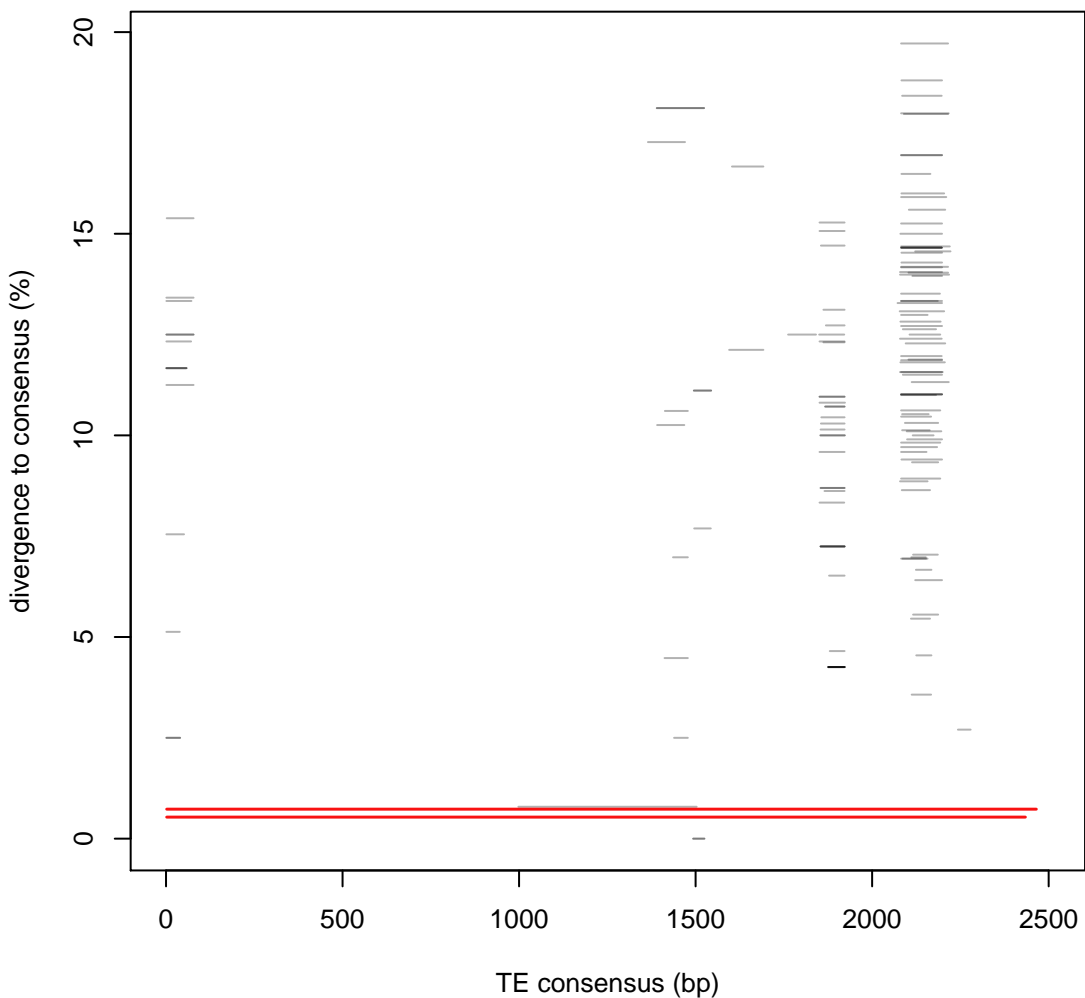
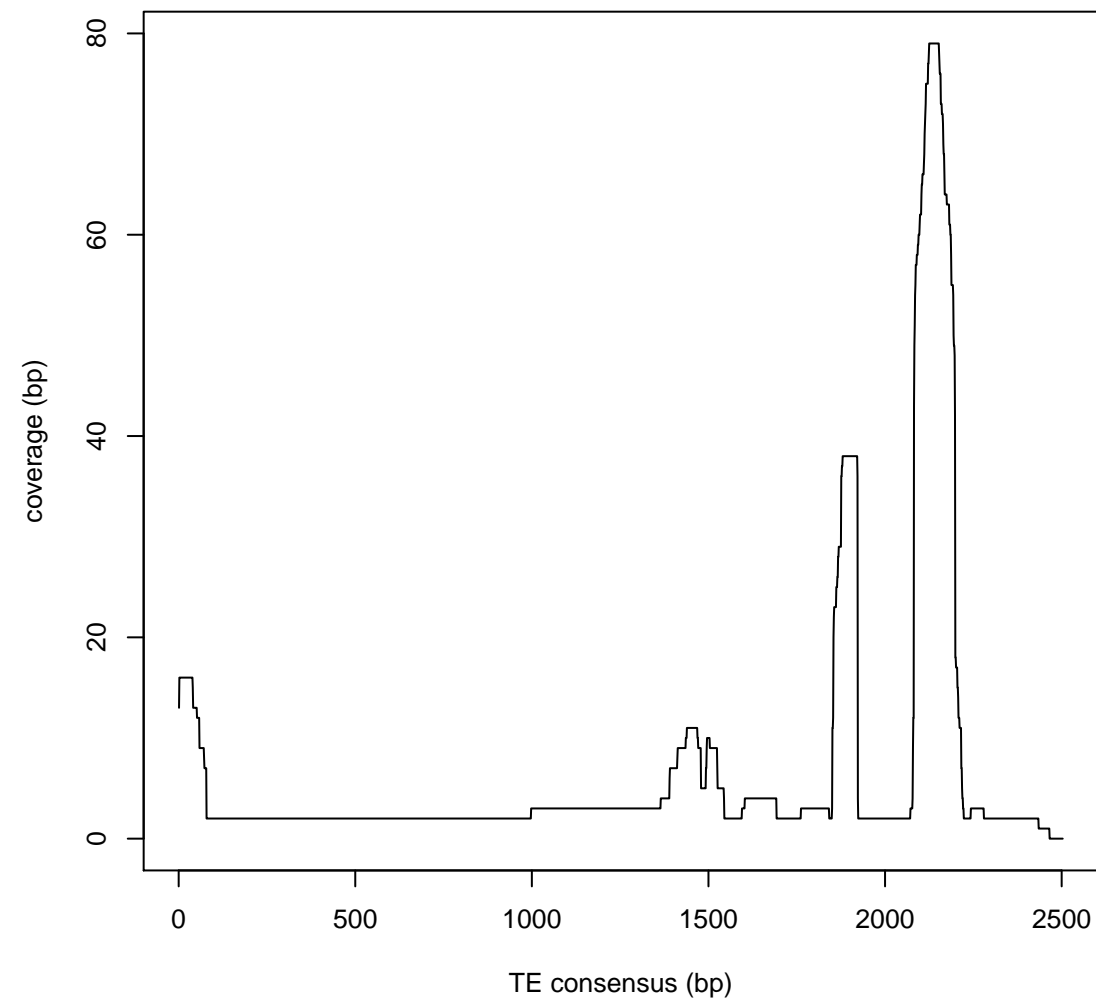


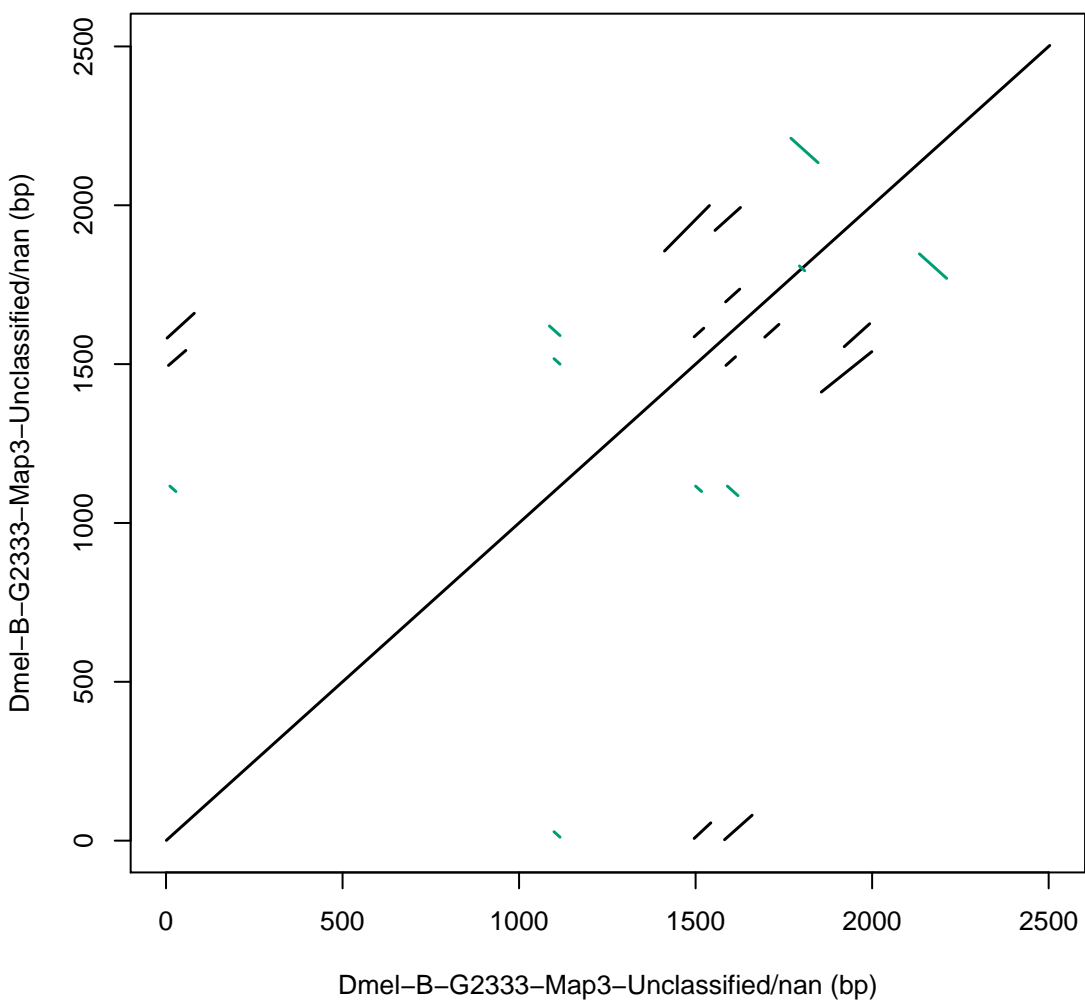
TE: Dmel-B-G2333-Map3-Unclassified/nan
consensus size: 2503bp; fragments: 147; full length: 2 (≥ 2252.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

