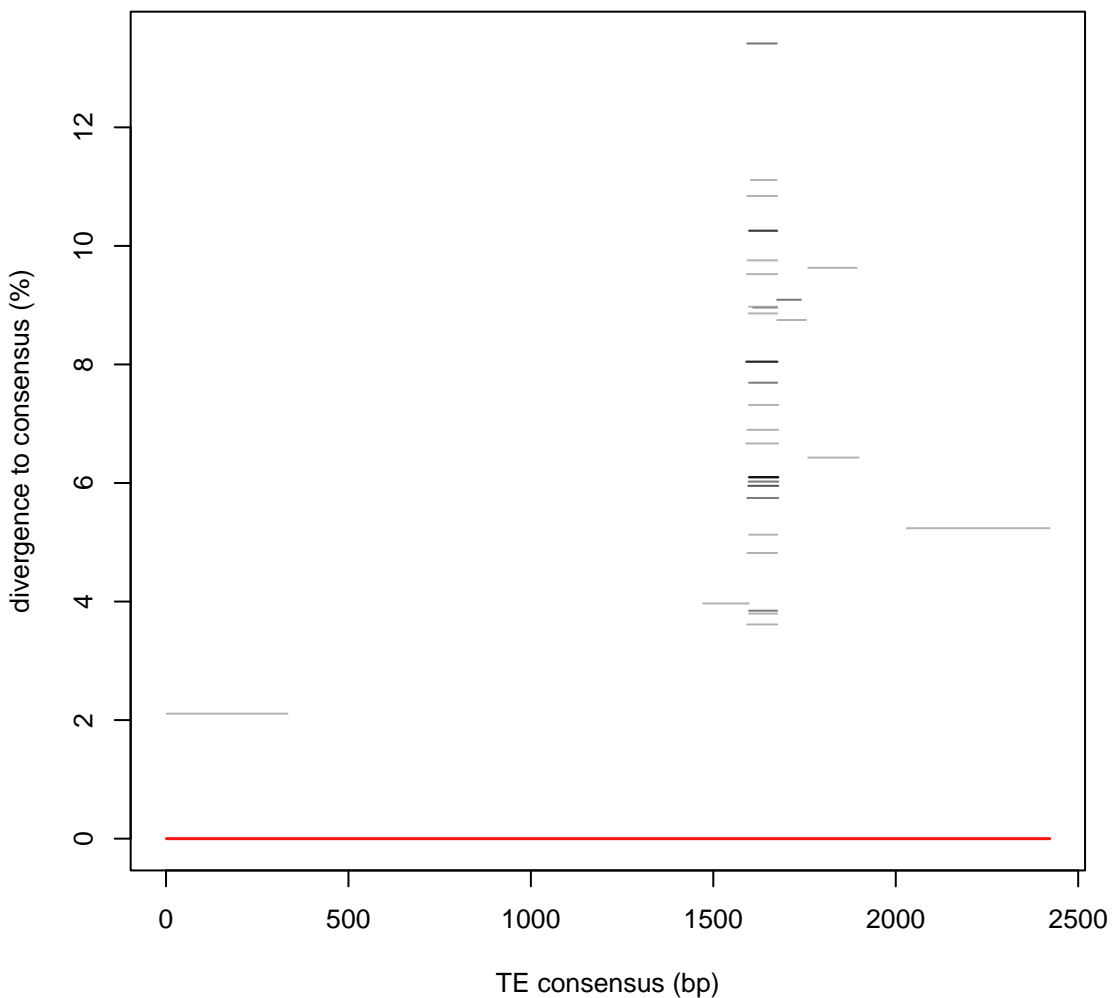
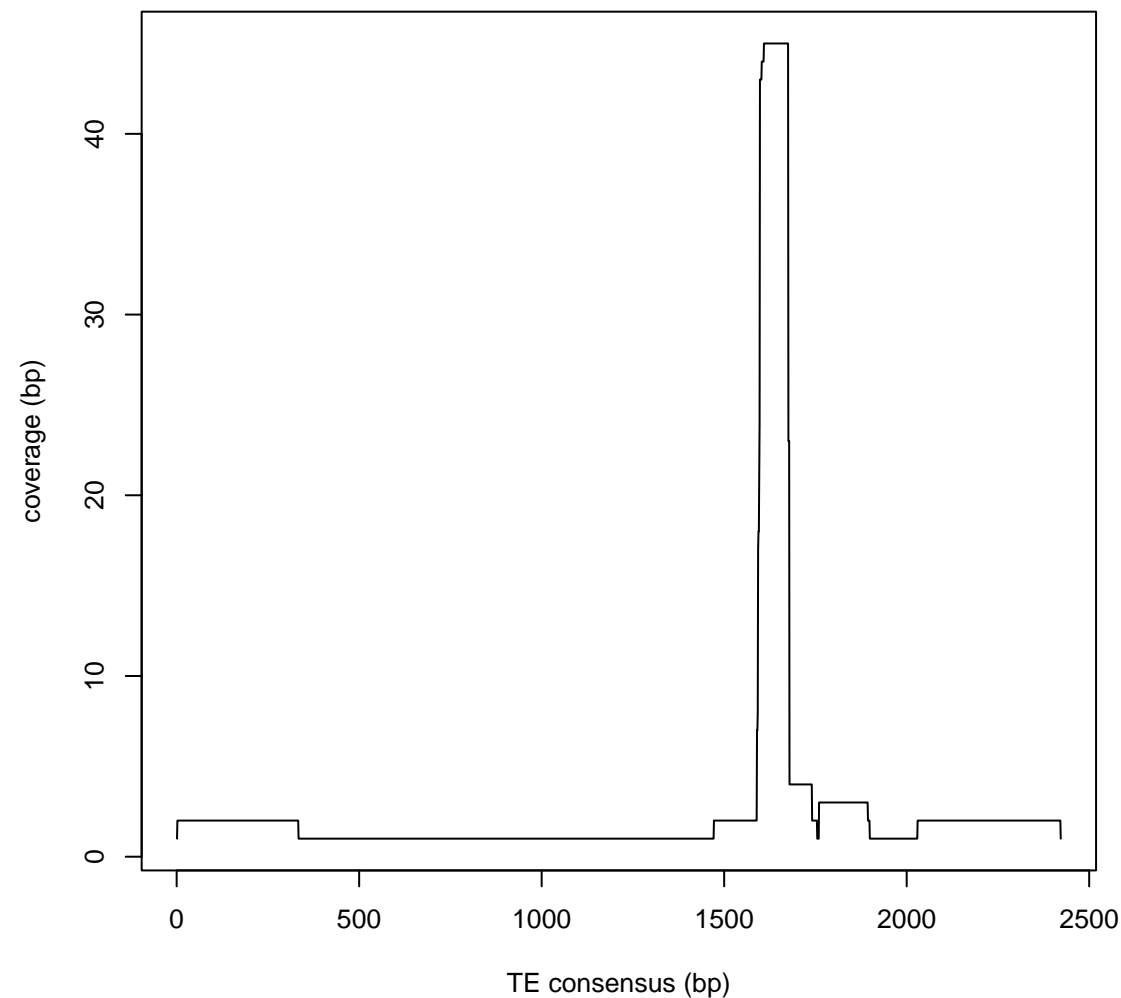


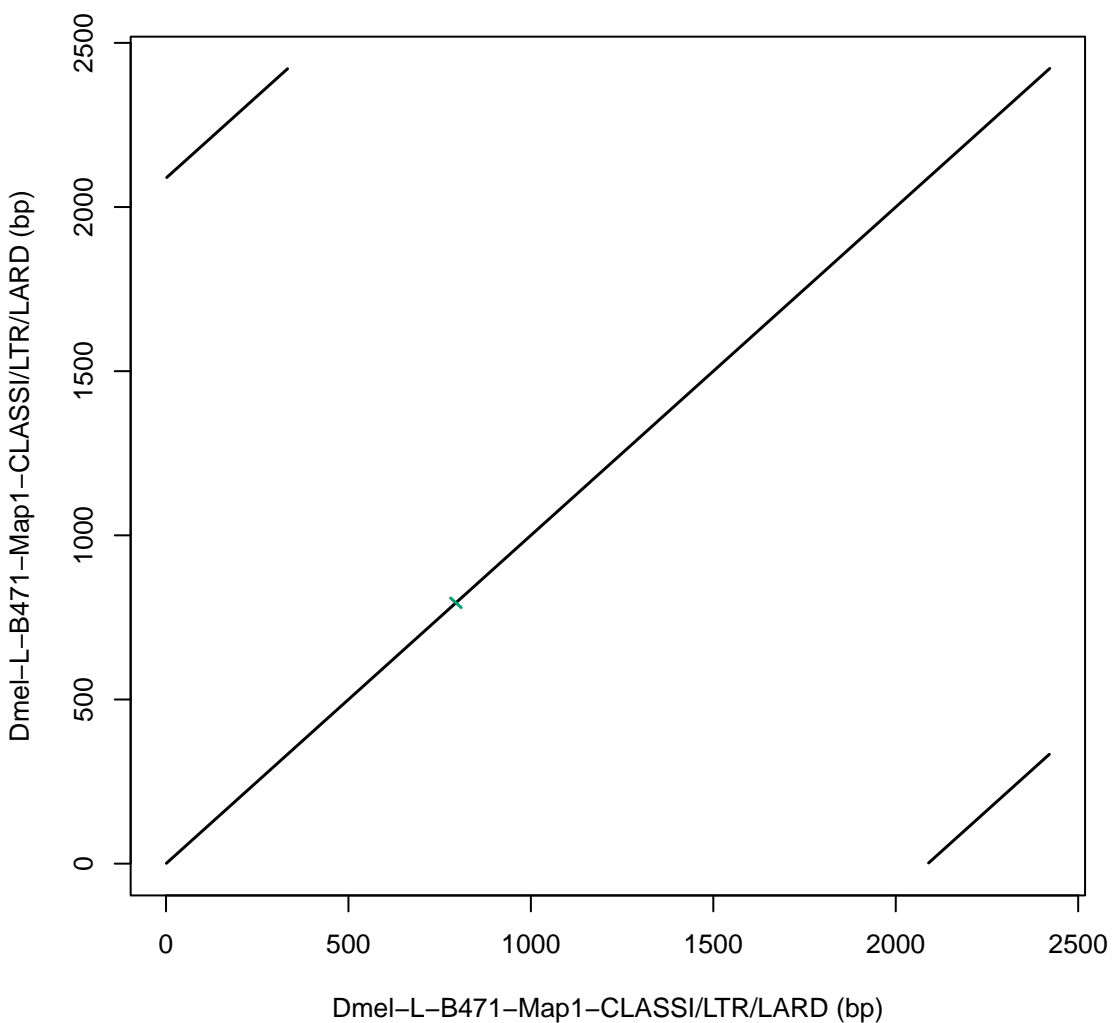
TE: Dmel-L-B471-Map1-CLASSI/LTR/LARD  
consensus size: 2422bp; fragments: 53; full length: 1 ( $\geq 2179.8\text{bp}$ )



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

