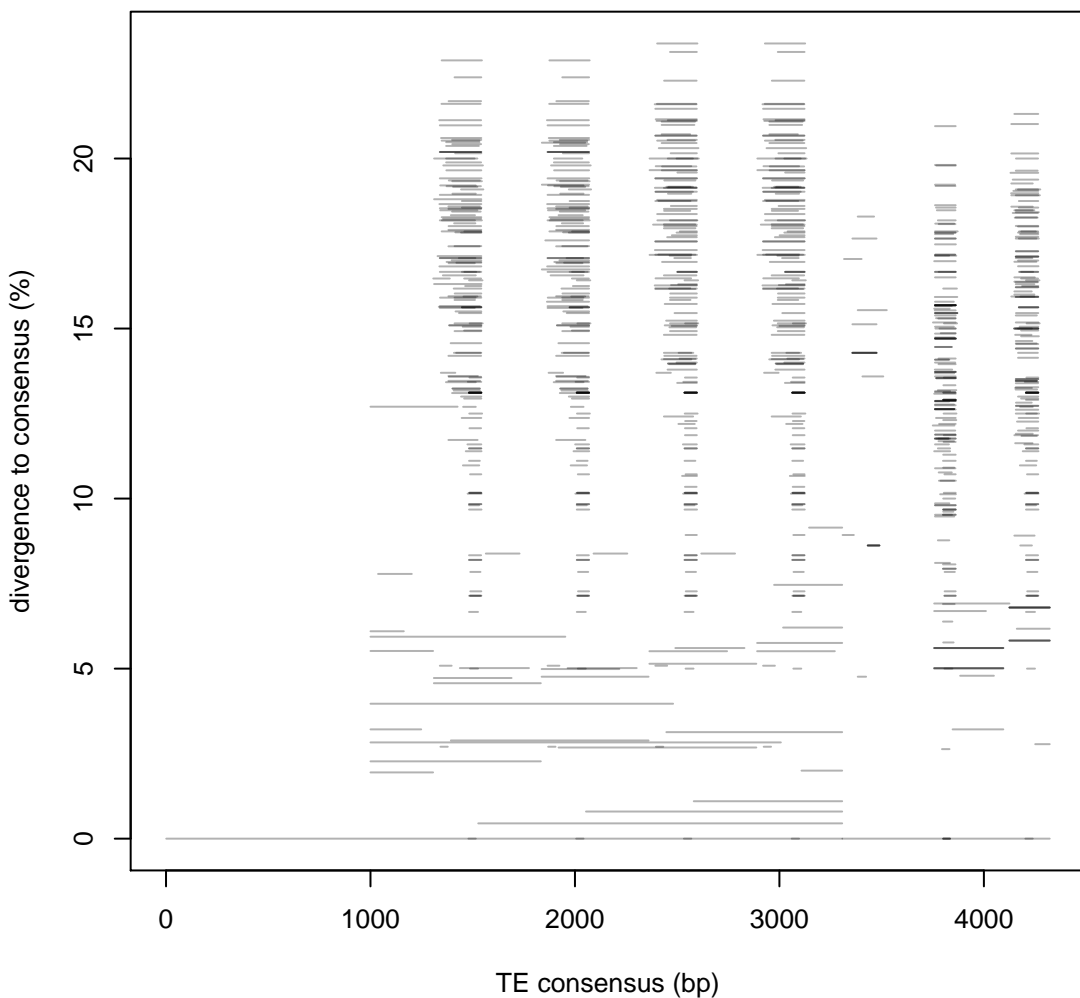
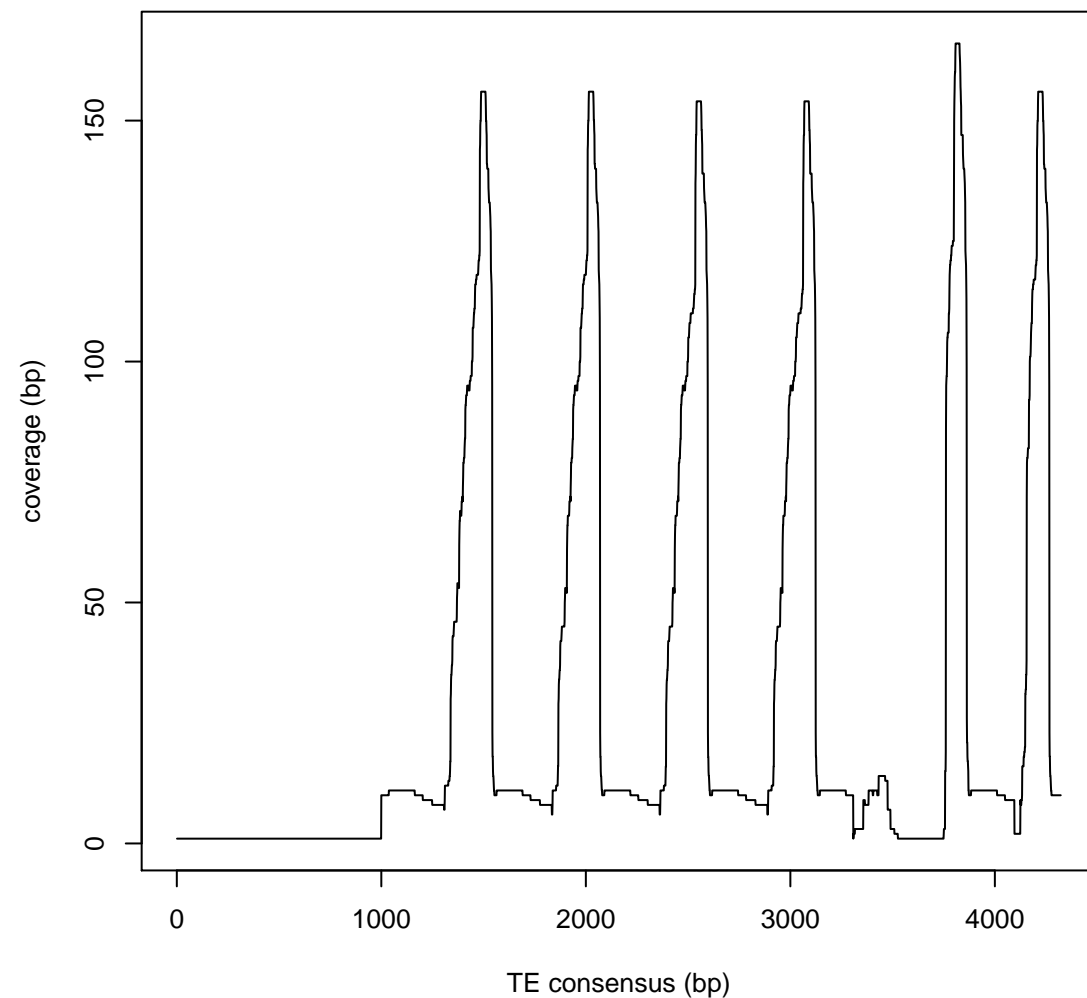


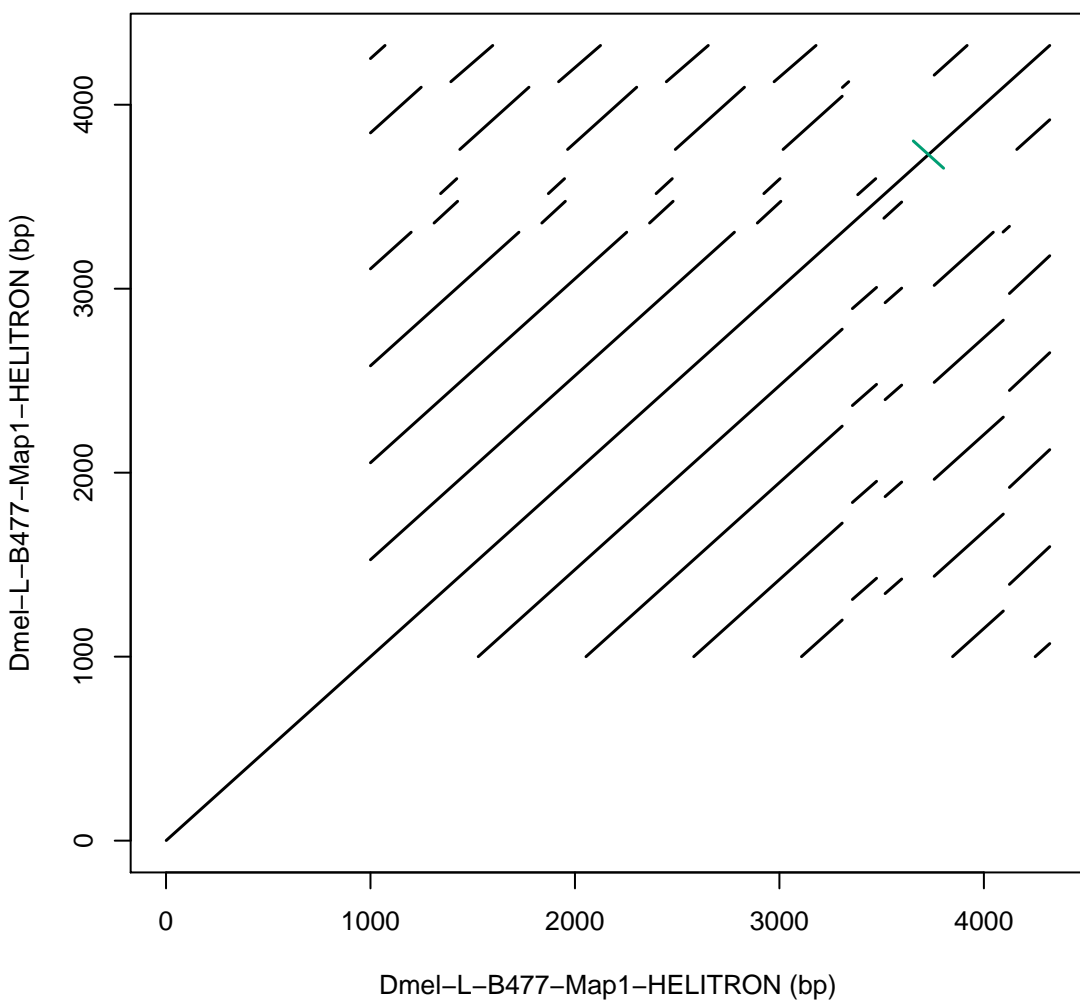
TE: Dmel-L-B477-Map1-HELITRON
consensus size: 4322bp; fragments: 972; full length: 0 (>=3889.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

