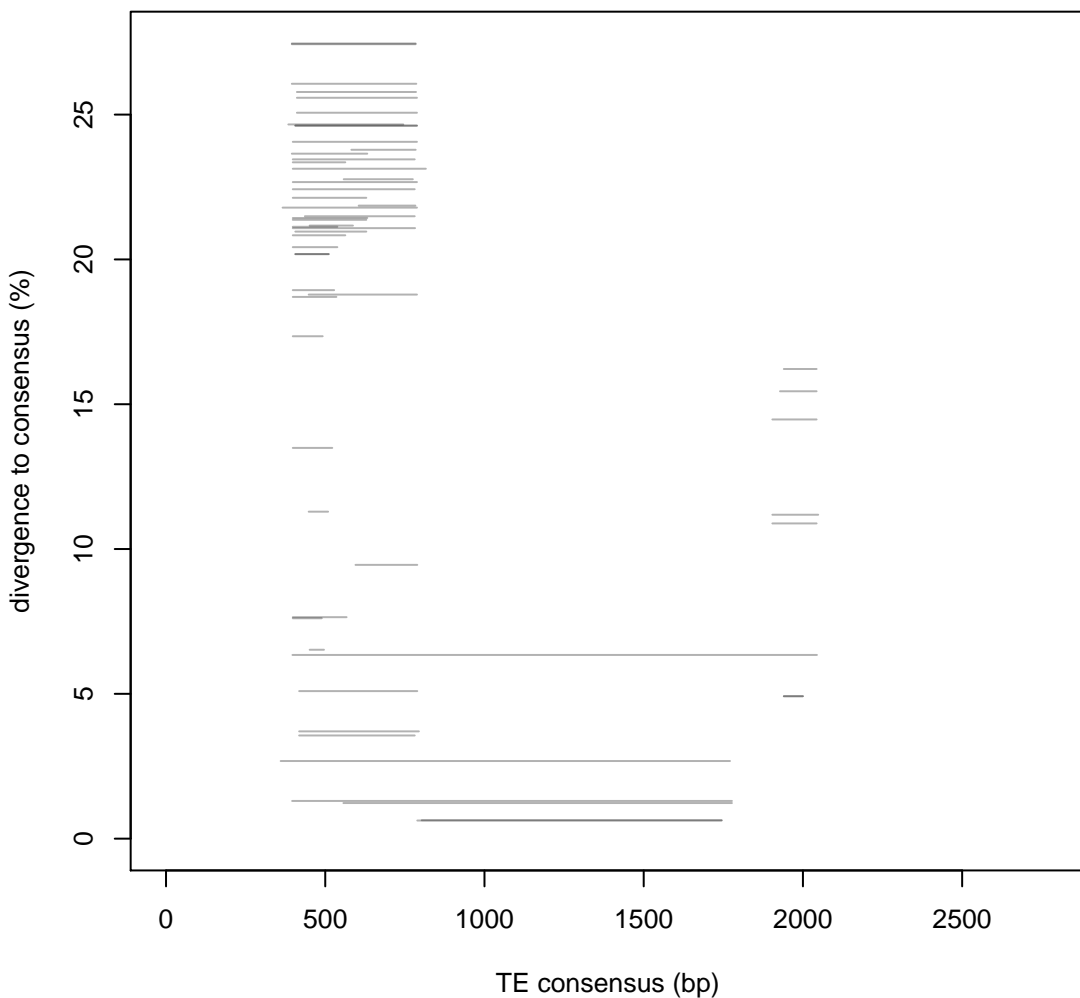
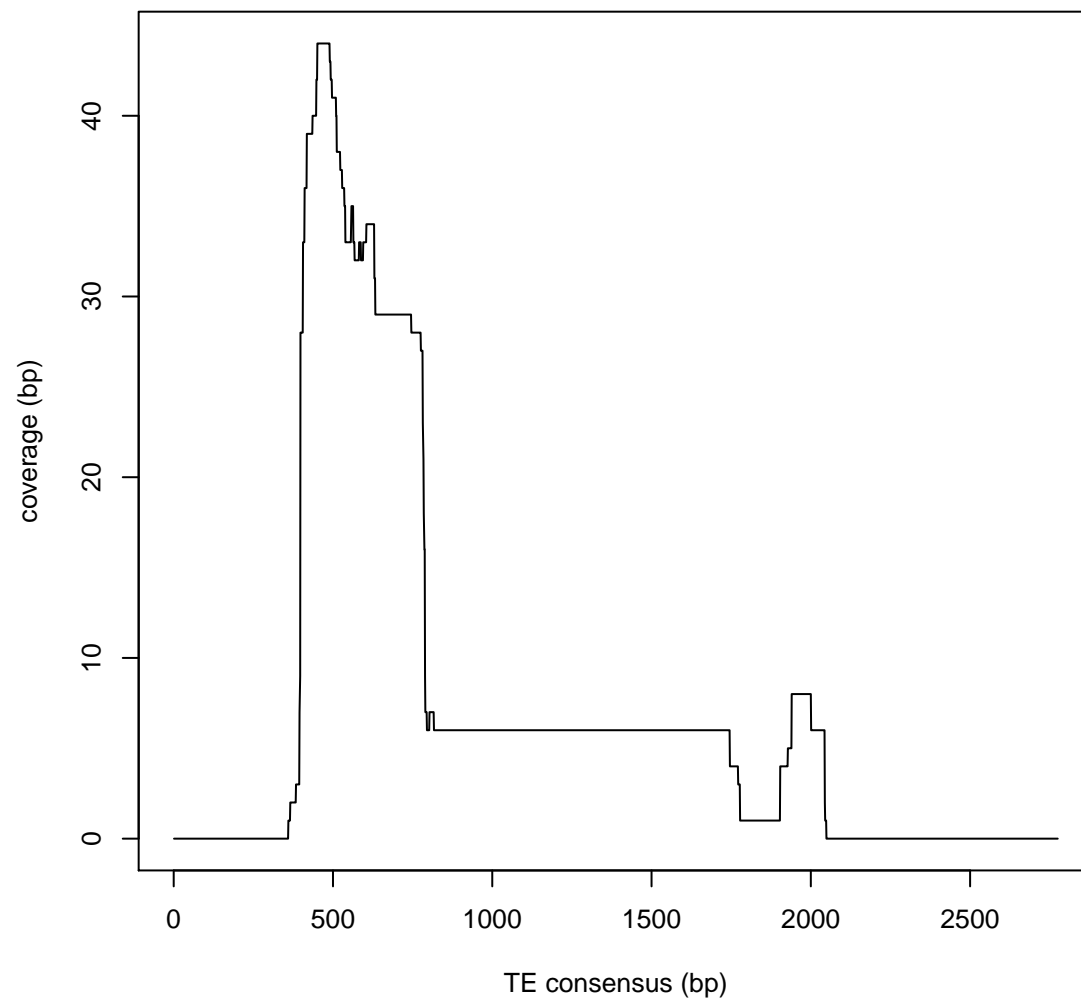


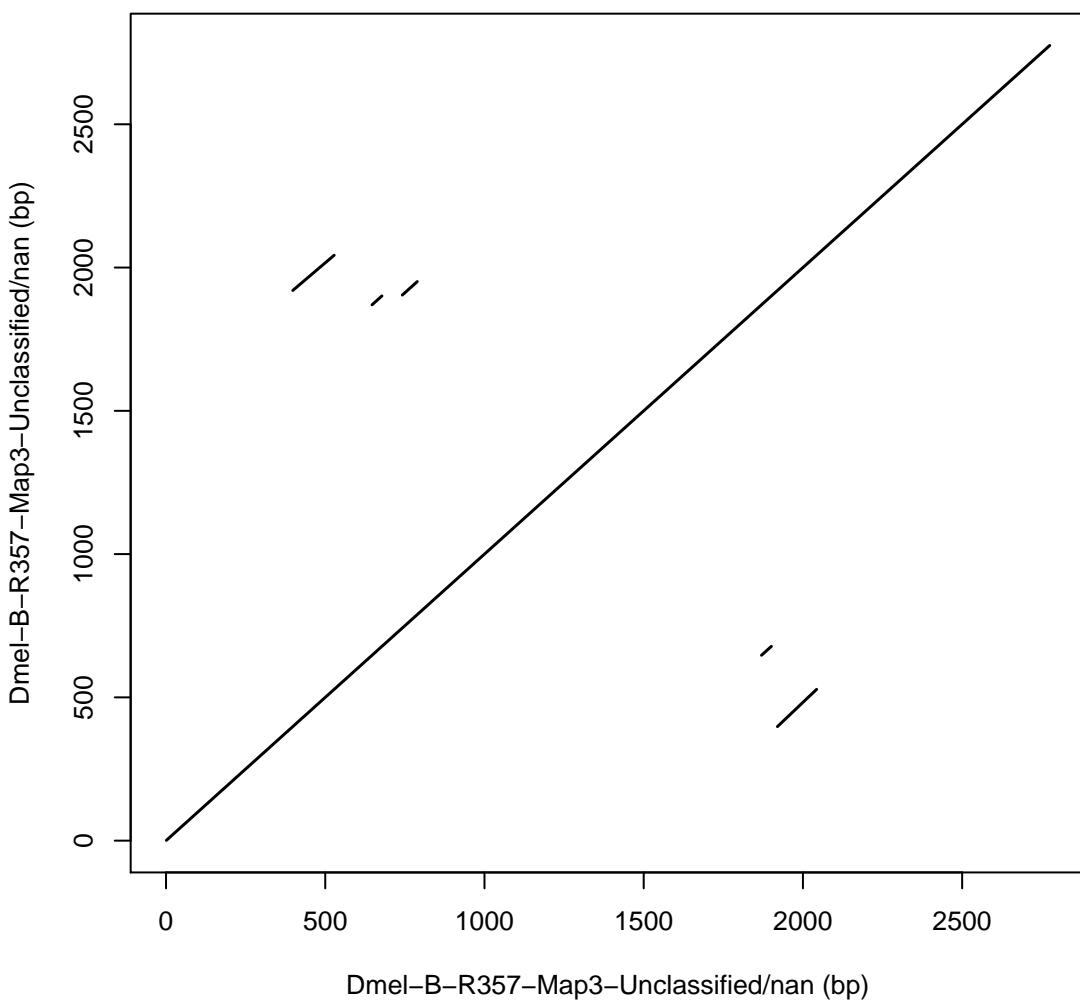
TE: Dmel-B-R357-Map3-Unclassified/nan  
consensus size: 2775bp; fragments: 58; full length: 0 ( $\geq 2497.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

