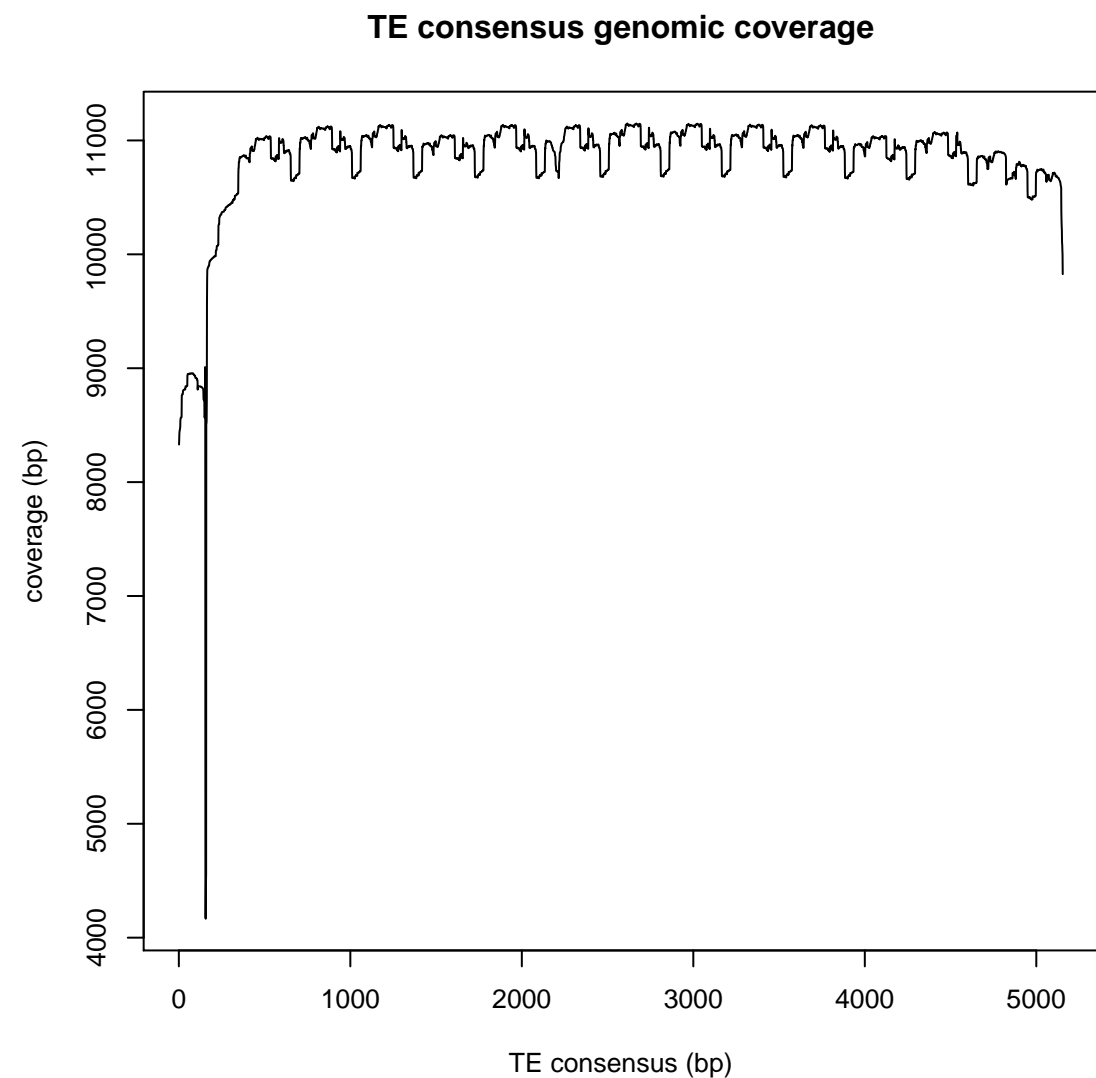
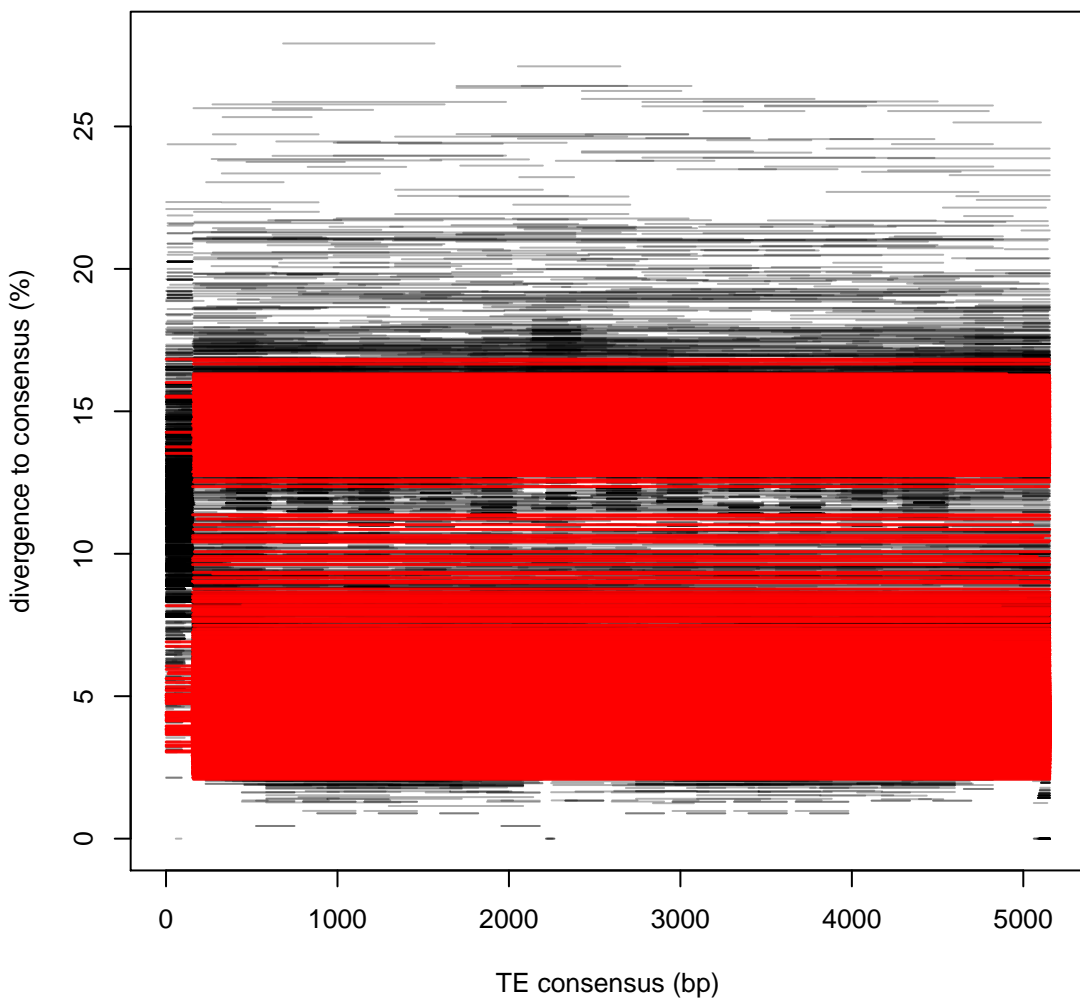
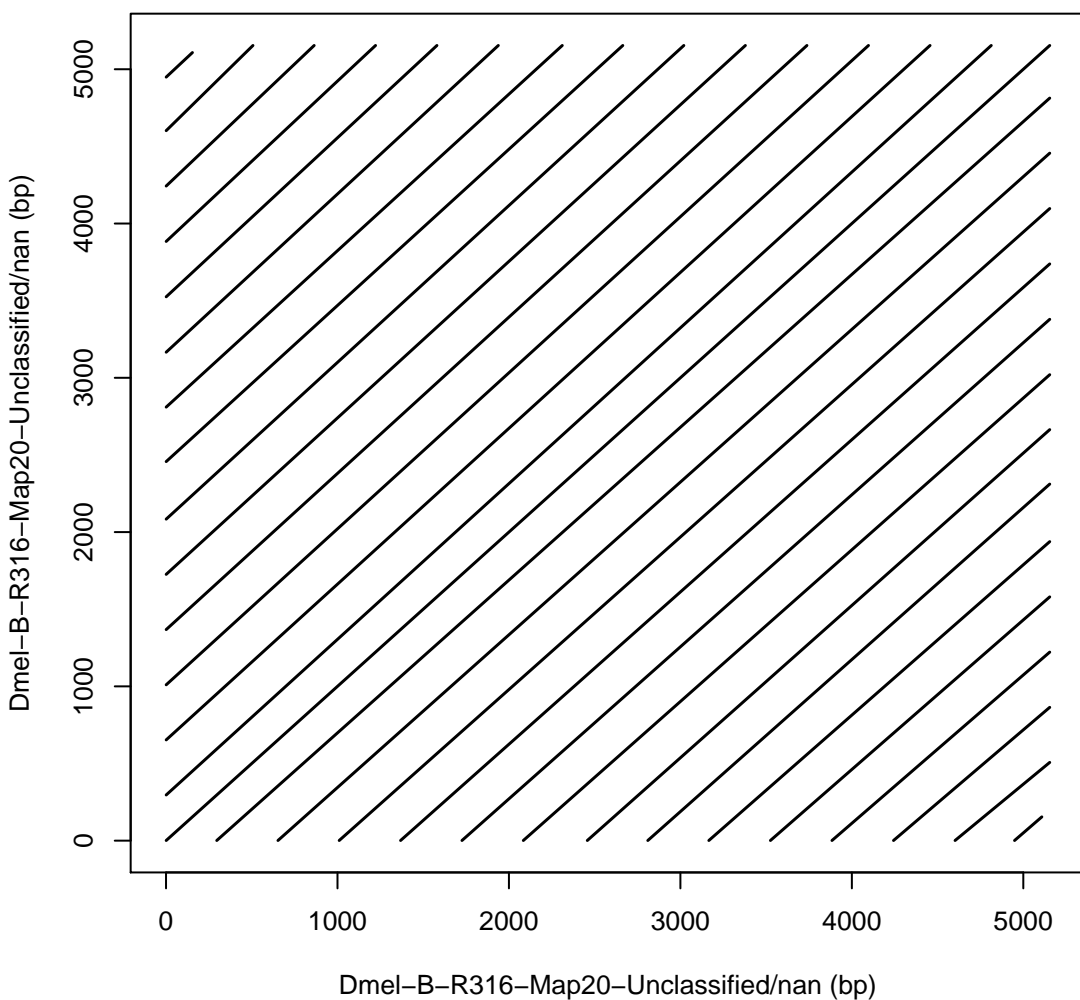


TE: Dmel-B-R316-Map20-Unclassified/nan
consensus size: 5154bp; fragments: 38114; full length: 4509 (≥ 4638.6 bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

