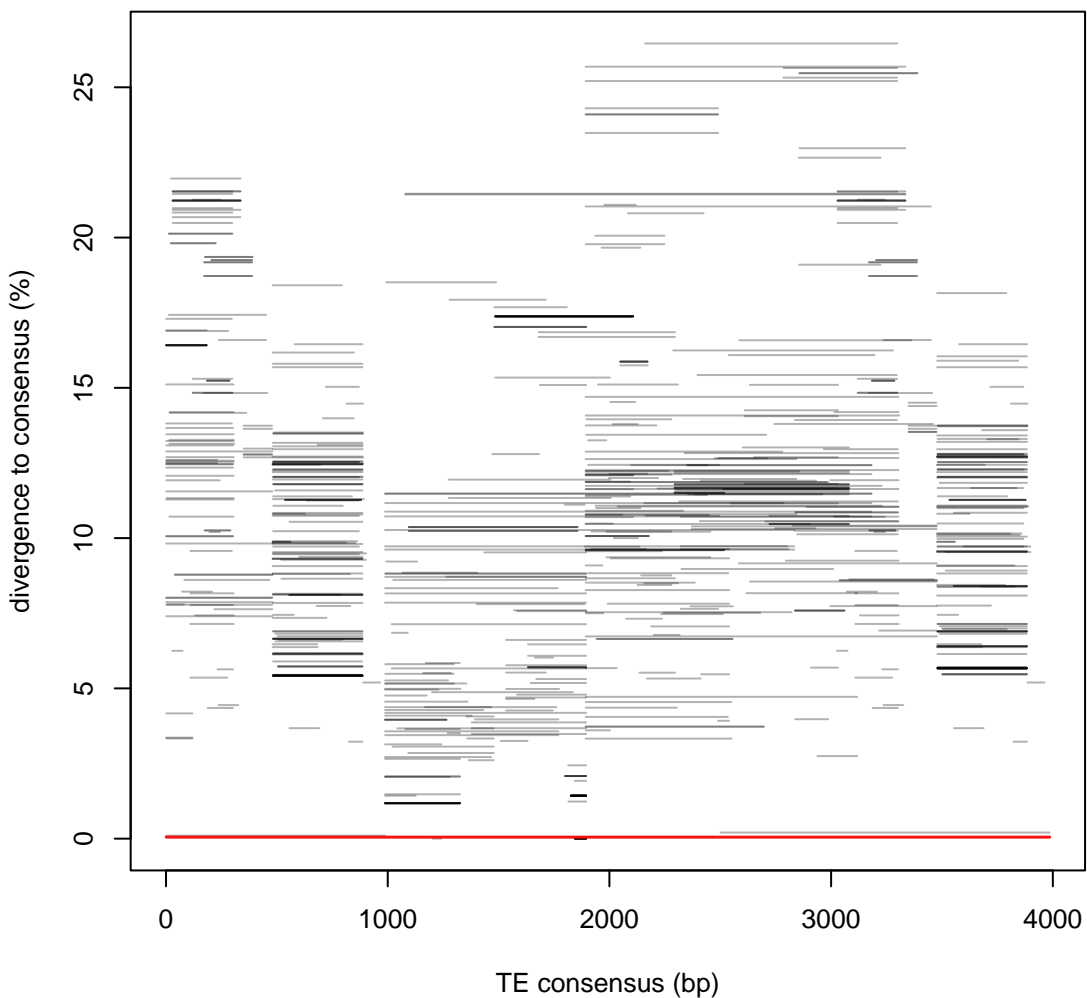
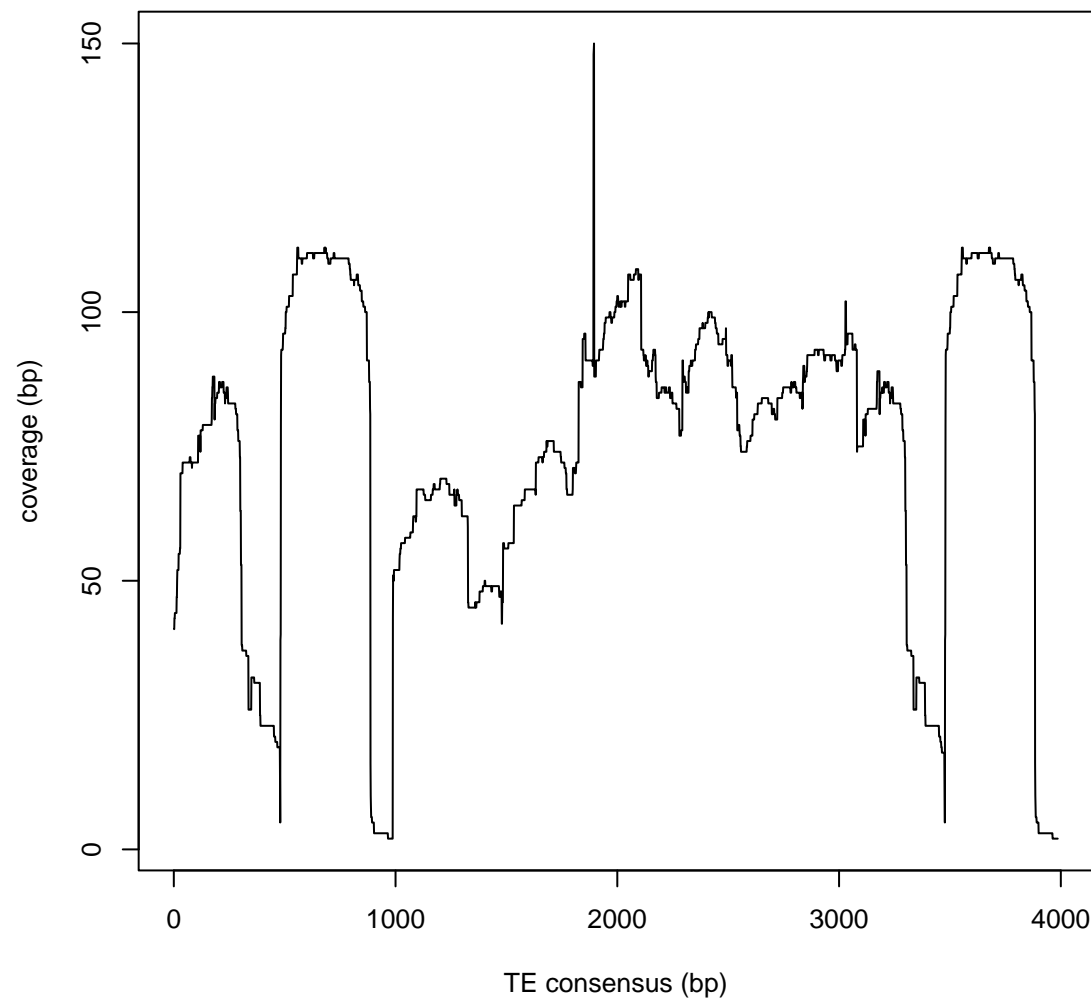


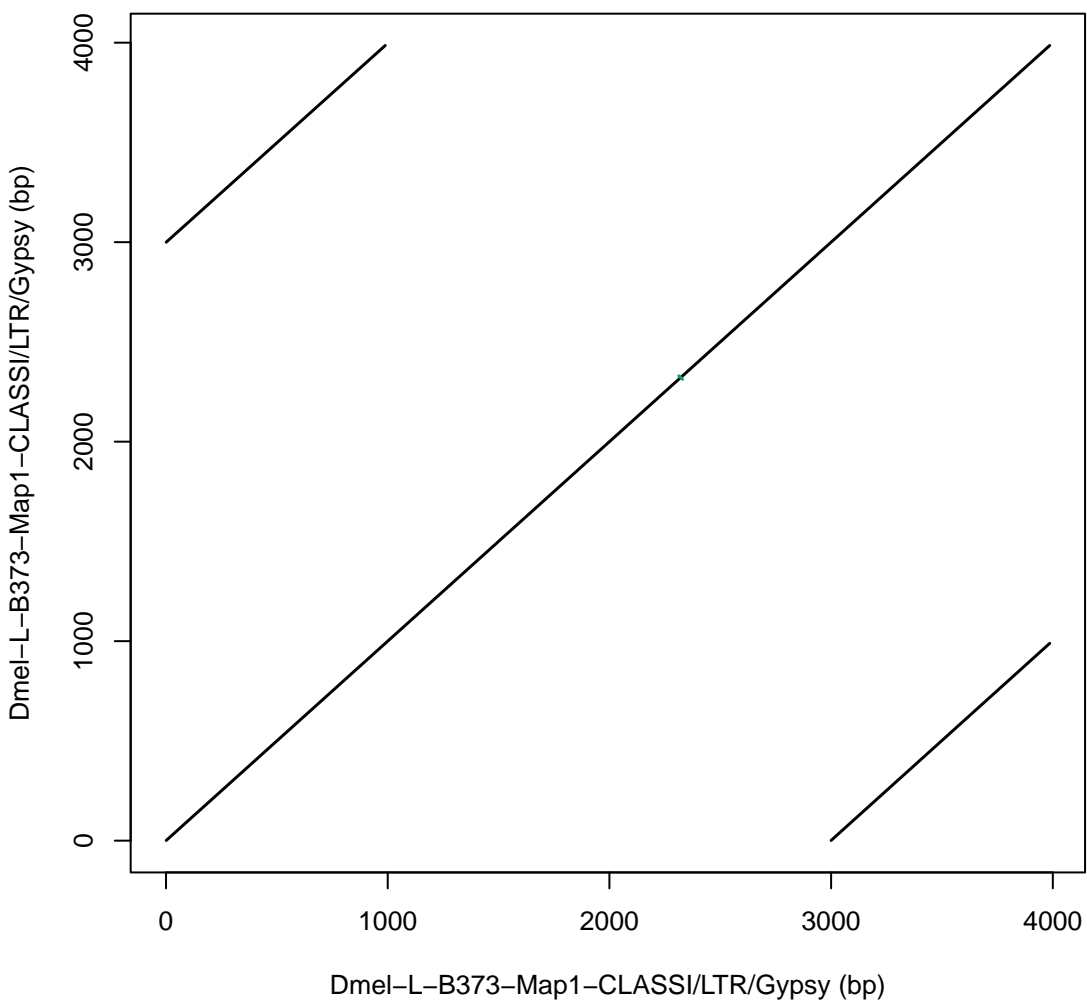
TE: Dmel-L-B373-Map1-CLASSI/LTR/Gypsy  
consensus size: 3986bp; fragments: 746; full length: 1 ( $\geq 3587.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

