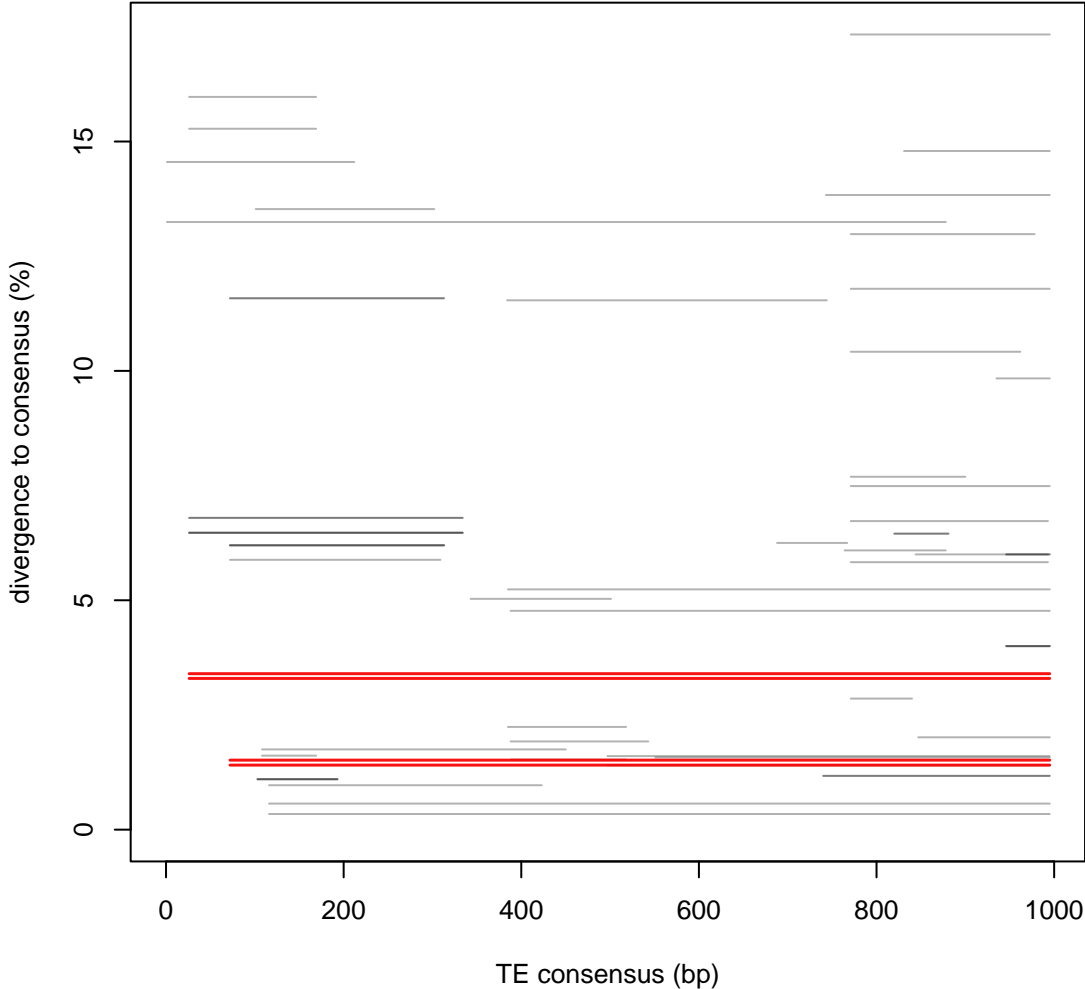
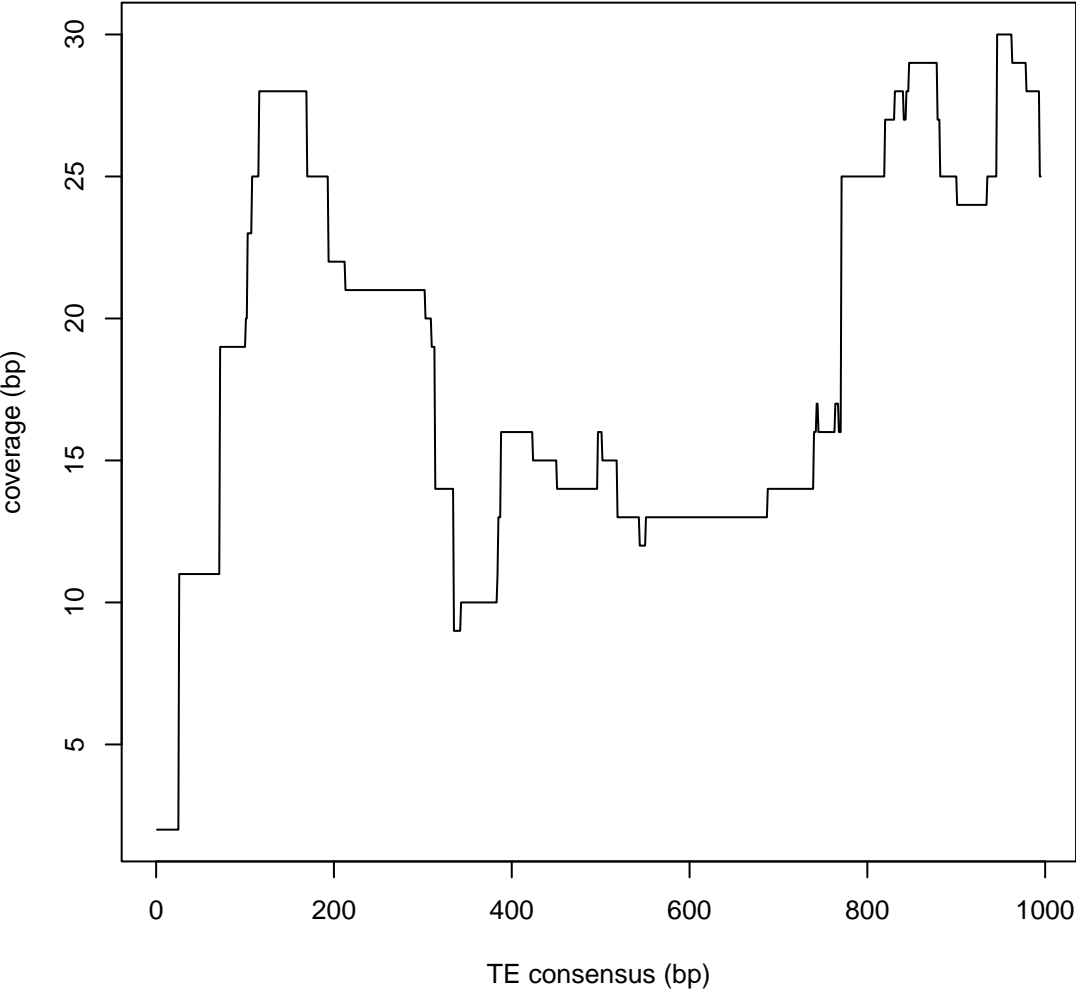


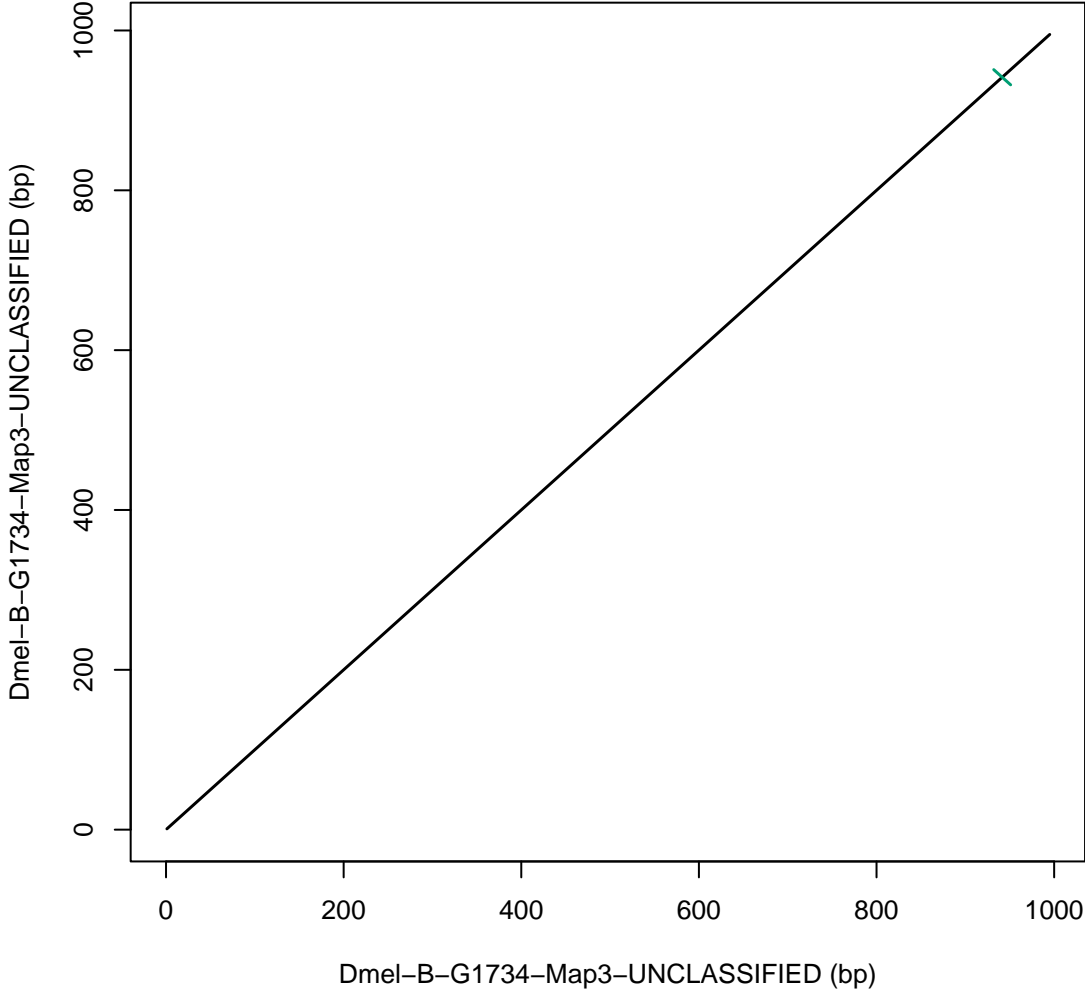
TE: Dmel-B-G1734-Map3-UNCLASSIFIED
consensus size: 995bp; fragments: 63; full length: 4 (>=895.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

