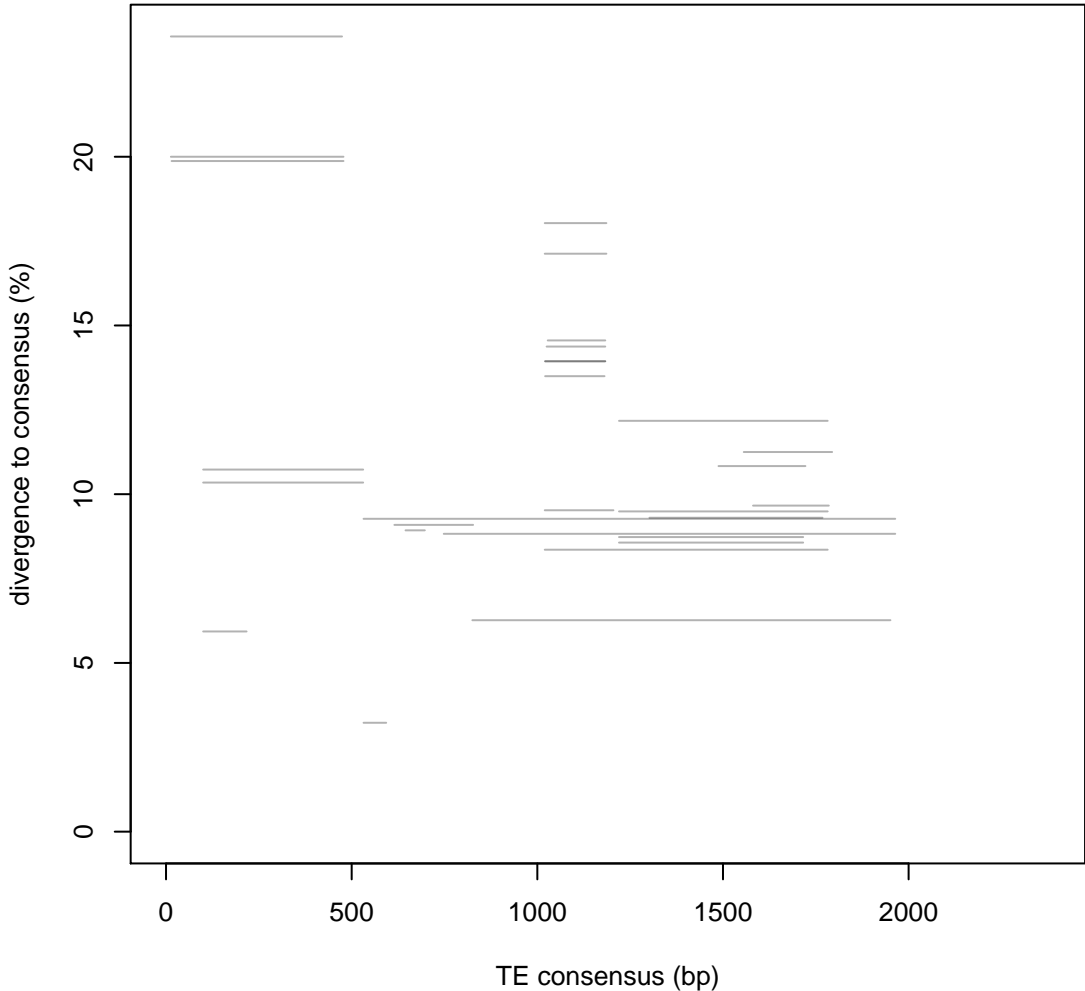
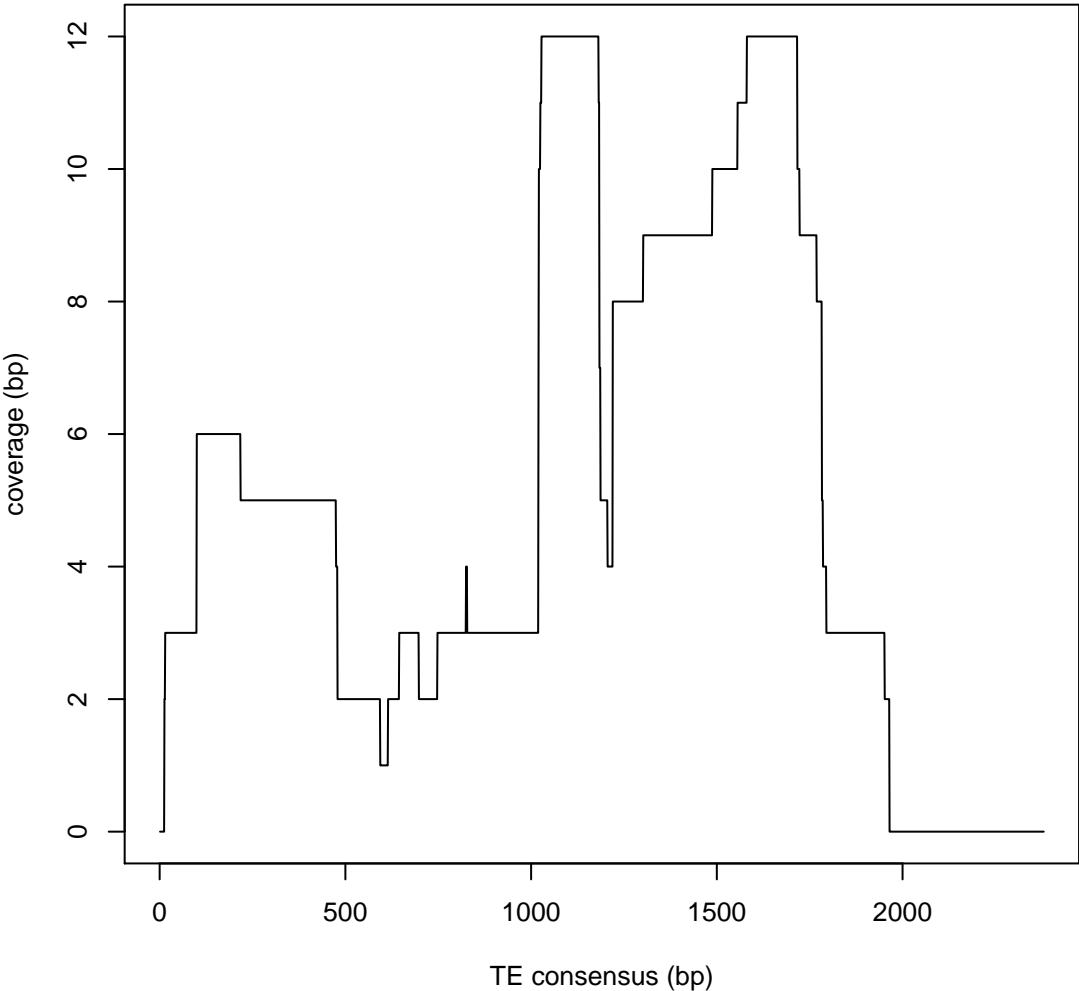


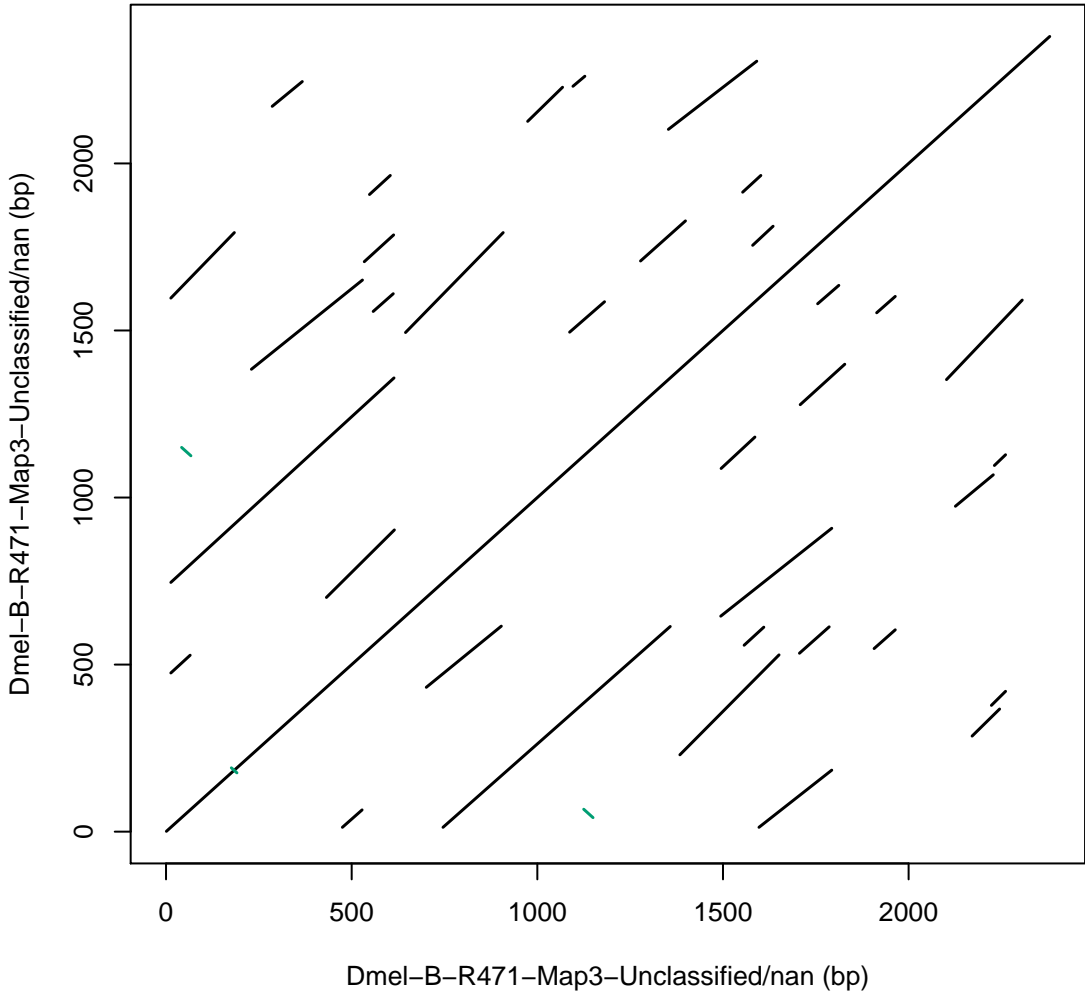
TE: Dmel-B-R471-Map3-Unclassified/nan  
consensus size: 2380bp; fragments: 29; full length: 0 (>=2142bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

