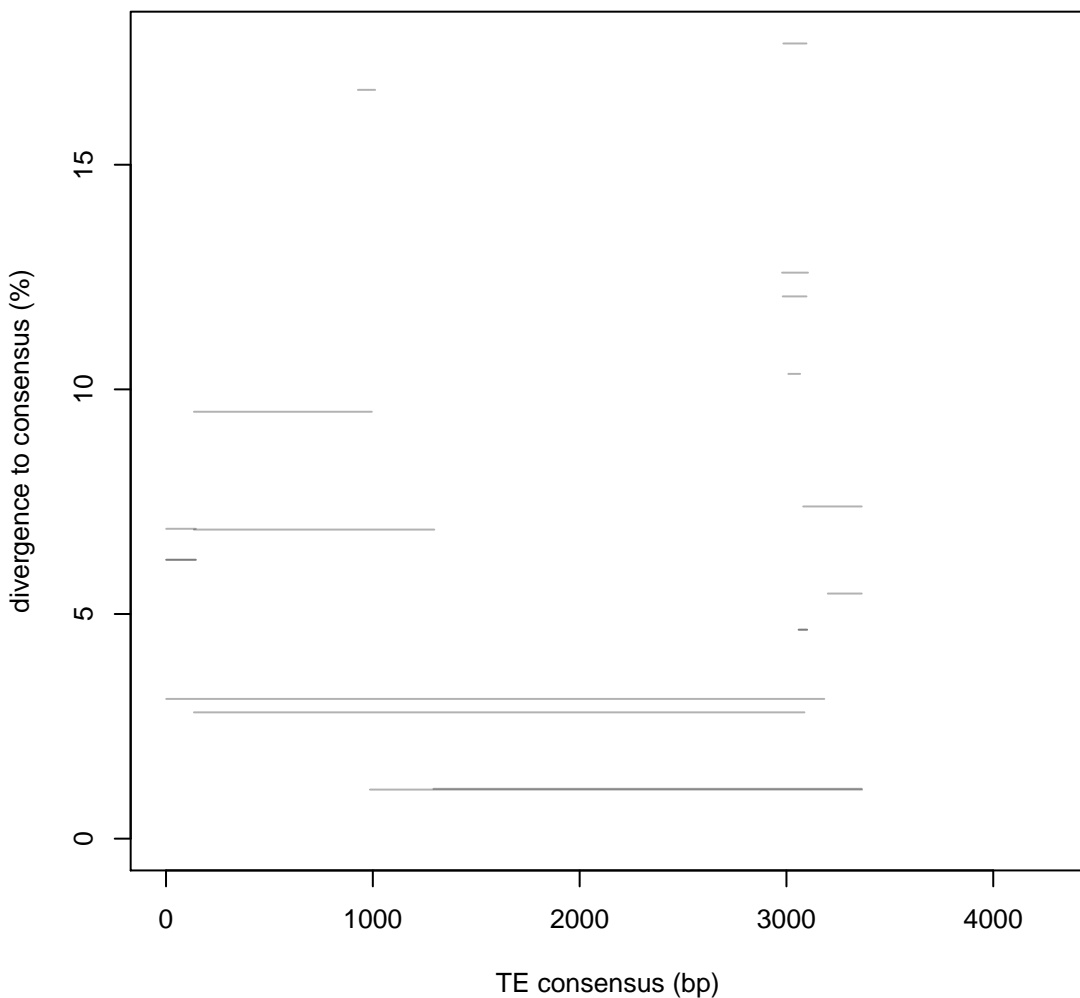
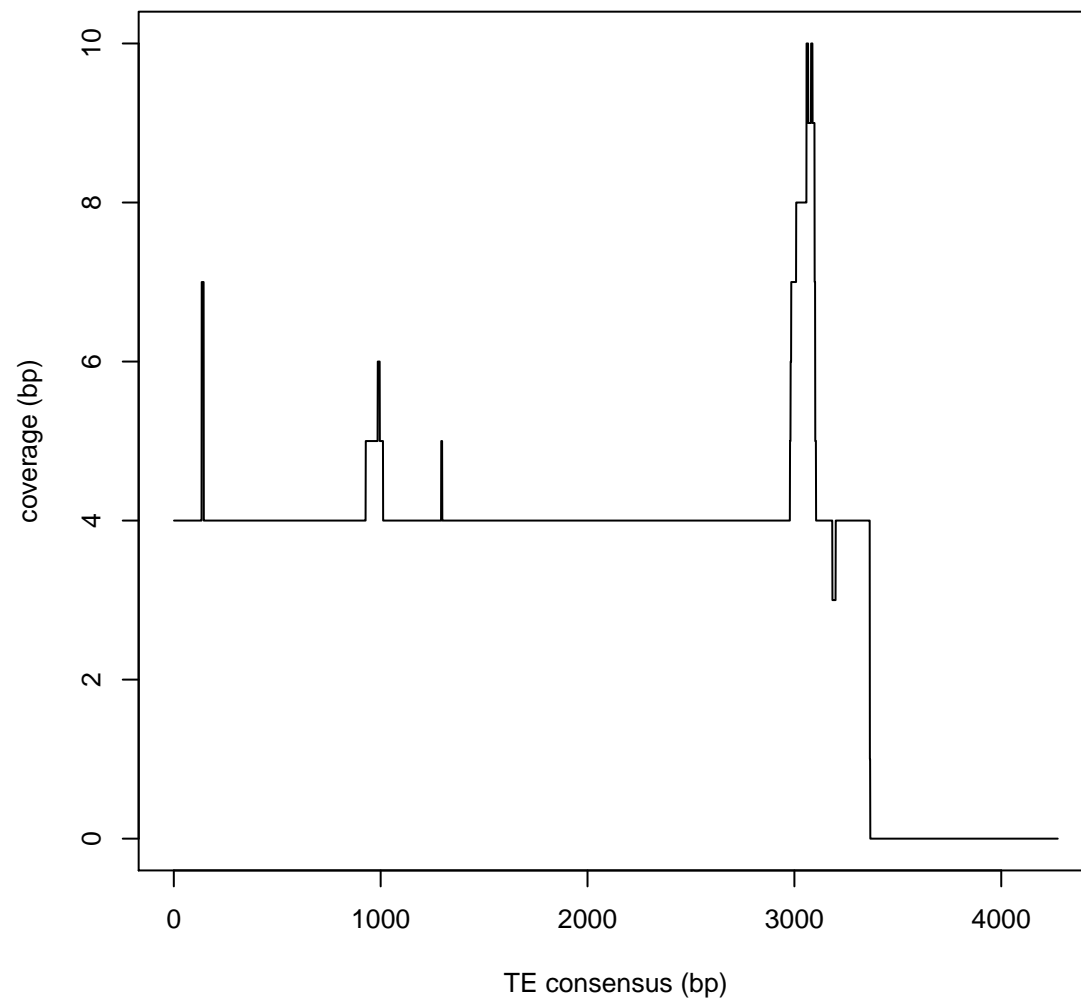


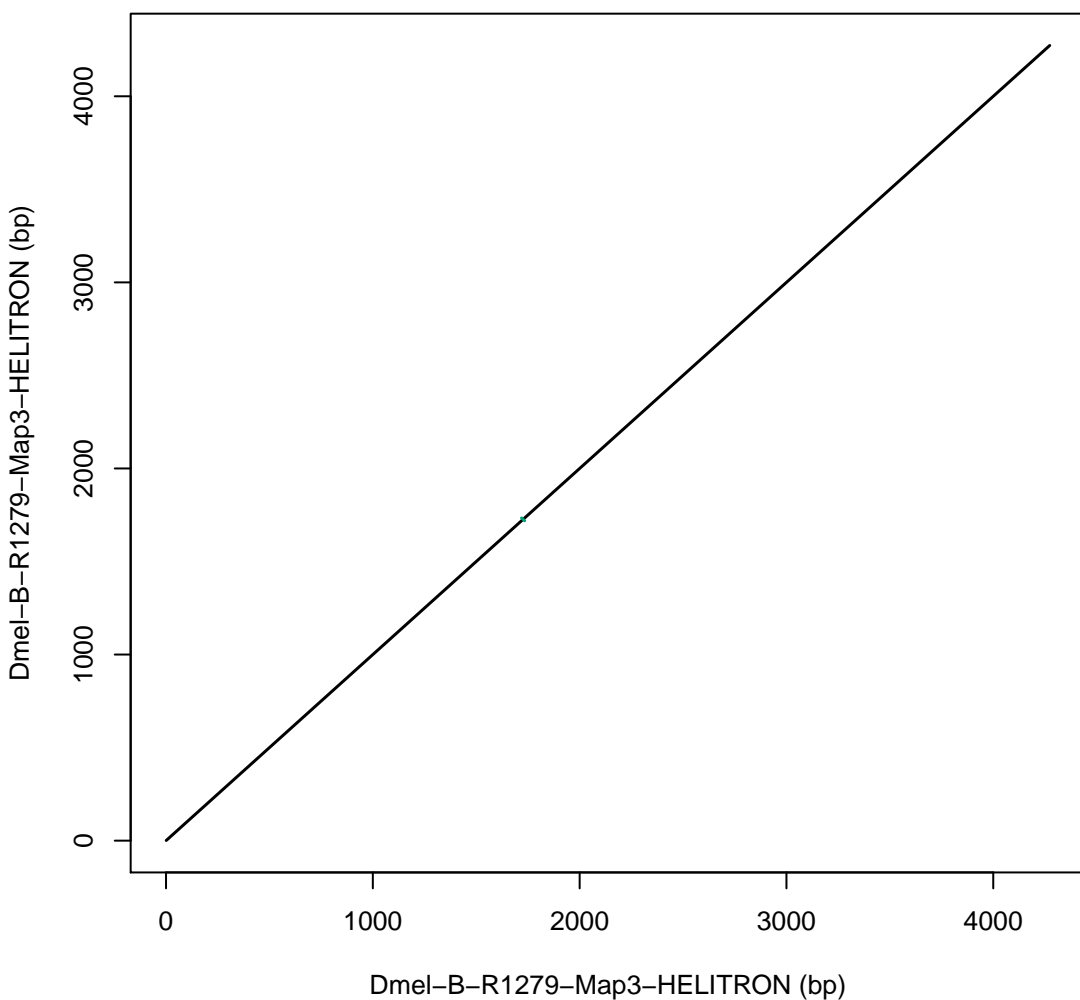
TE: Dmel-B-R1279-Map3-HELITRON
consensus size: 4273bp; fragments: 18; full length: 0 (≥ 3845.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

