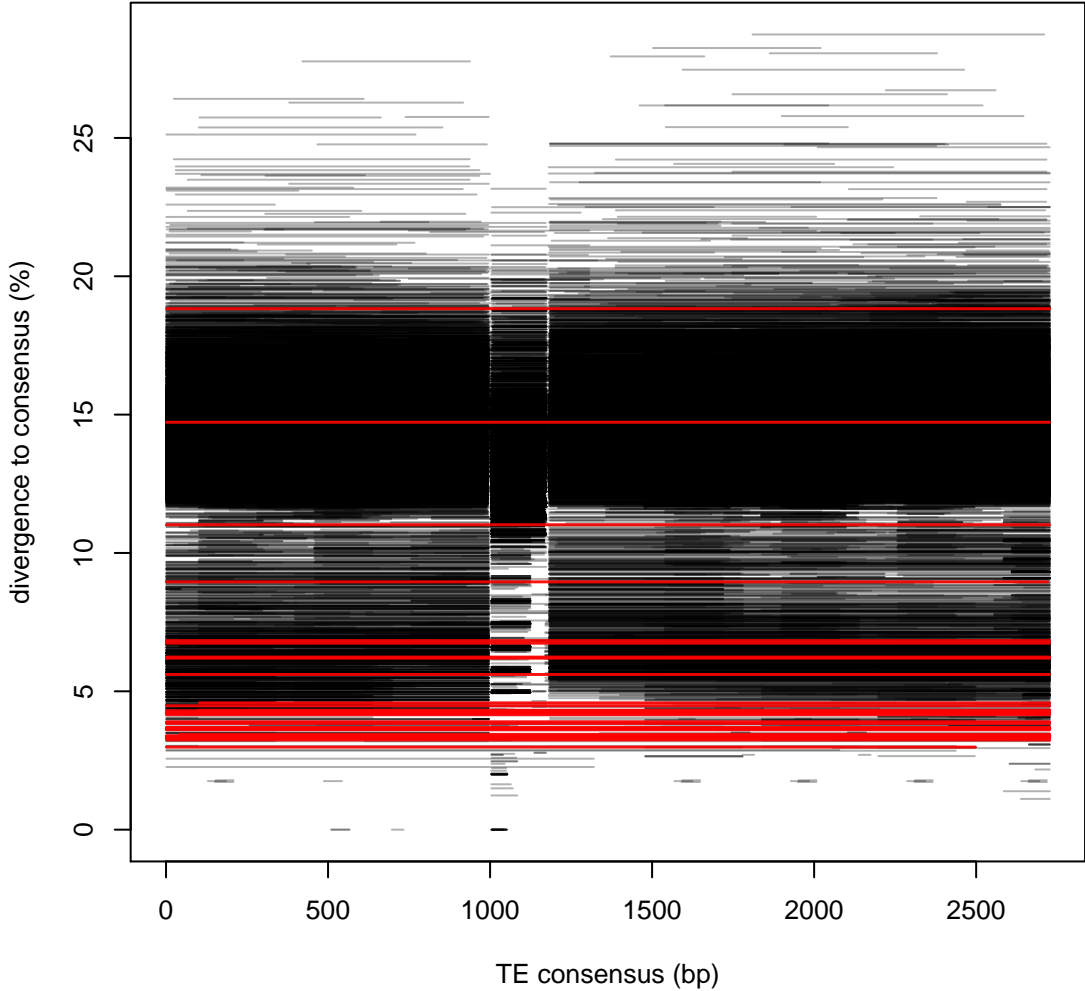
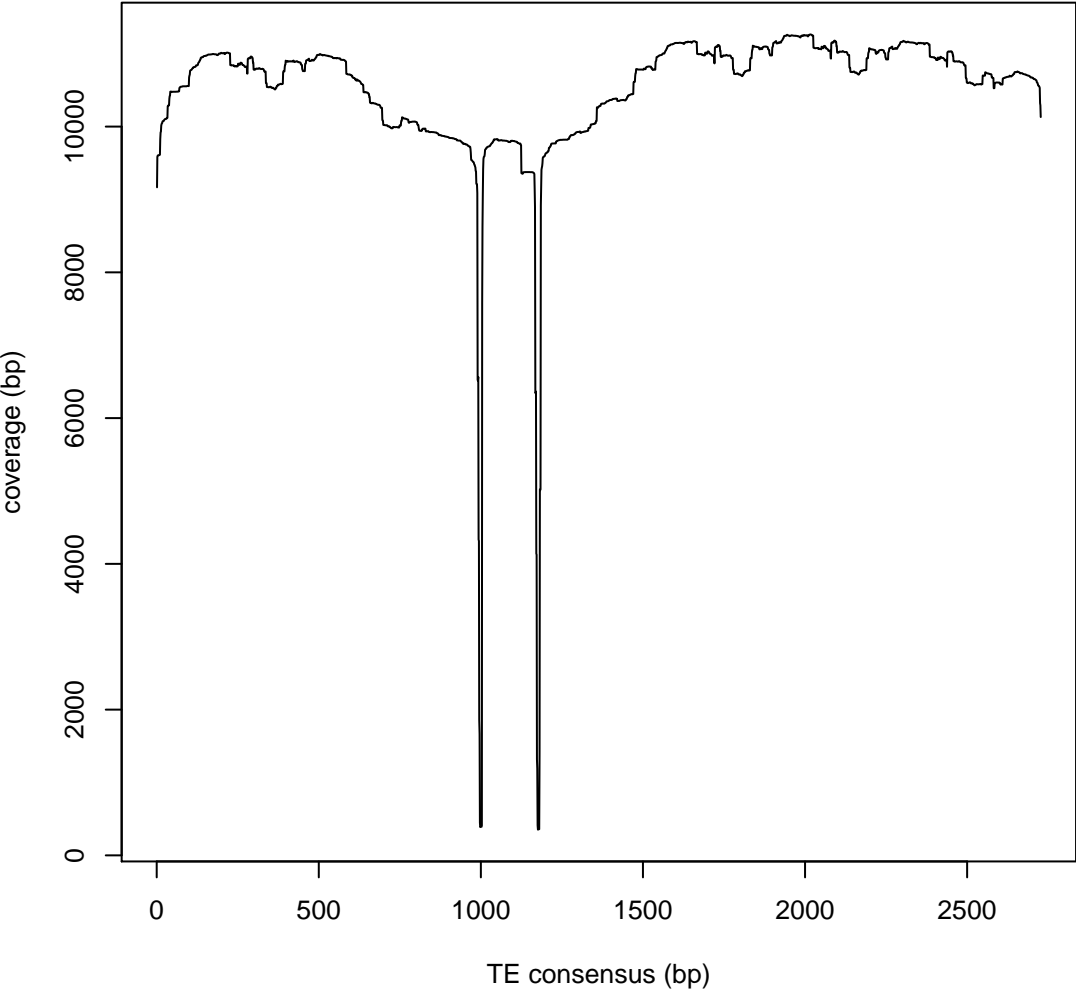


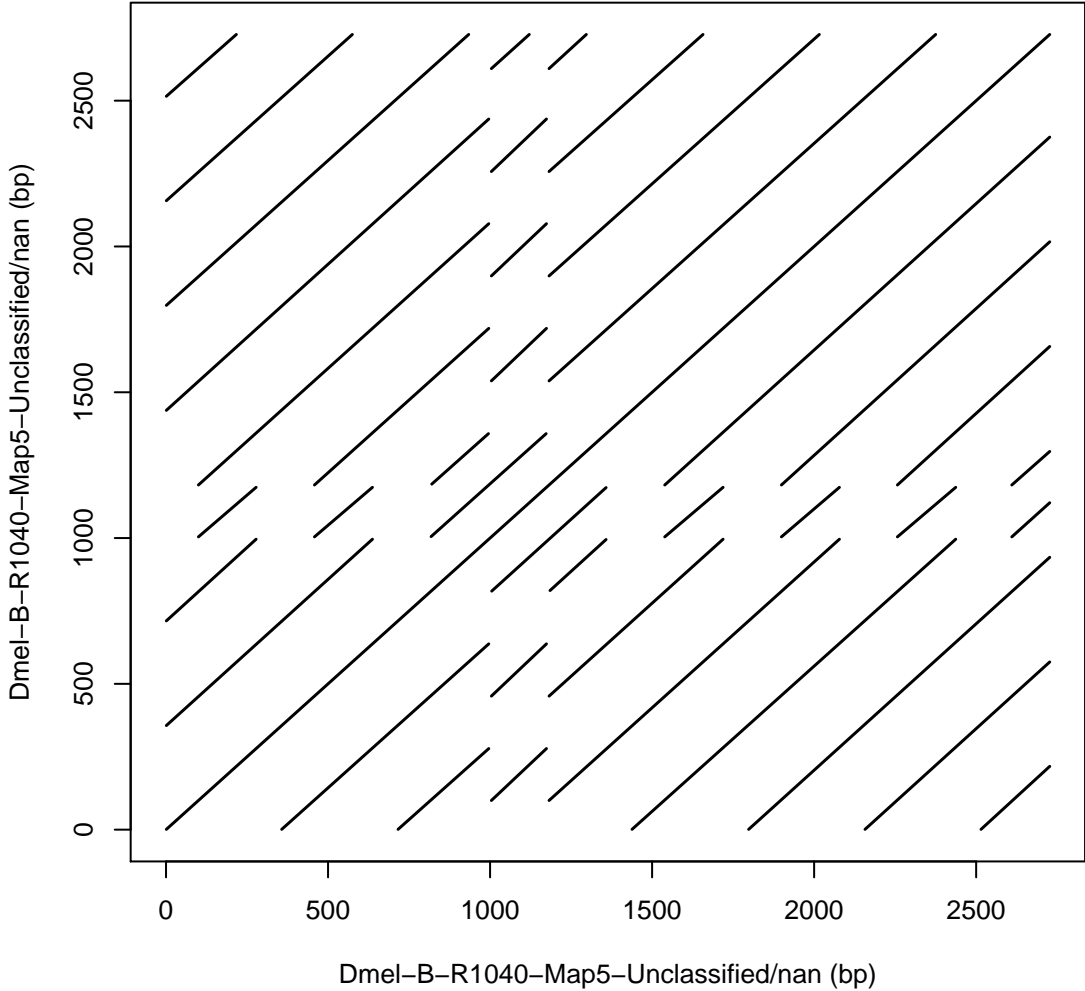
TE: Dmel-B-R1040-Map5-Unclassified/nan
consensus size: 2727bp; fragments: 37220; full length: 31 (>=2454.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

