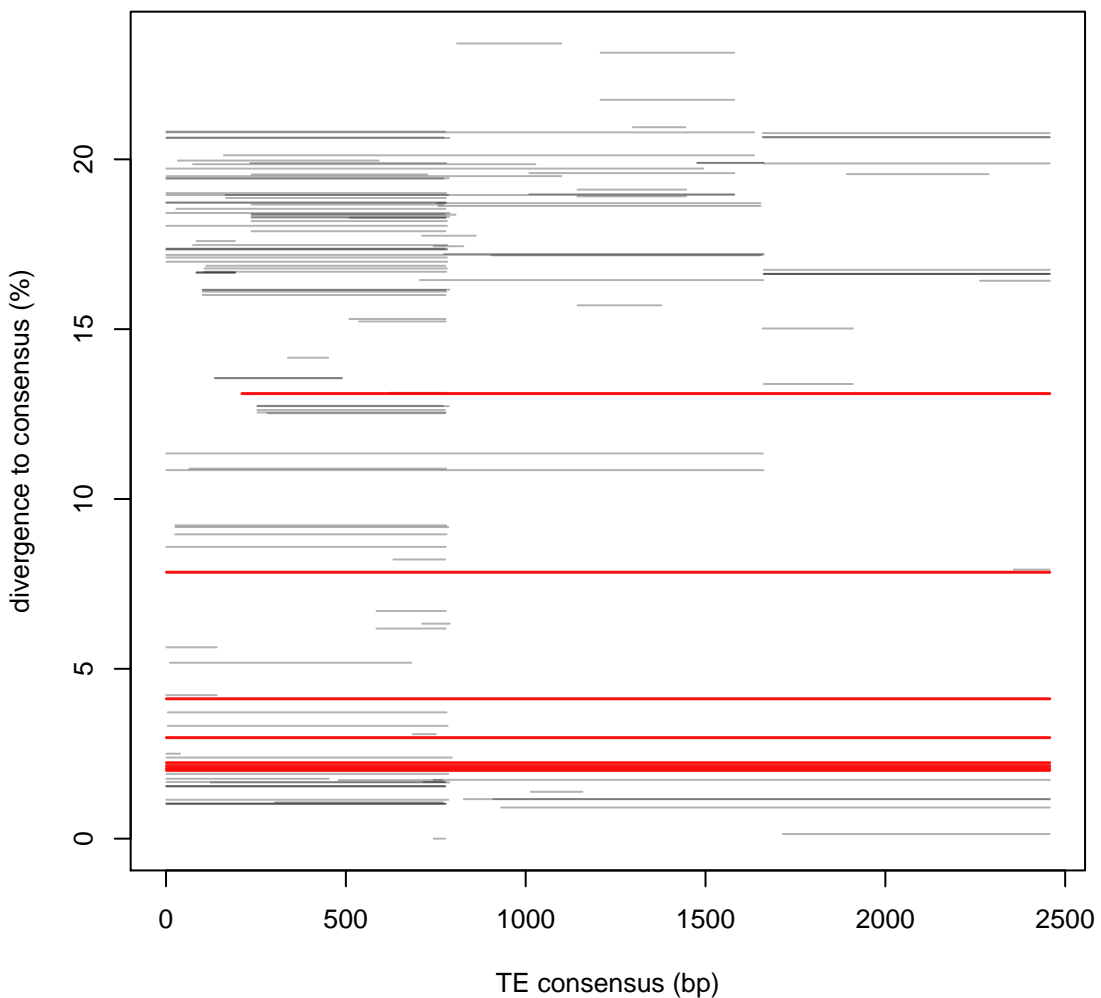
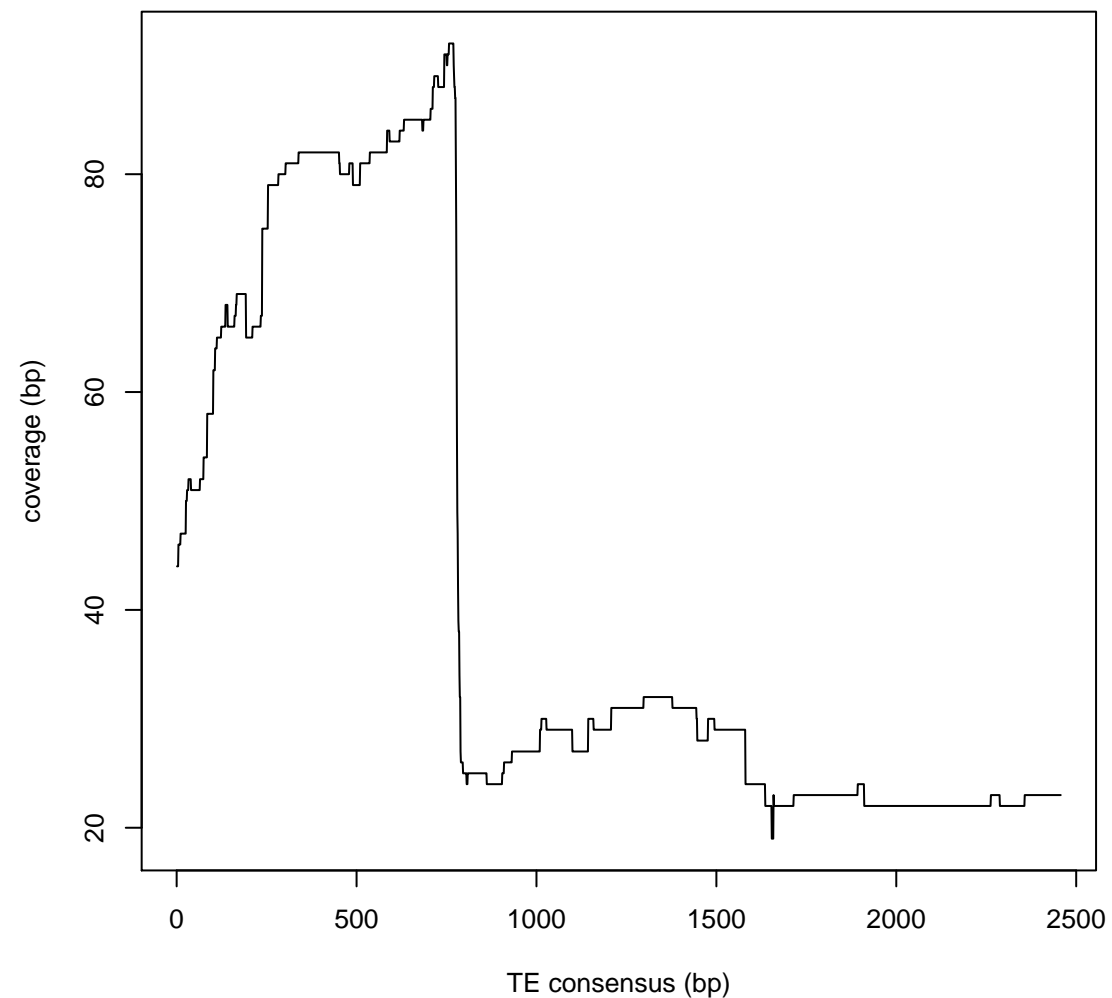


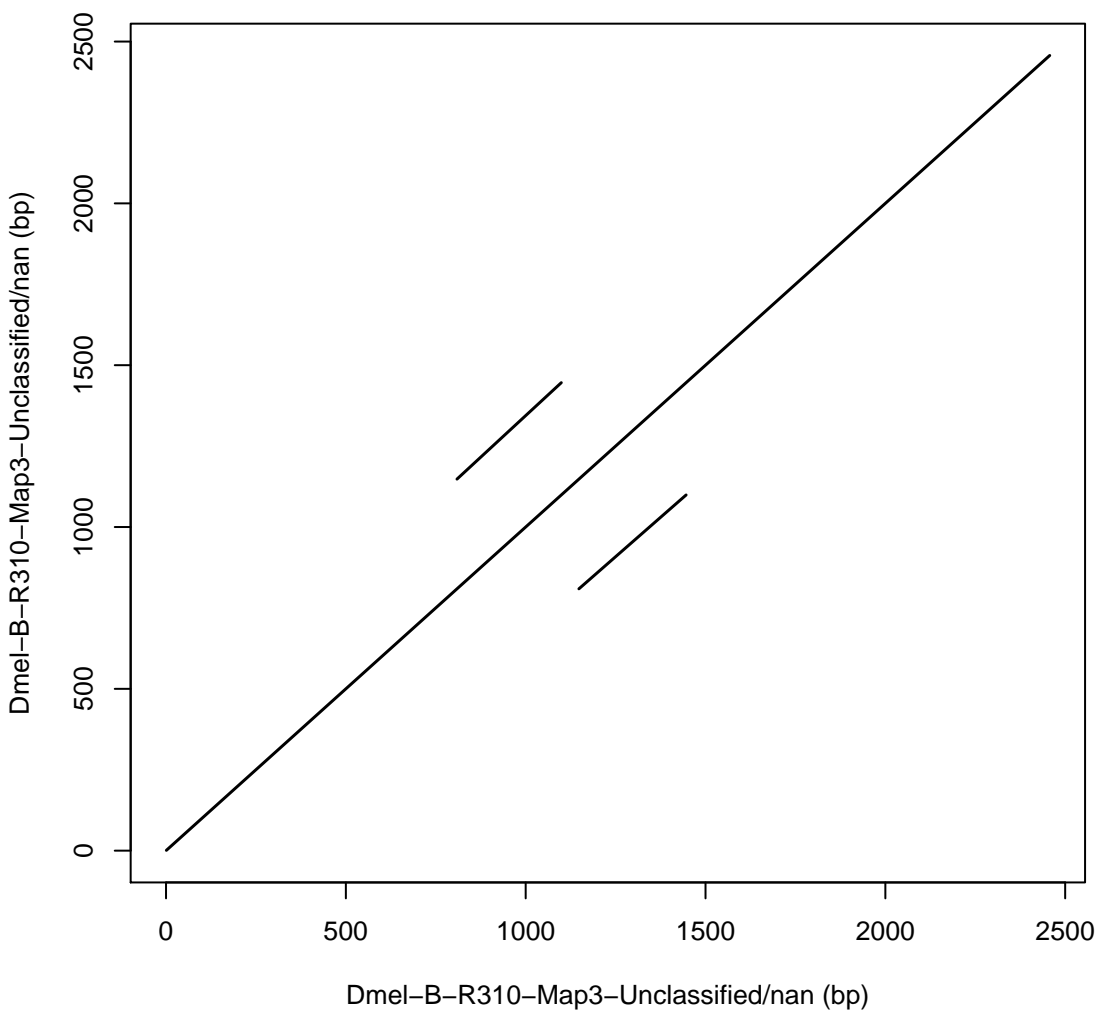
TE: Dmel-B-R310-Map3-Unclassified/nan
consensus size: 2457bp; fragments: 138; full length: 8 (>=2211.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

