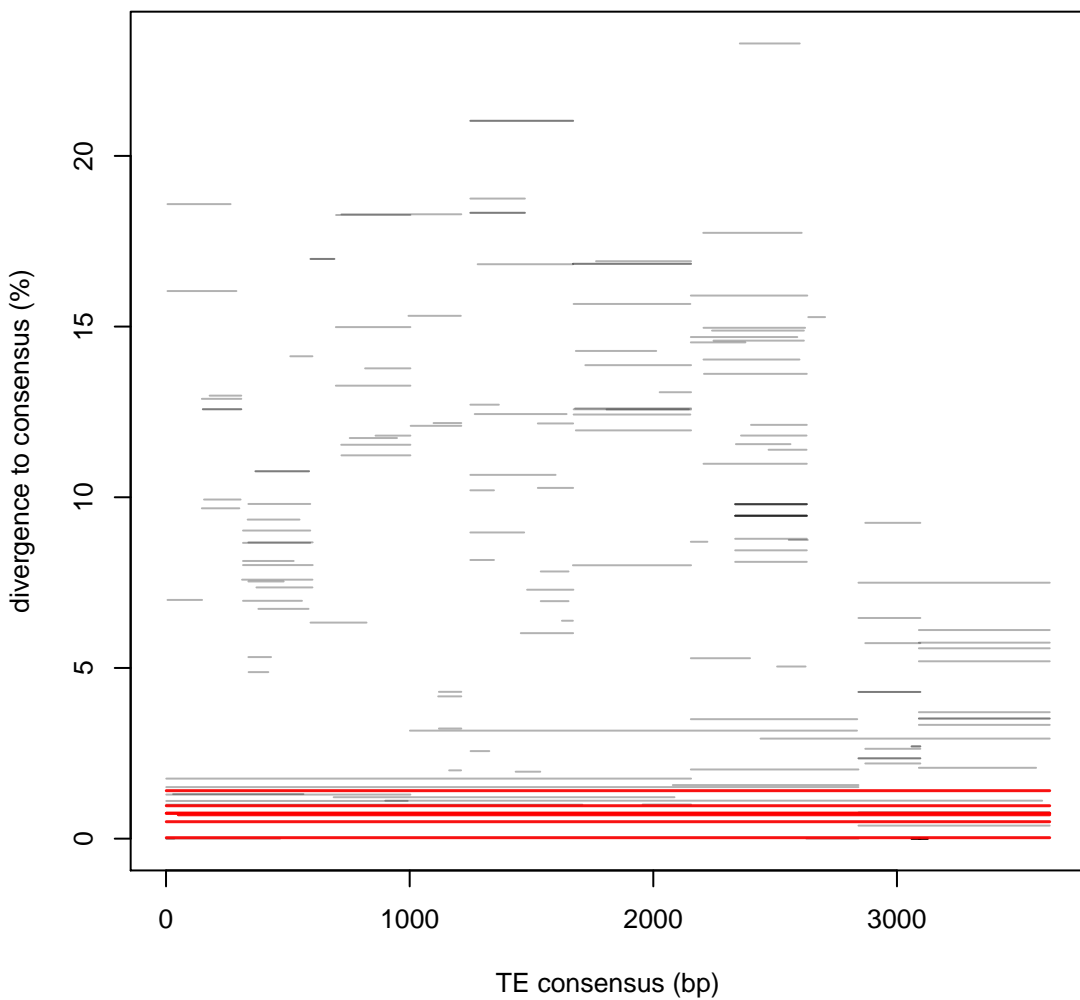
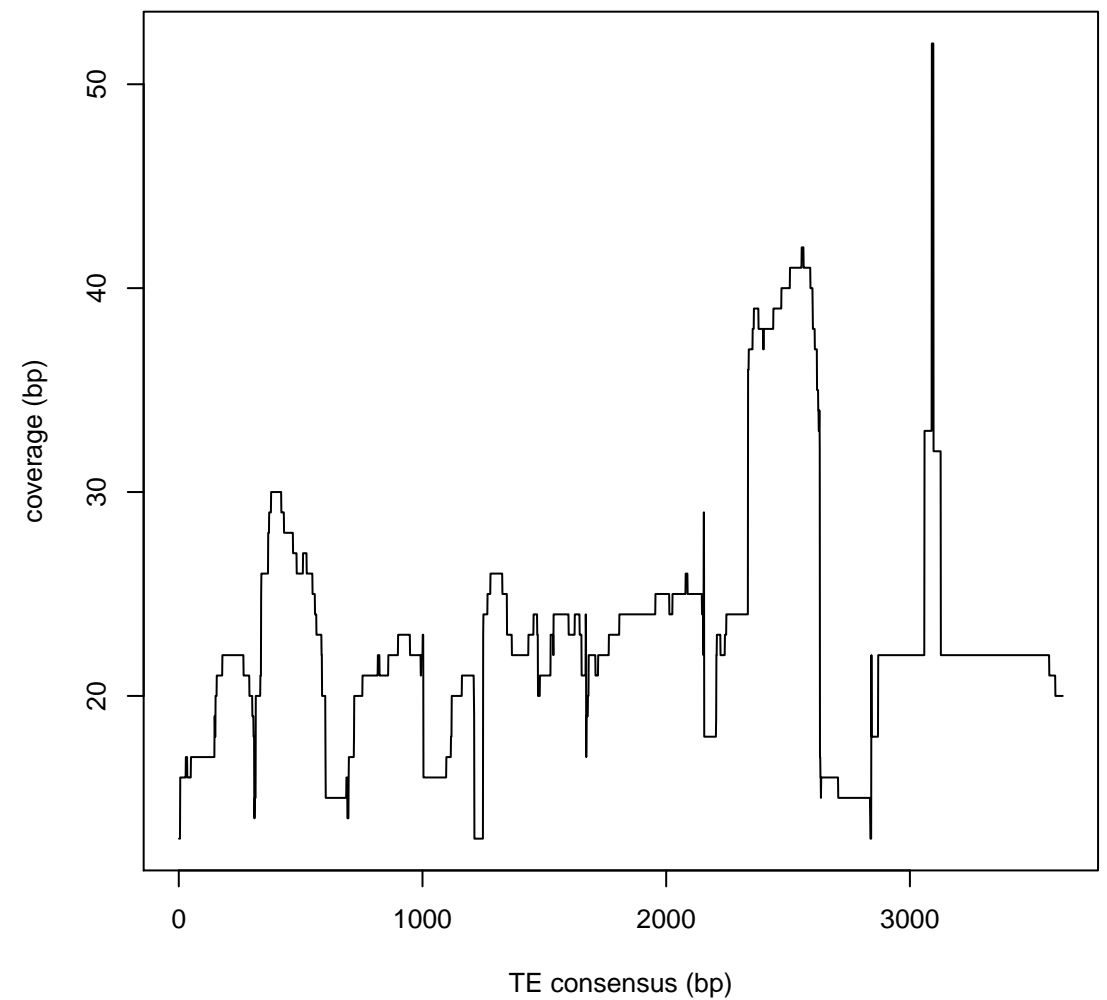


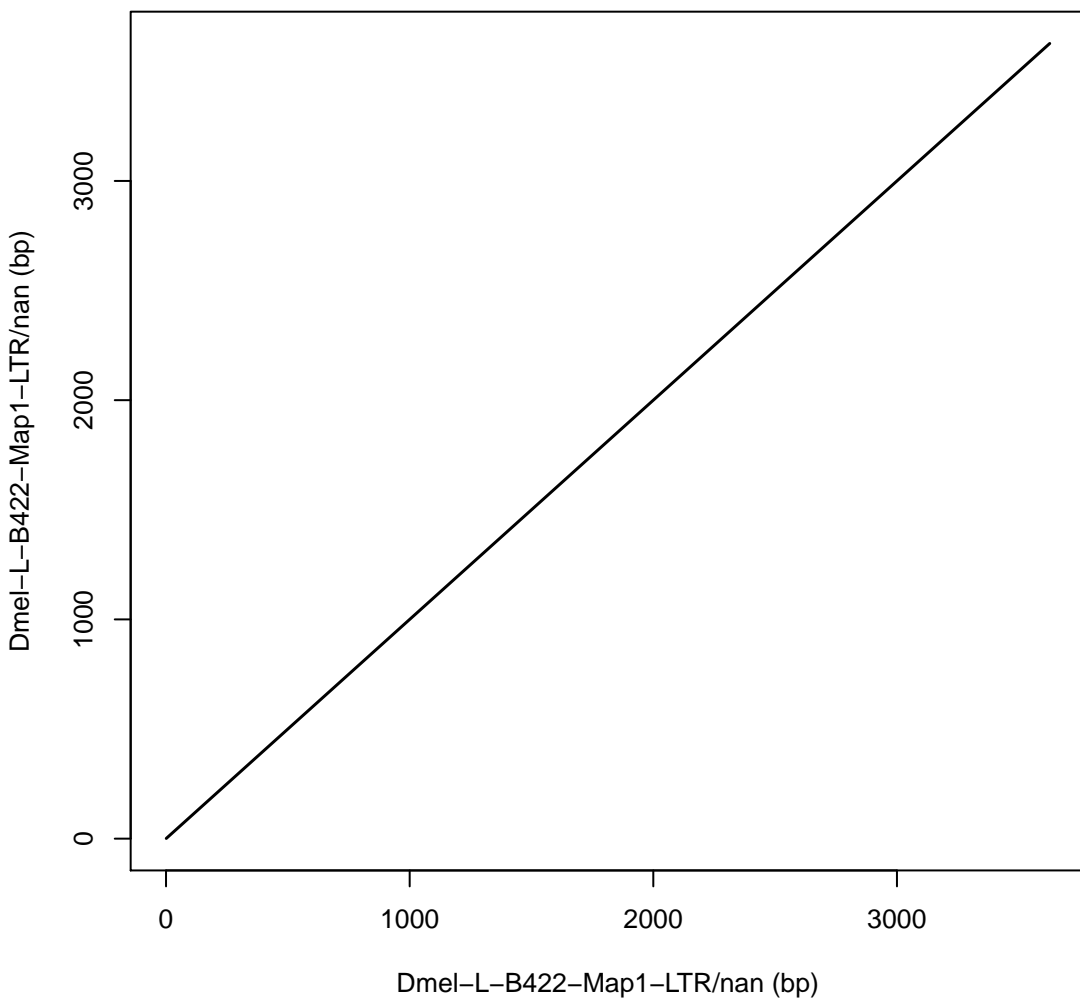
TE: Dmel-L-B422-Map1-LTR/nan  
consensus size: 3627bp; fragments: 178; full length: 7 ( $\geq 3264.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

