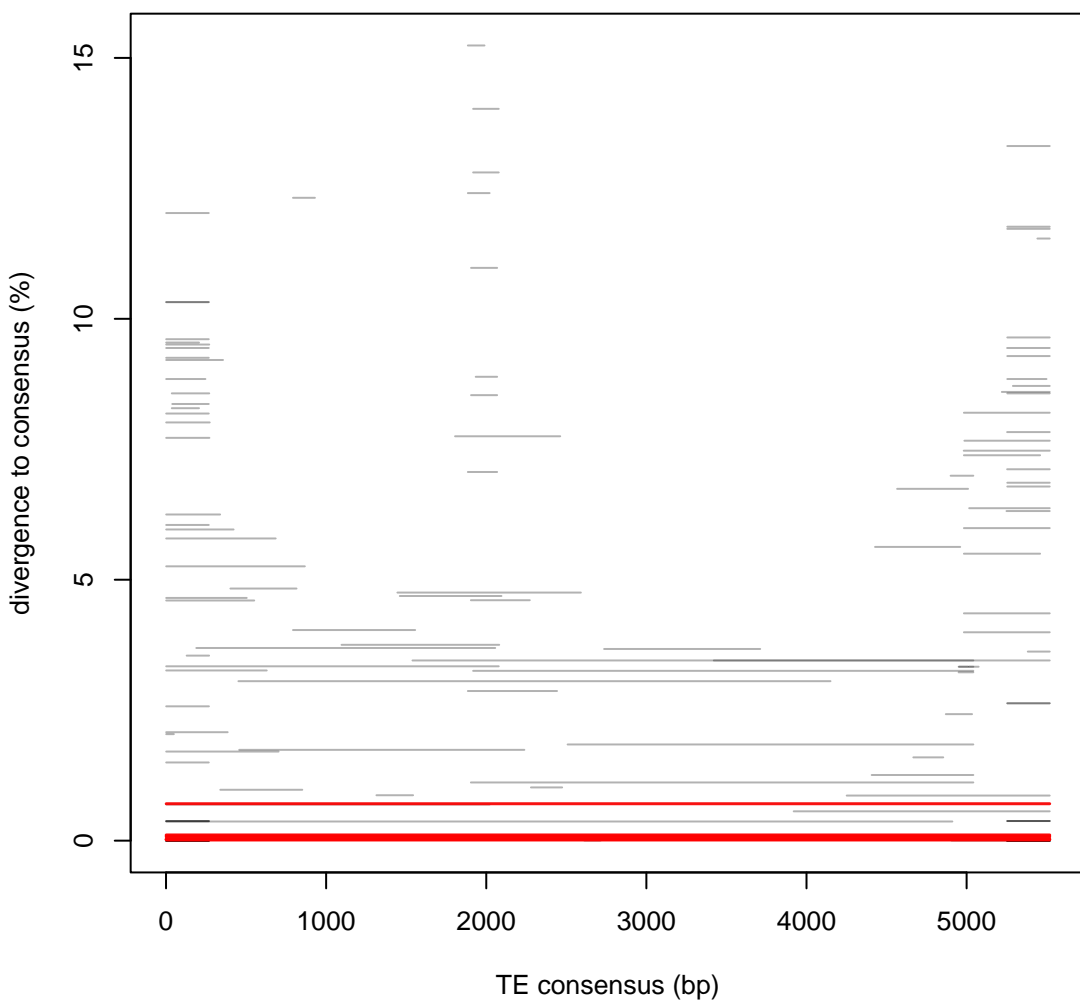
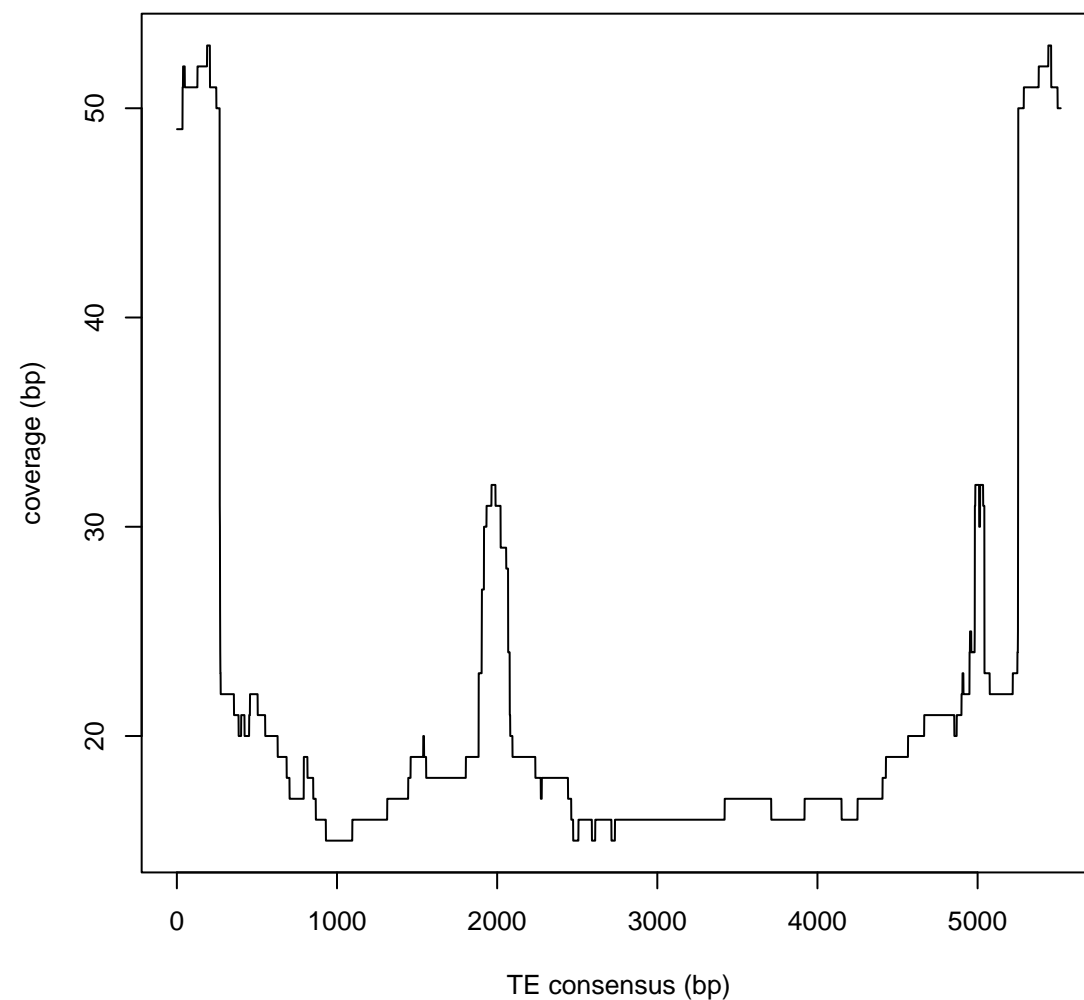


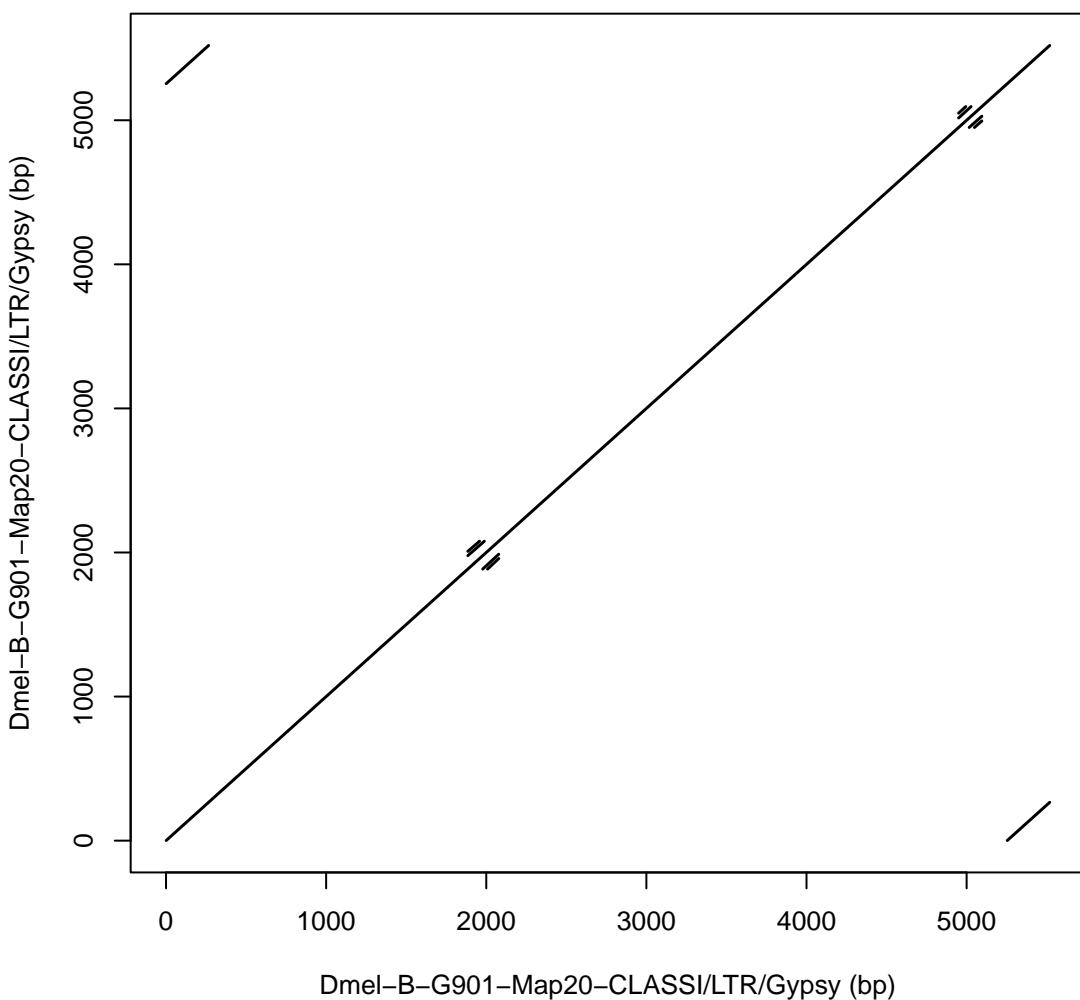
TE: Dmel-B-G901-Map20-CLASSI/LTR/Gypsy  
consensus size: 5519bp; fragments: 137; full length: 8 ( $\geq 4967.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

