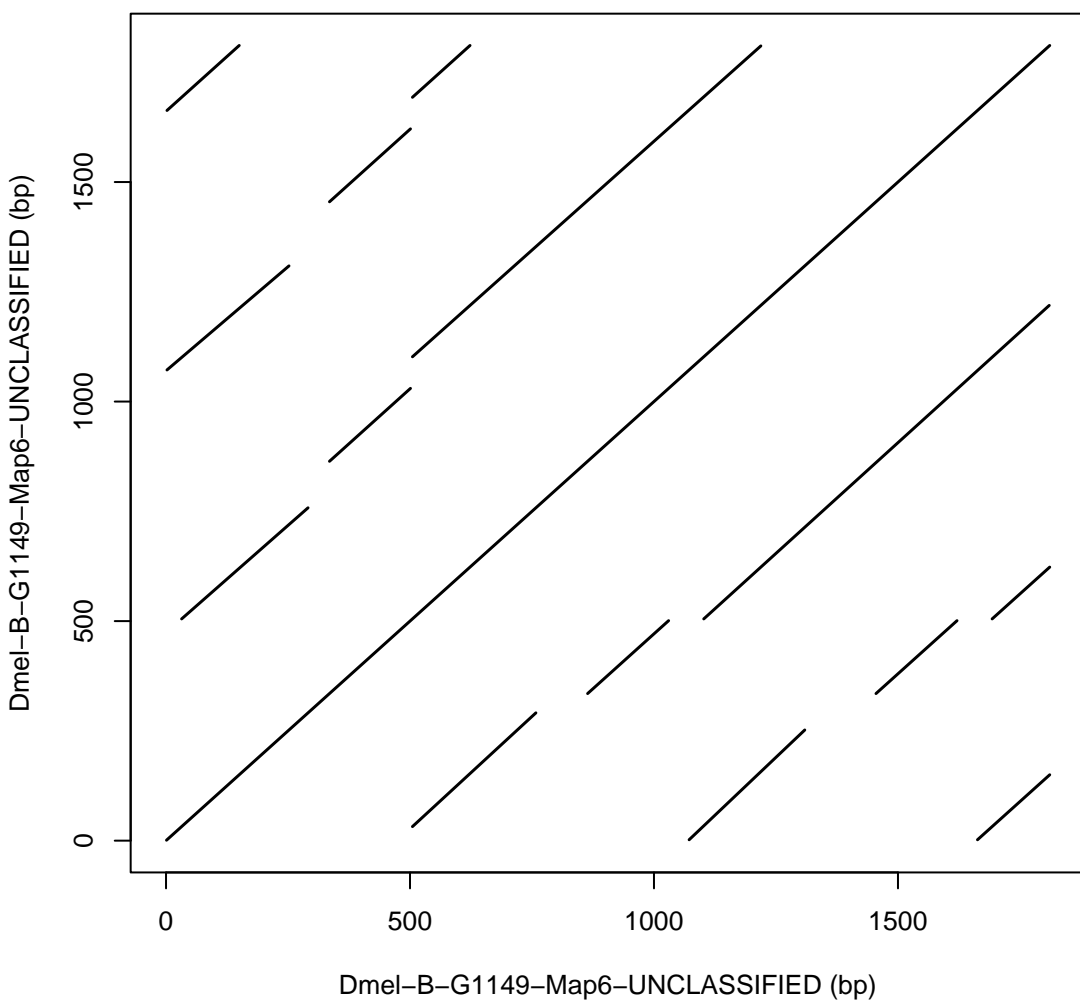
TE: Dmel-B-G1149-Map6-UNCLASSIFIED
consensus size: 1811bp; fragments: 257; full length: 1 (>=1629.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

