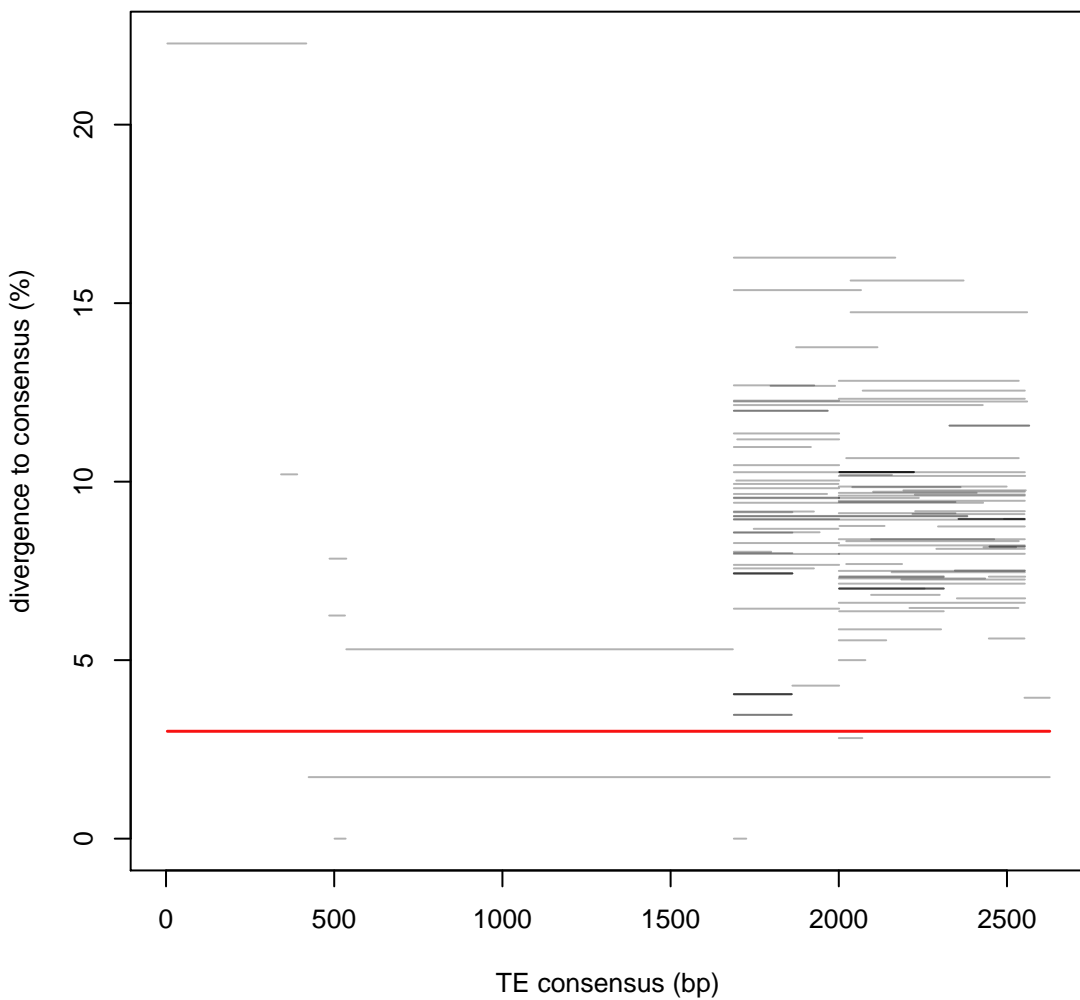
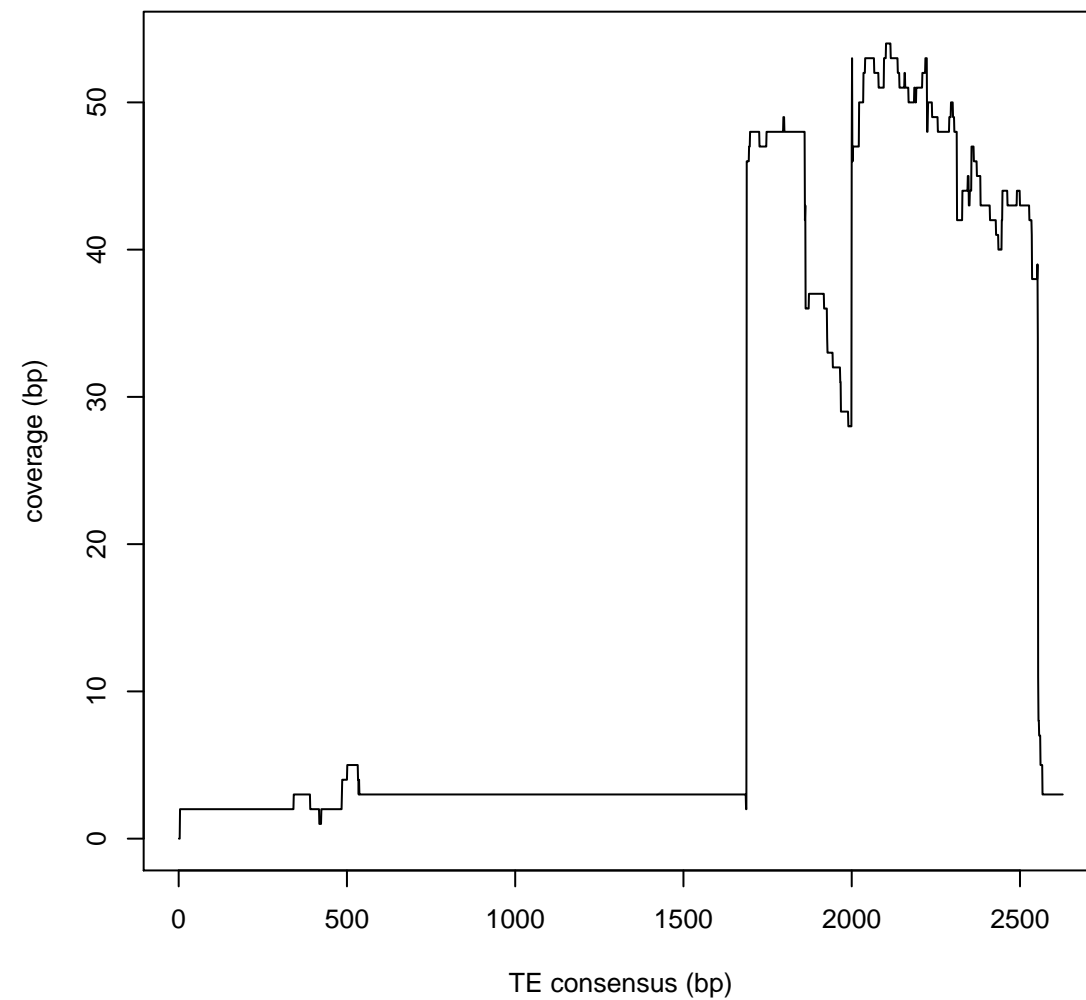


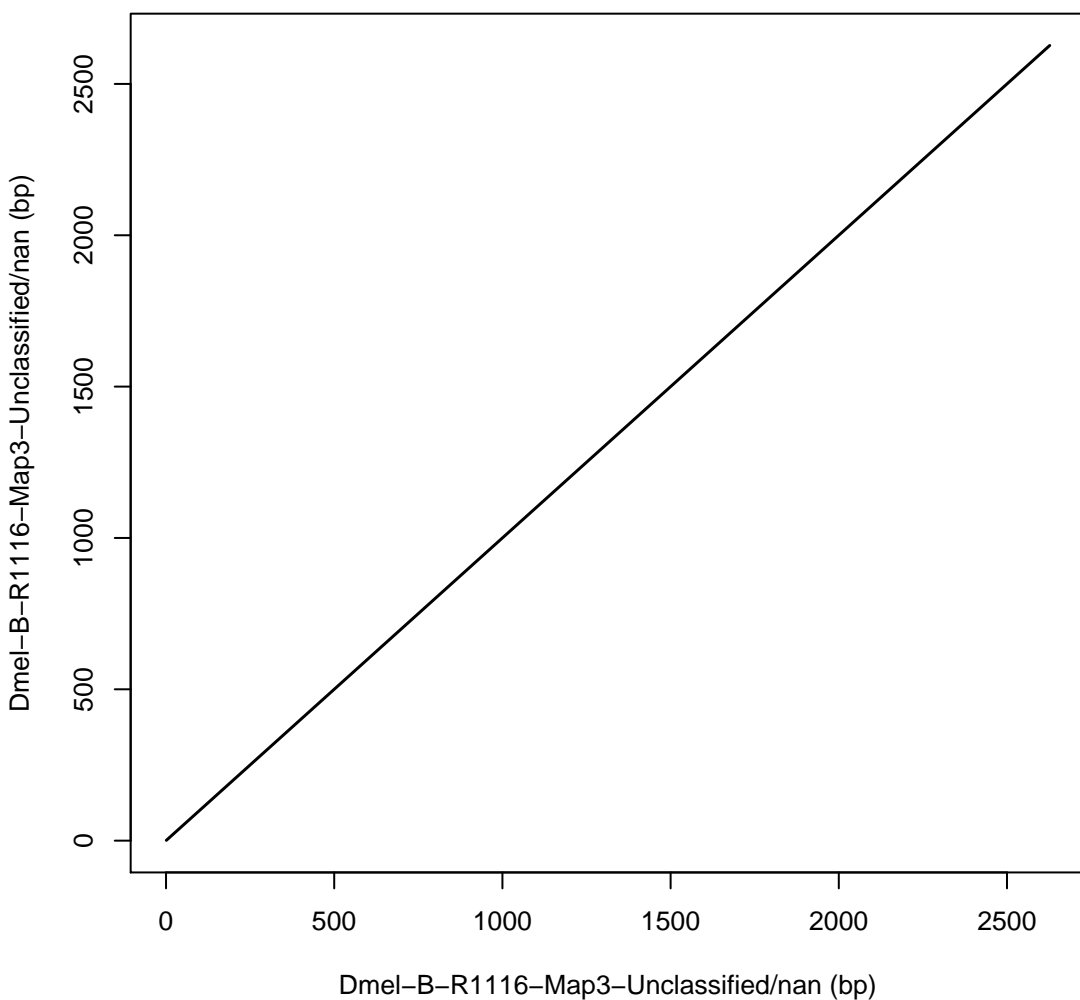
TE: Dmel-B-R1116-Map3-Unclassified/nan
consensus size: 2627bp; fragments: 126; full length: 1 (>=2364.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

