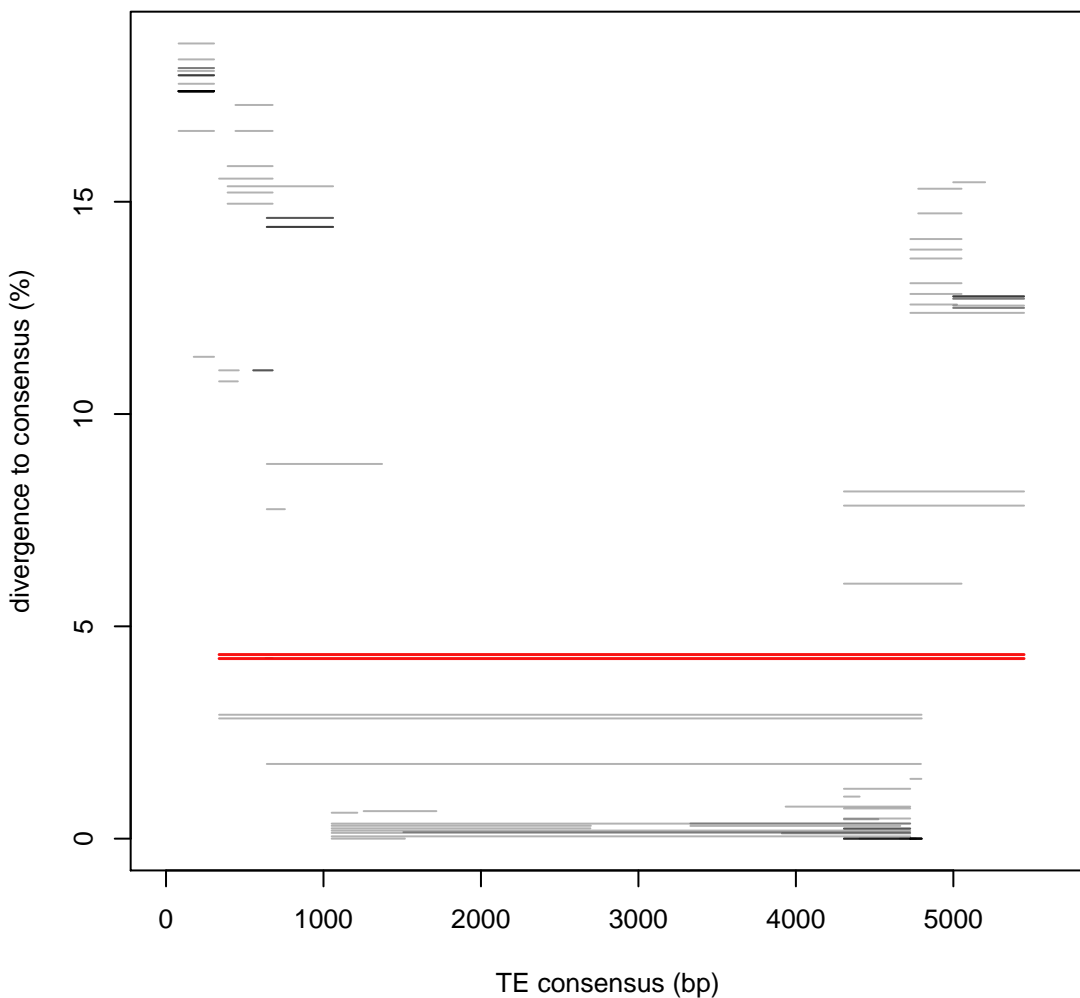
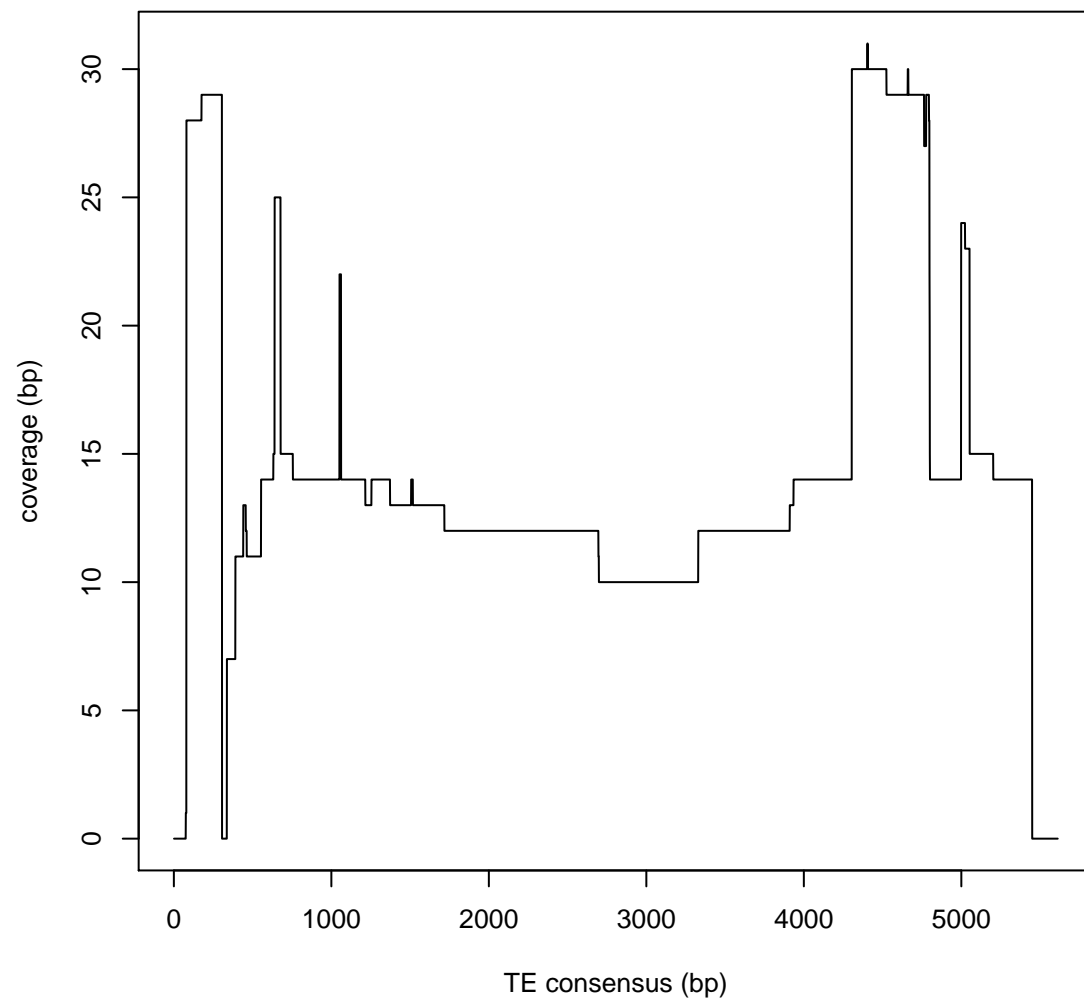


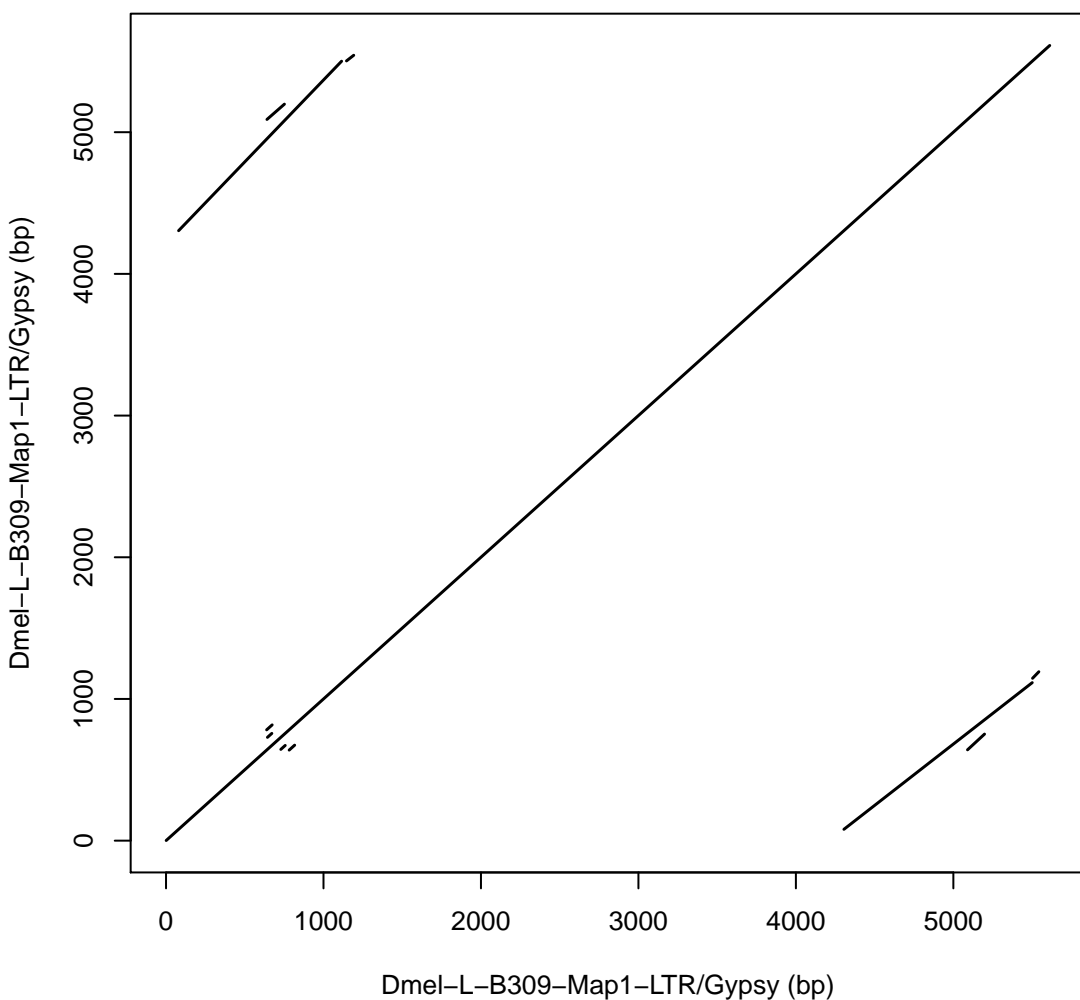
TE: Dmel-L-B309-Map1-LTR/Gypsy  
consensus size: 5612bp; fragments: 121; full length: 2 ( $\geq 5050.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

