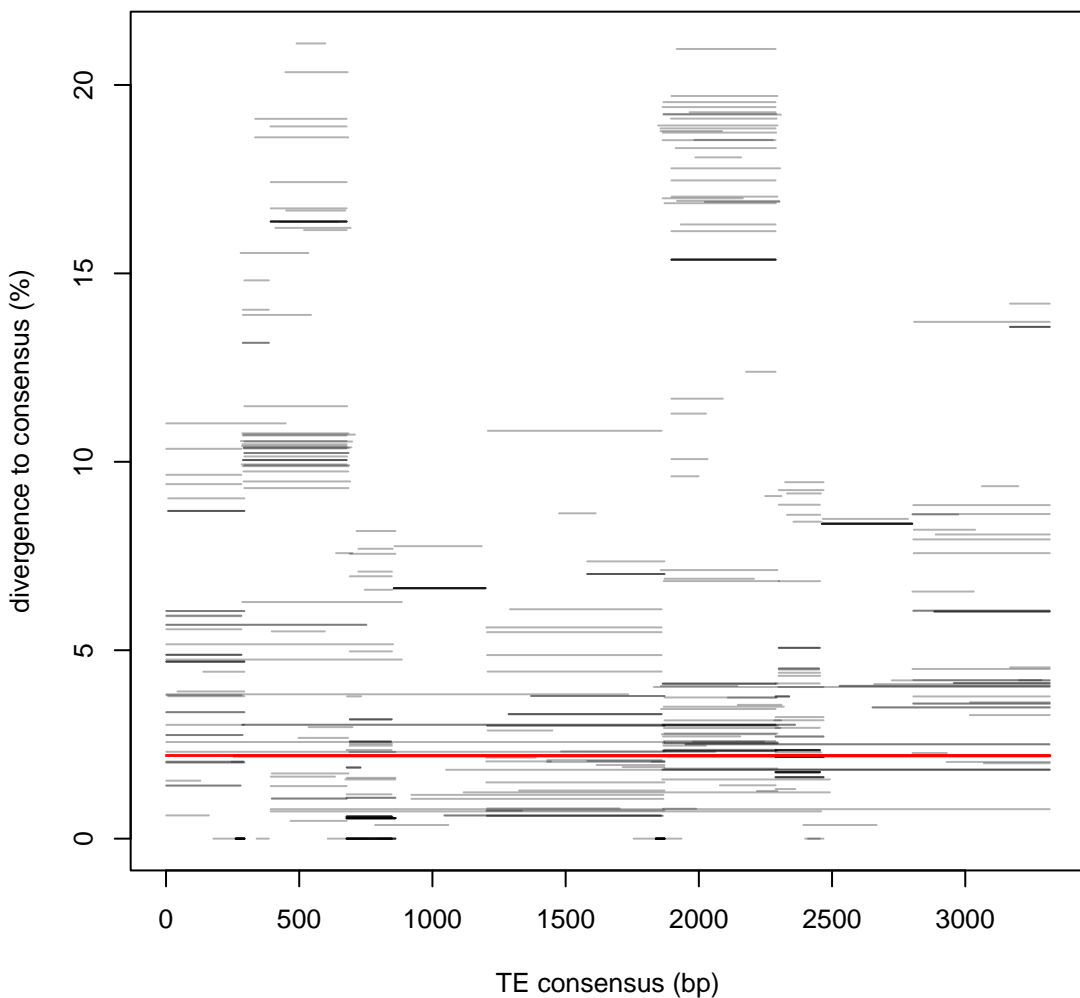
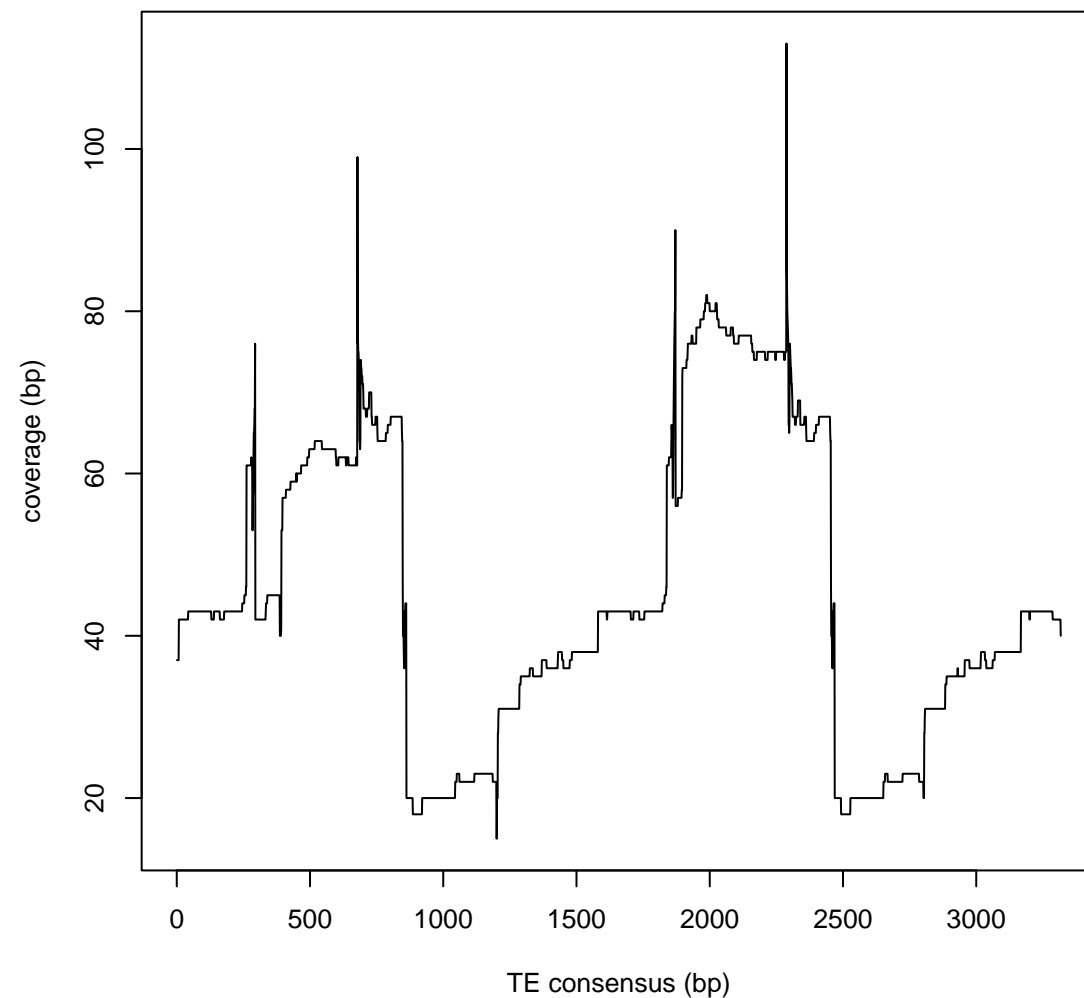


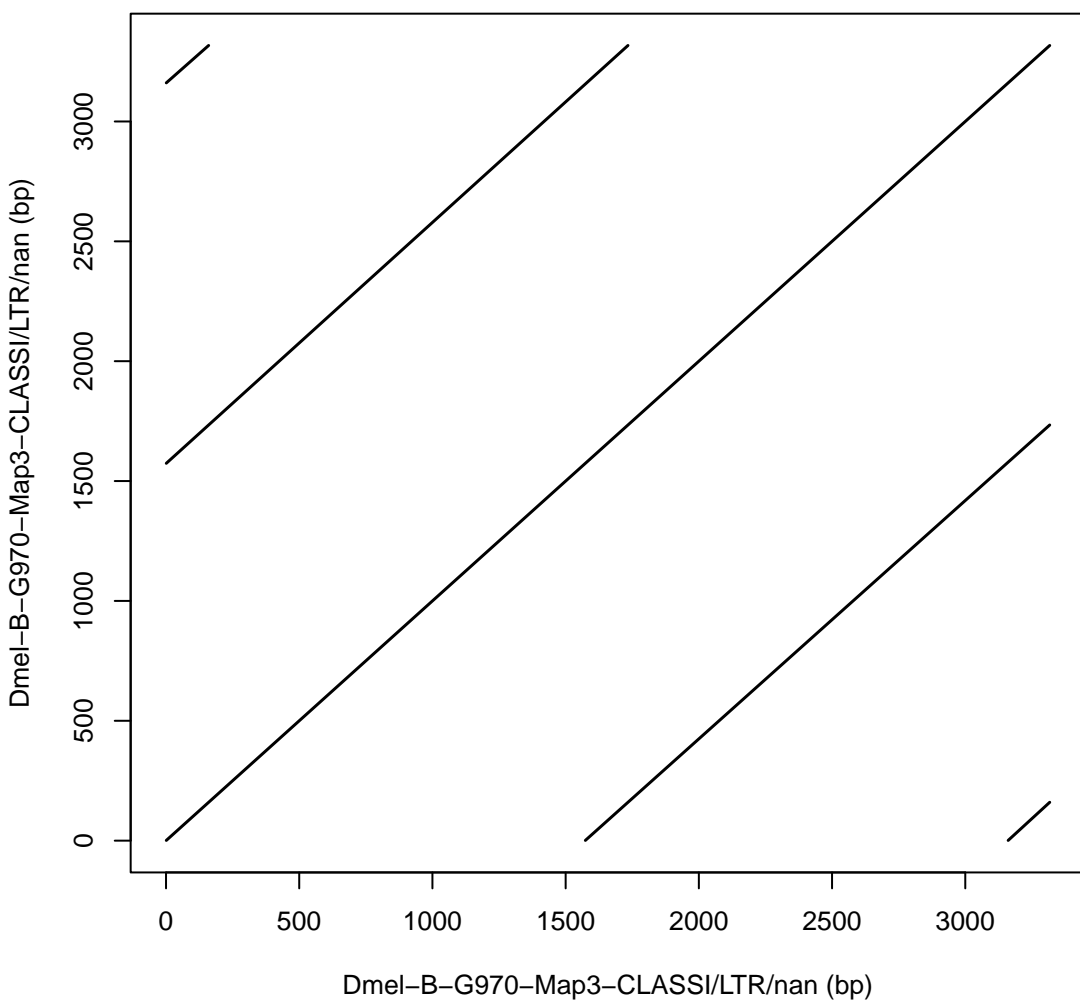
TE: Dmel-B-G970-Map3-CLASSI/LTR/nan  
consensus size: 3317bp; fragments: 439; full length: 2 ( $\geq 2985.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

