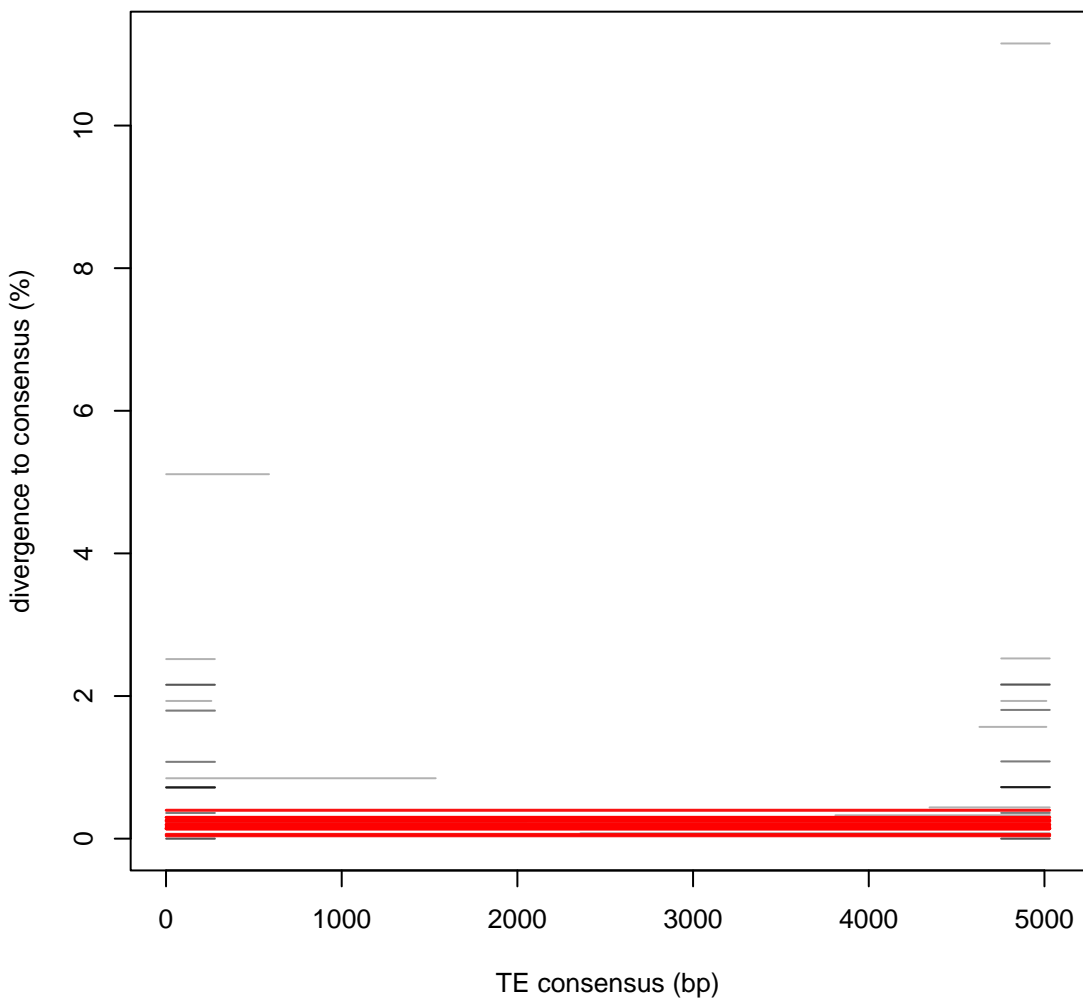
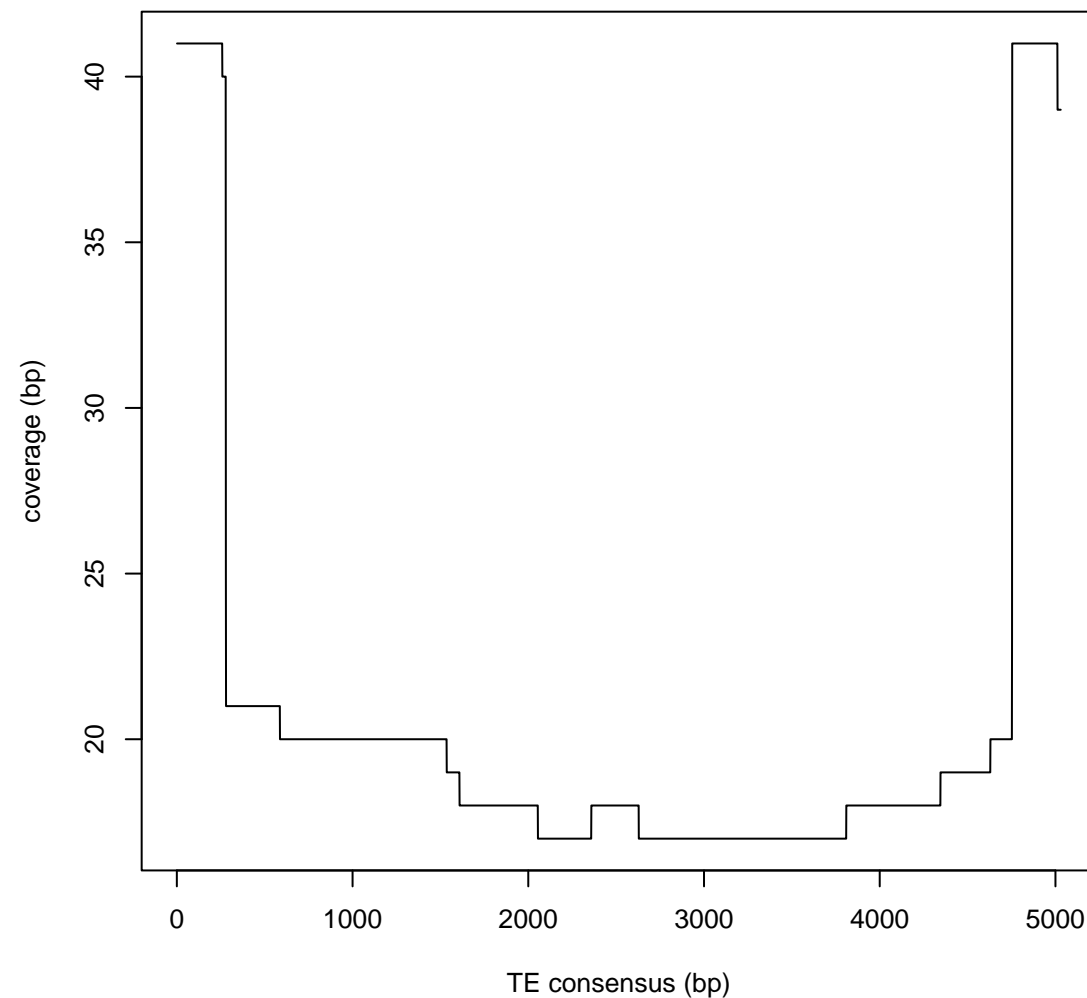


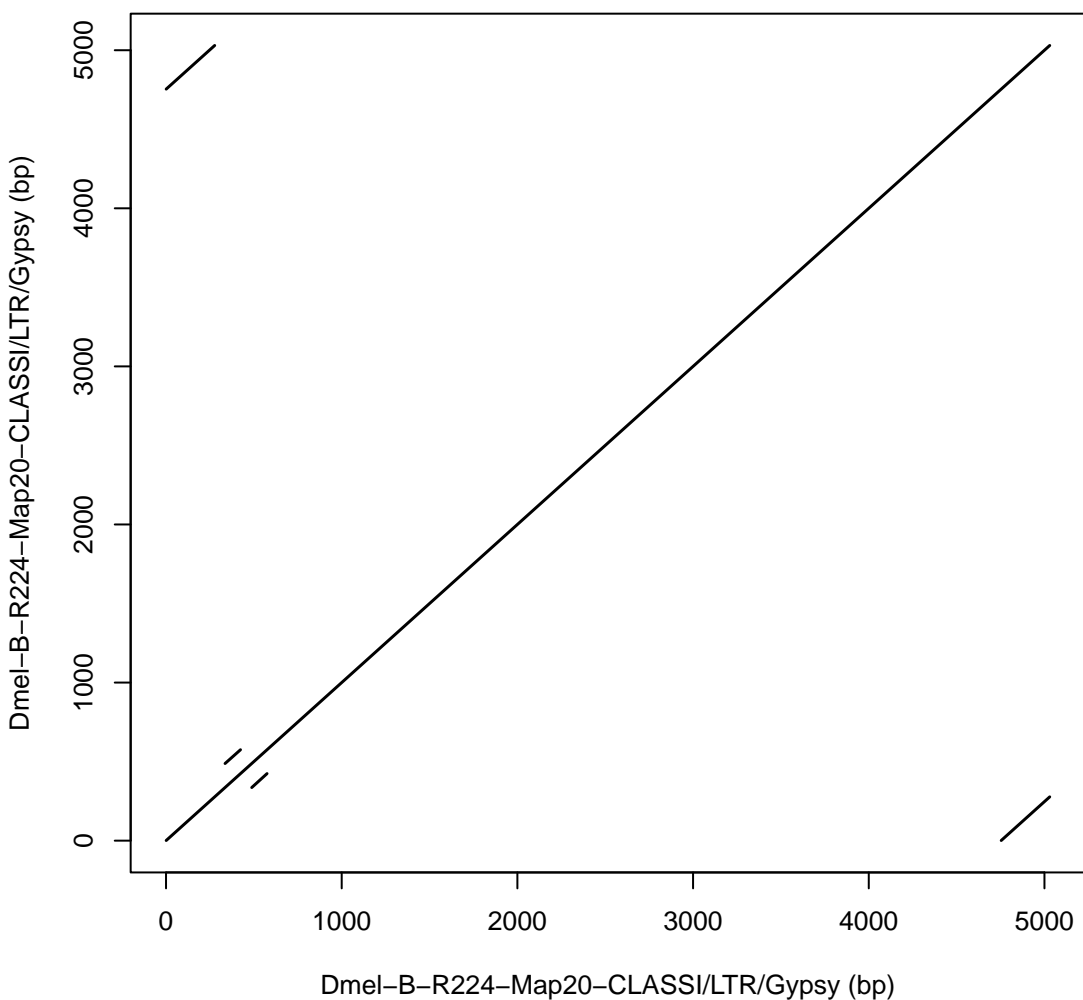
TE: Dmel-B-R224-Map20-CLASSI/LTR/Gypsy  
consensus size: 5030bp; fragments: 66; full length: 16 ( $\geq 4527$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

