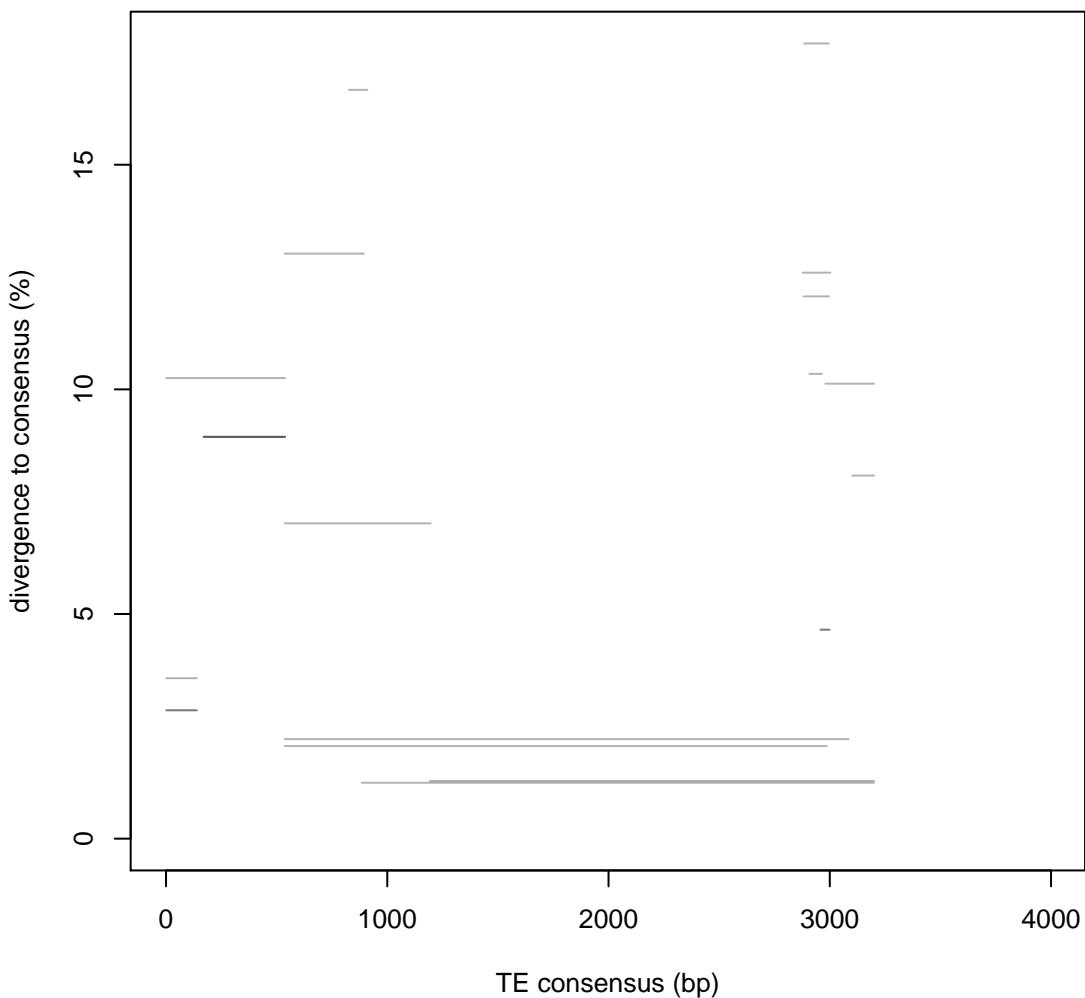
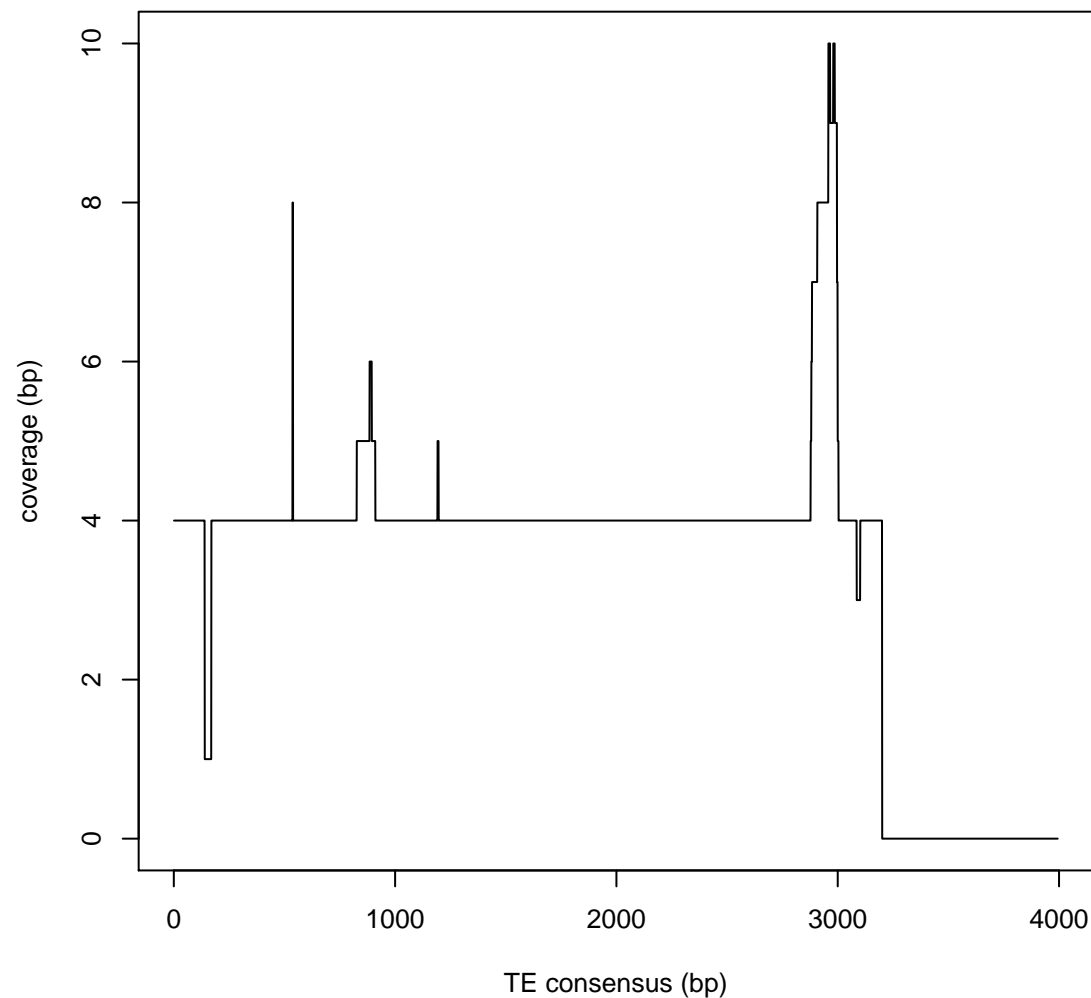


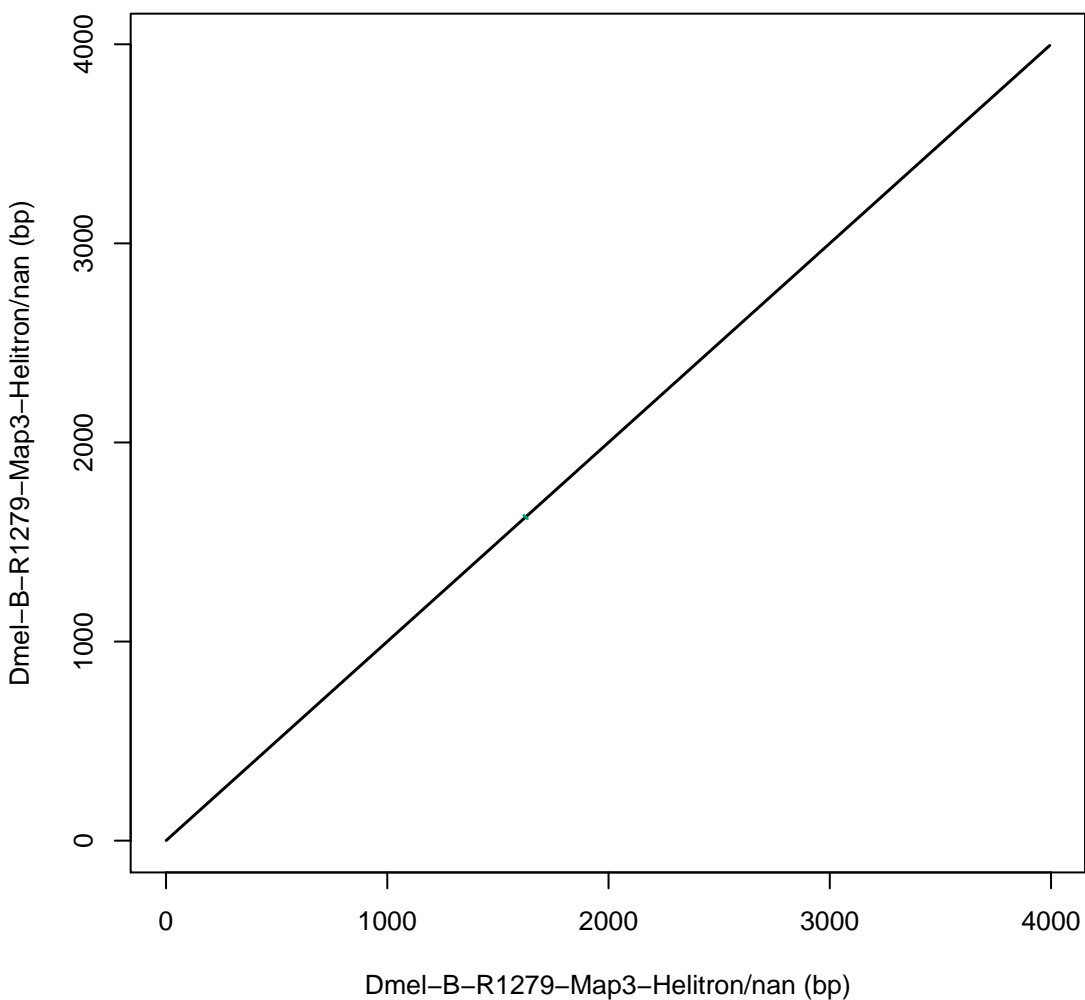
TE: Dmel-B-R1279-Map3-Helitron/nan  
consensus size: 3994bp; fragments: 22; full length: 0 ( $\geq 3594.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

