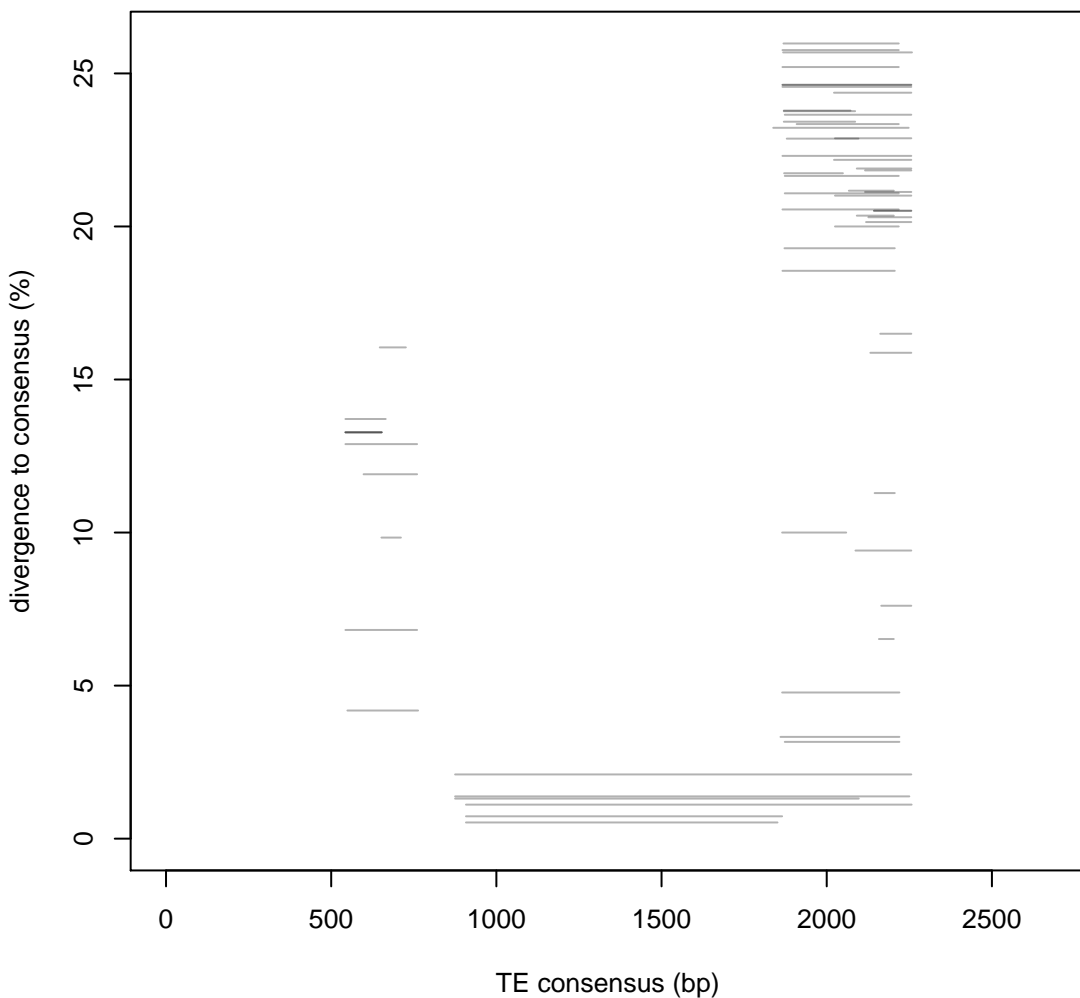
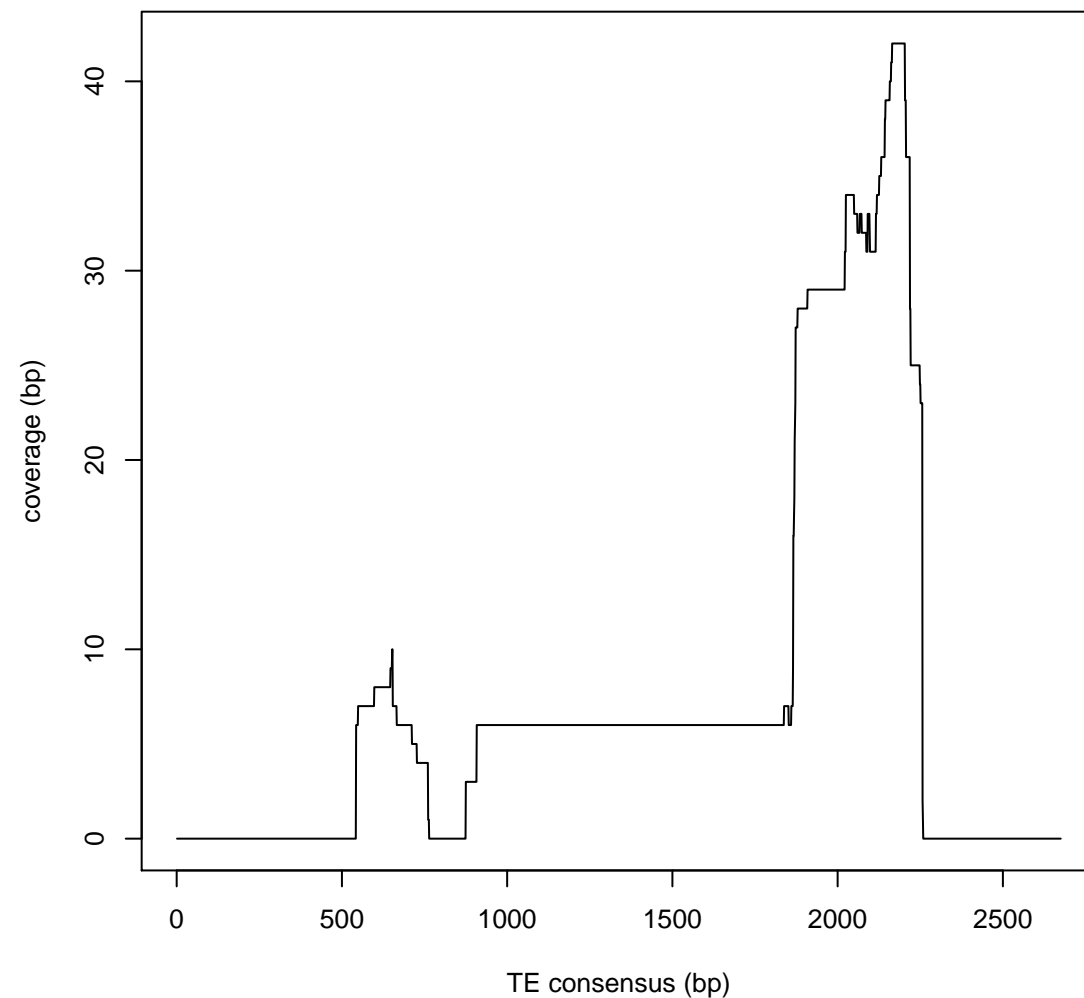


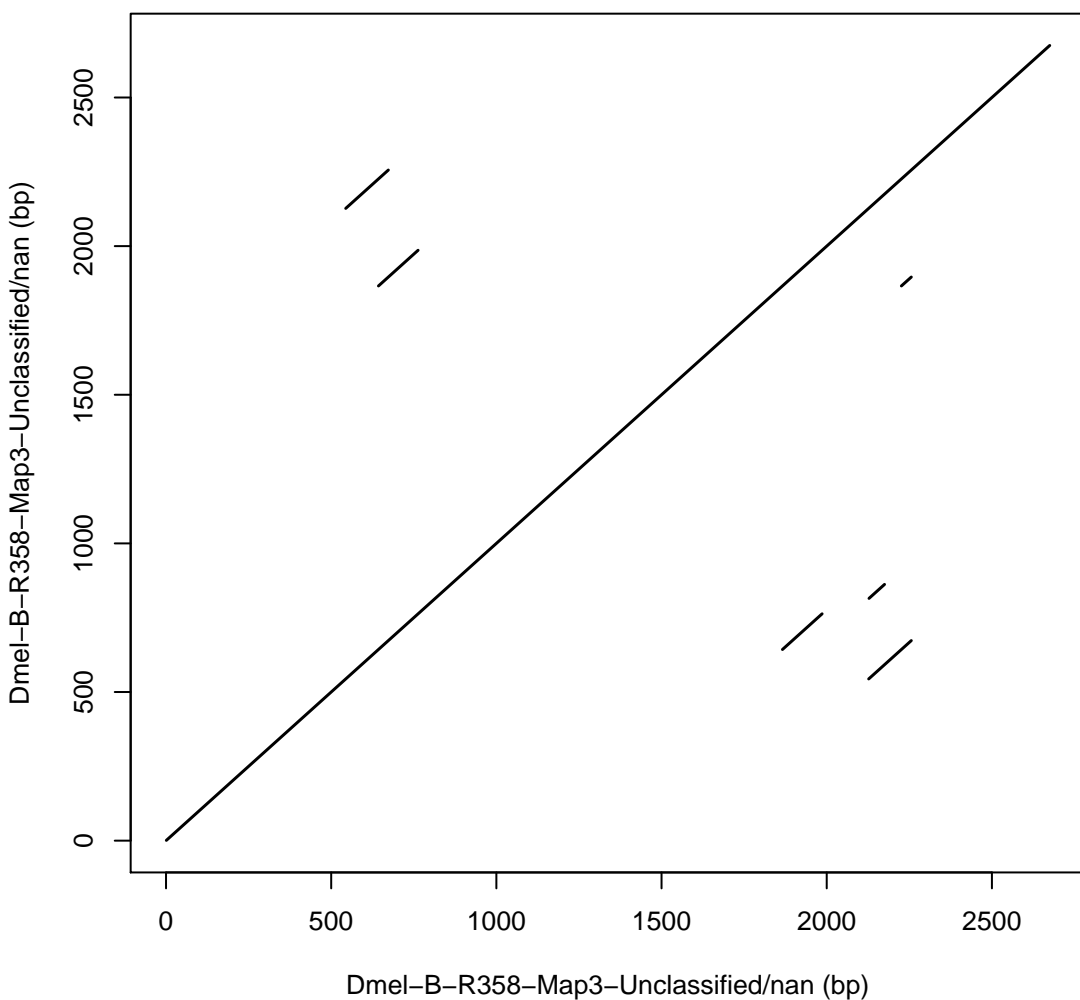
TE: Dmel-B-R358-Map3-Unclassified/nan  
consensus size: 2675bp; fragments: 61; full length: 0 ( $\geq 2407.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

