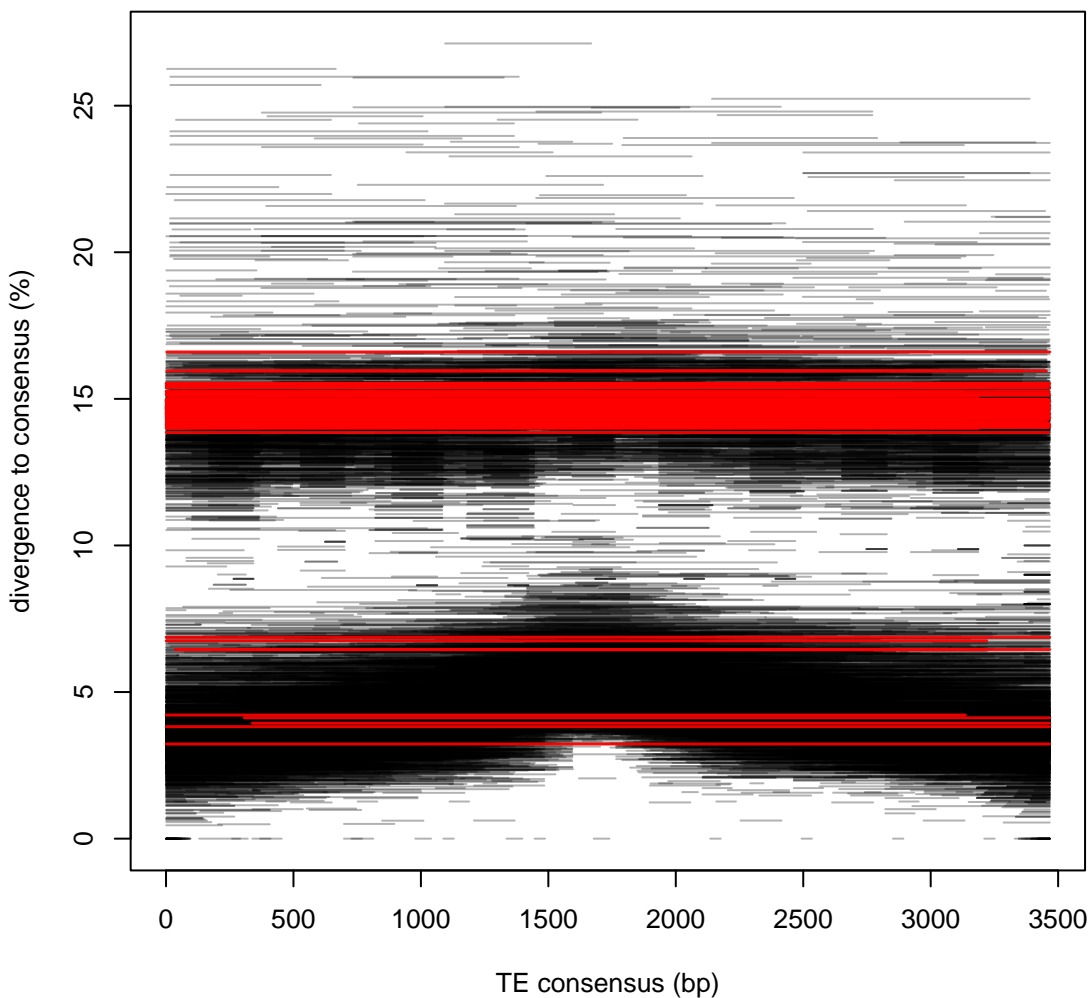
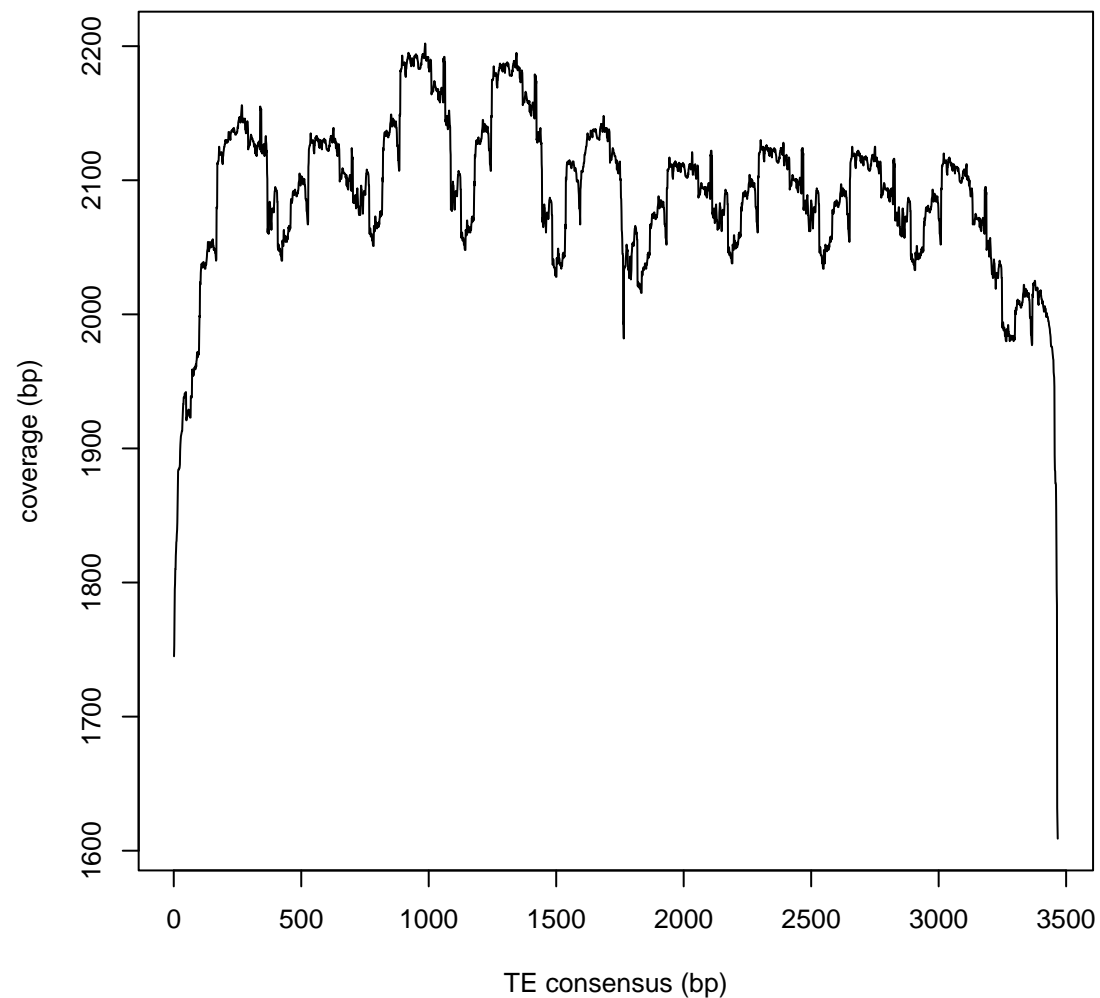


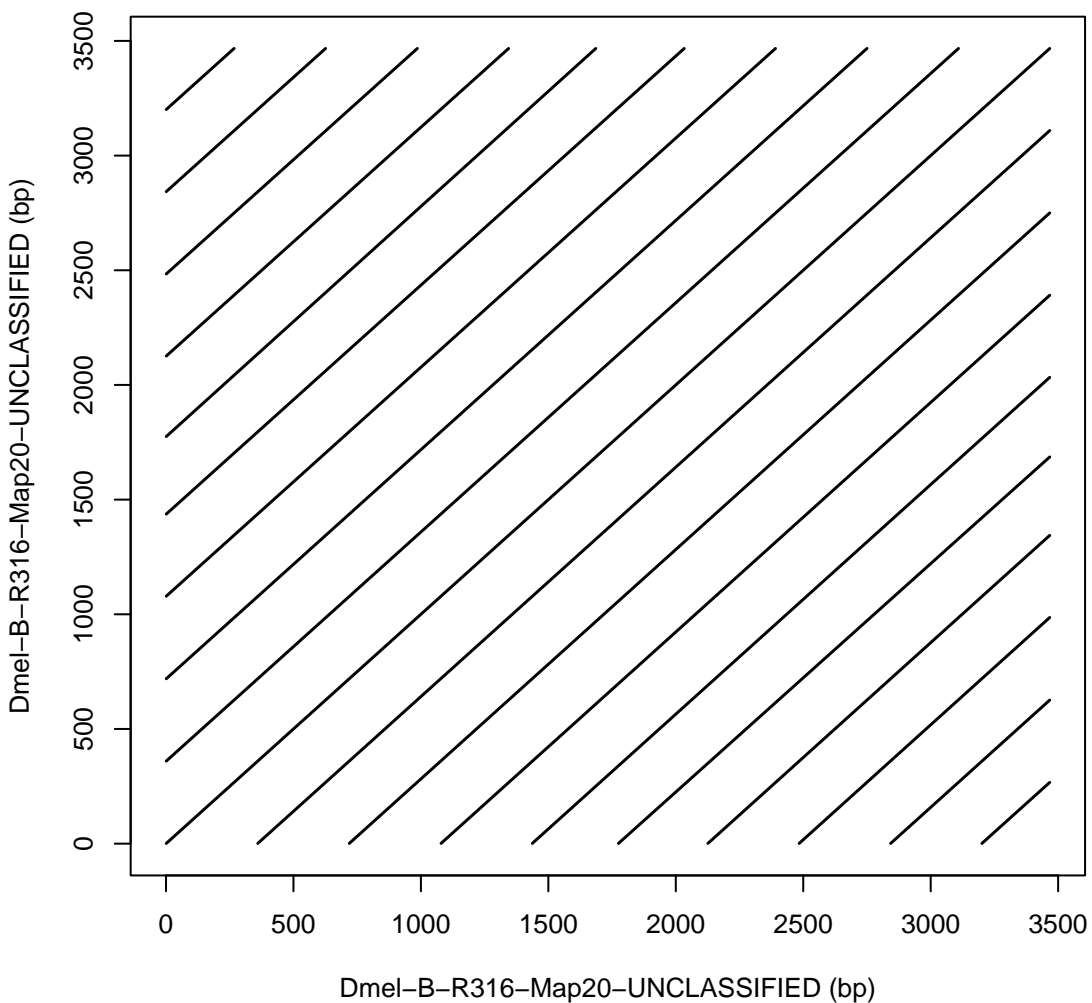
TE: Dmel-B-R316-Map20-UNCLASSIFIED
consensus size: 3467bp; fragments: 8350; full length: 94 (>=3120.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

