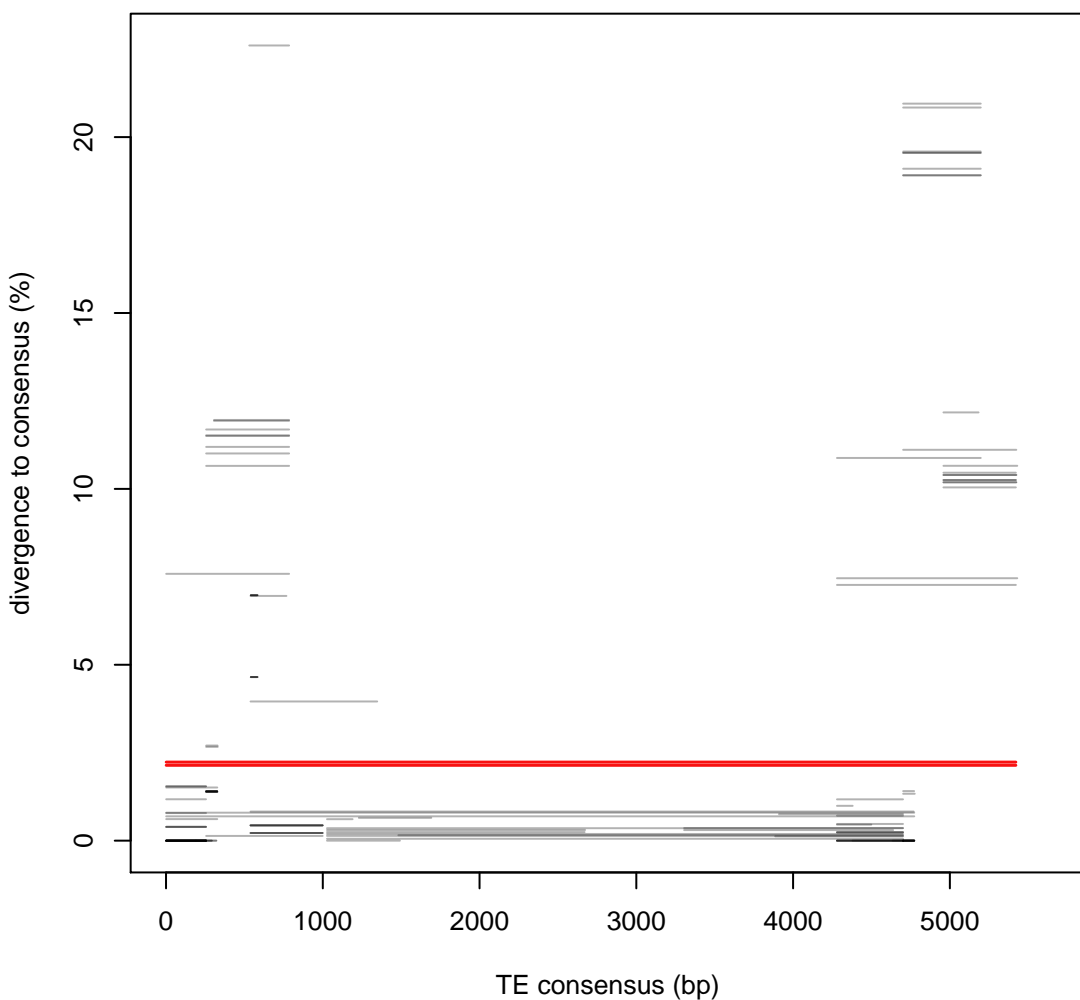
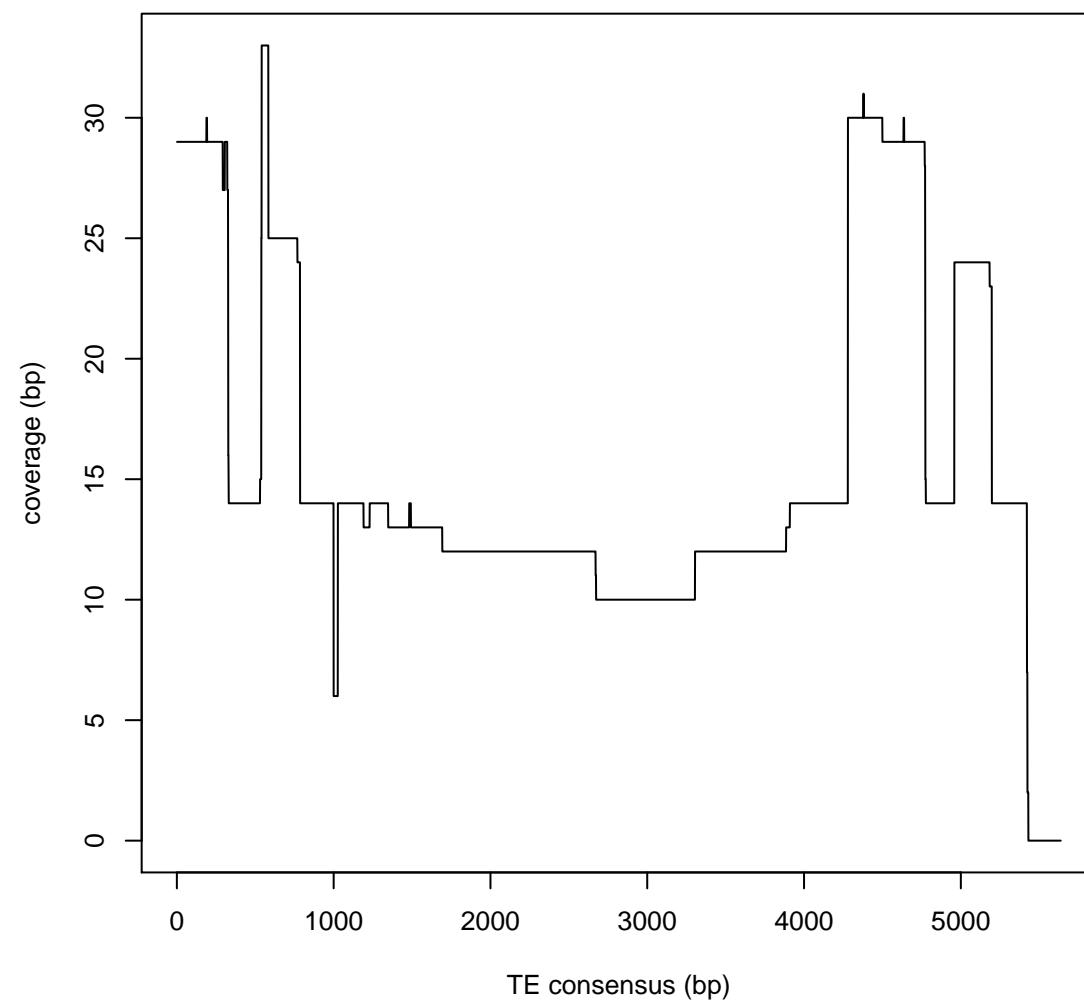


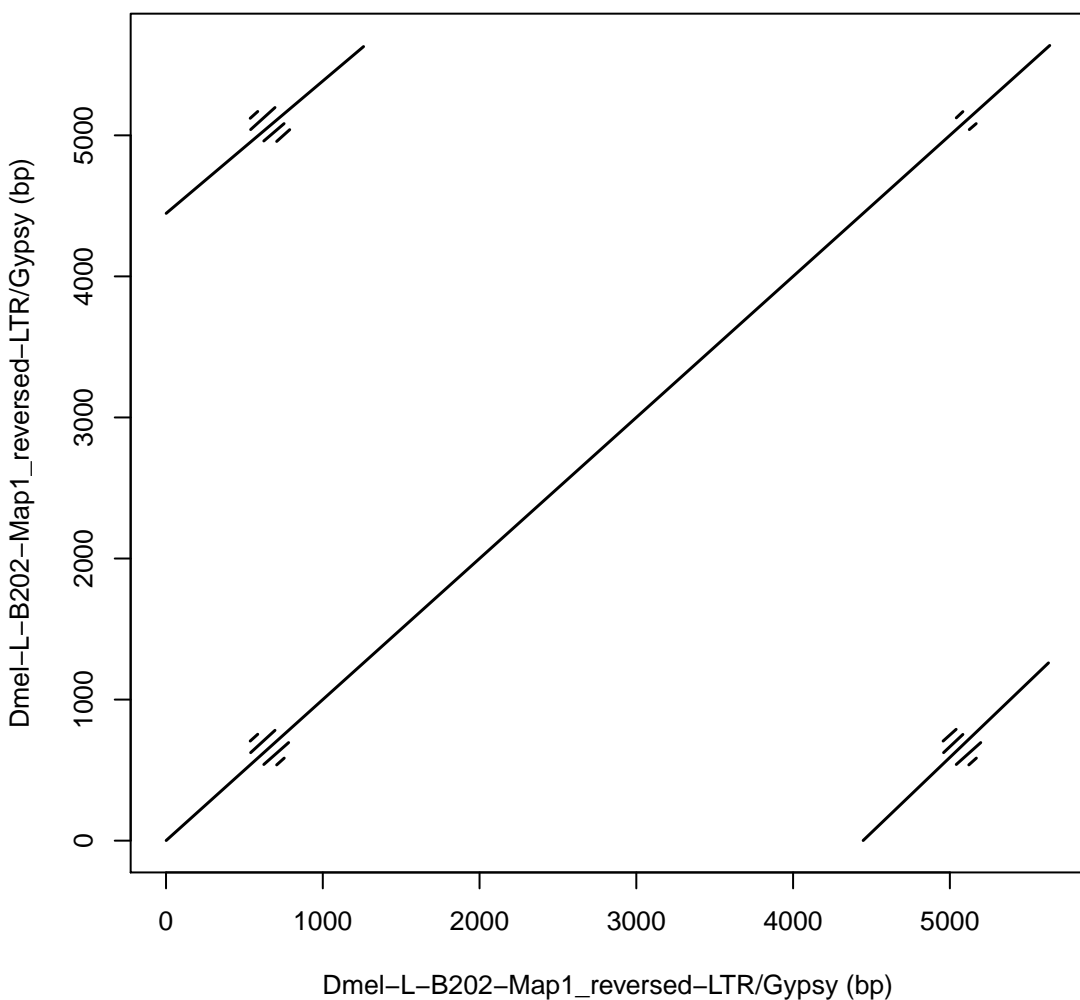
TE: Dmel-L-B202-Map1\_reversed-LTR/Gypsy  
consensus size: 5637bp; fragments: 135; full length: 2 ( $\geq 5073.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

