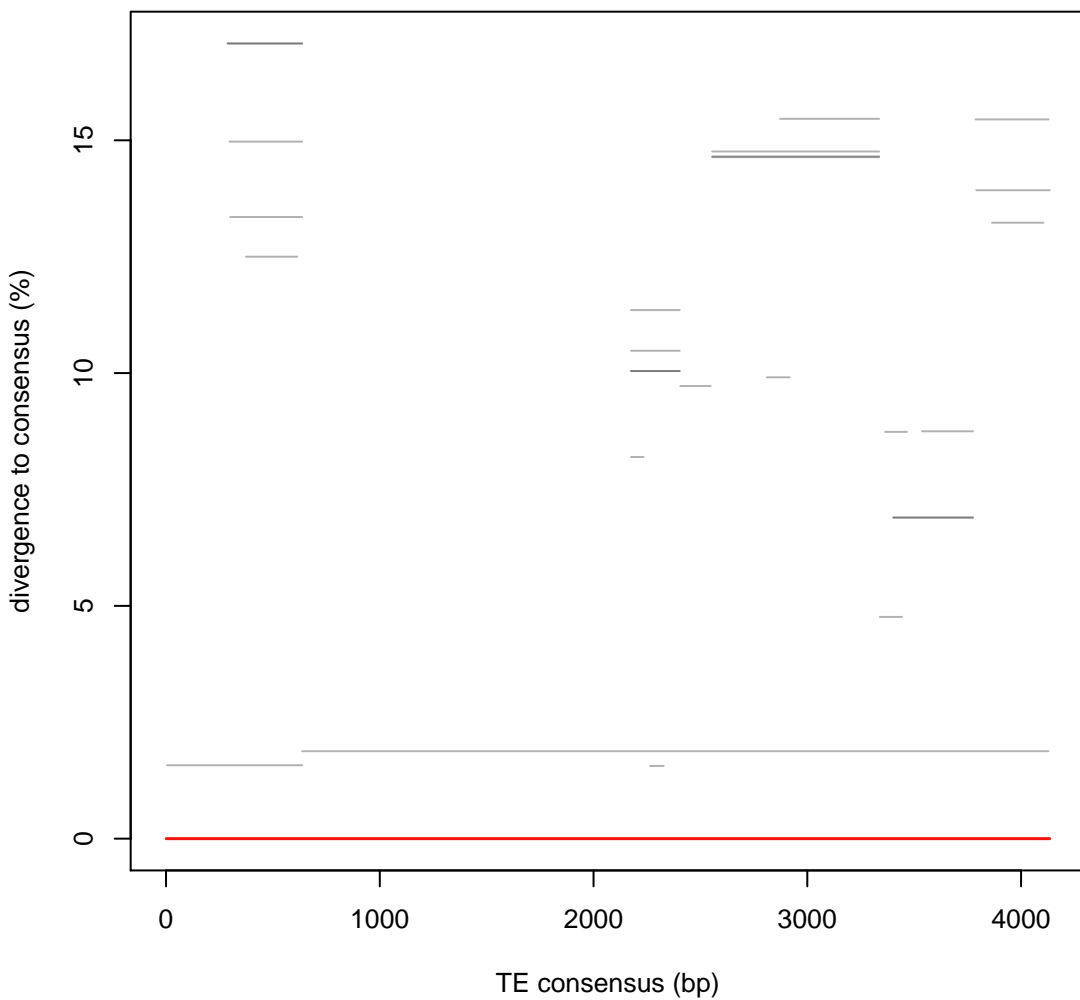
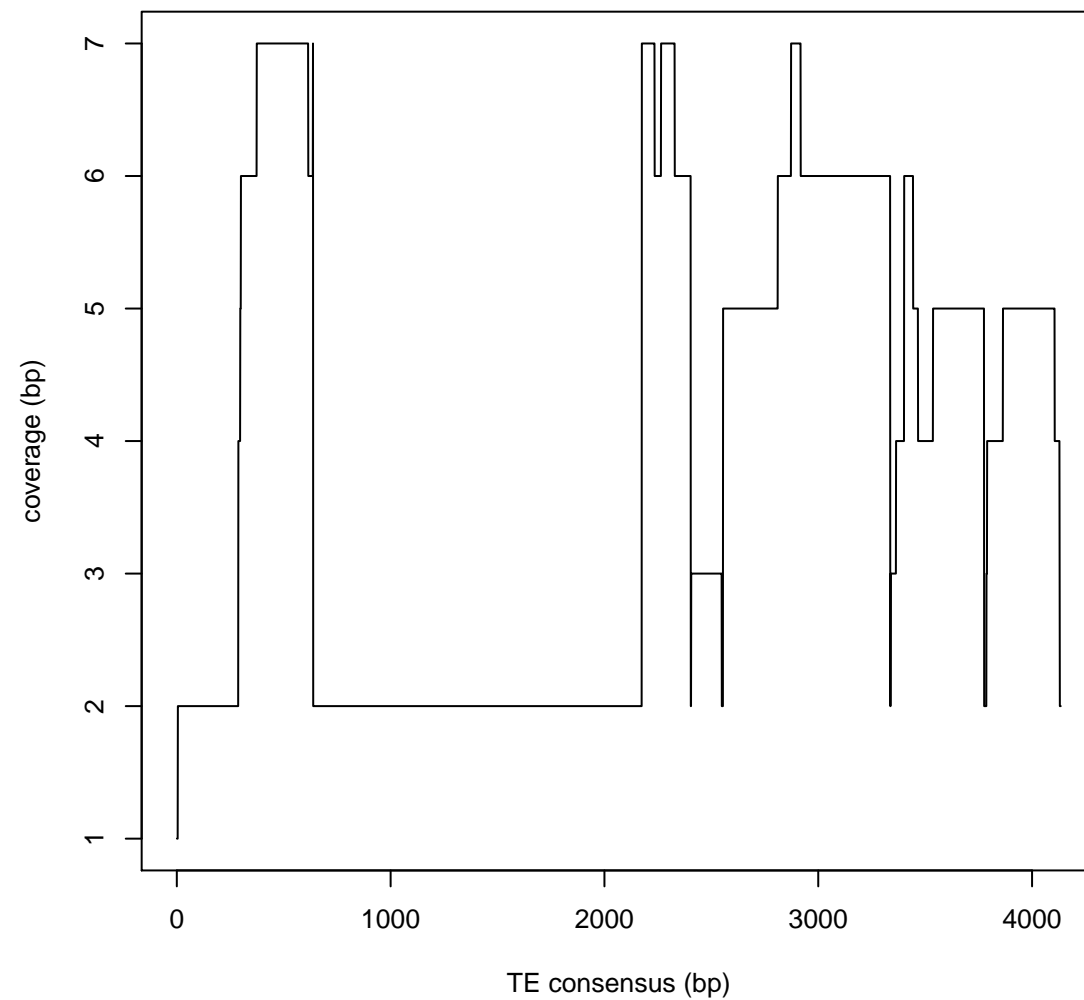


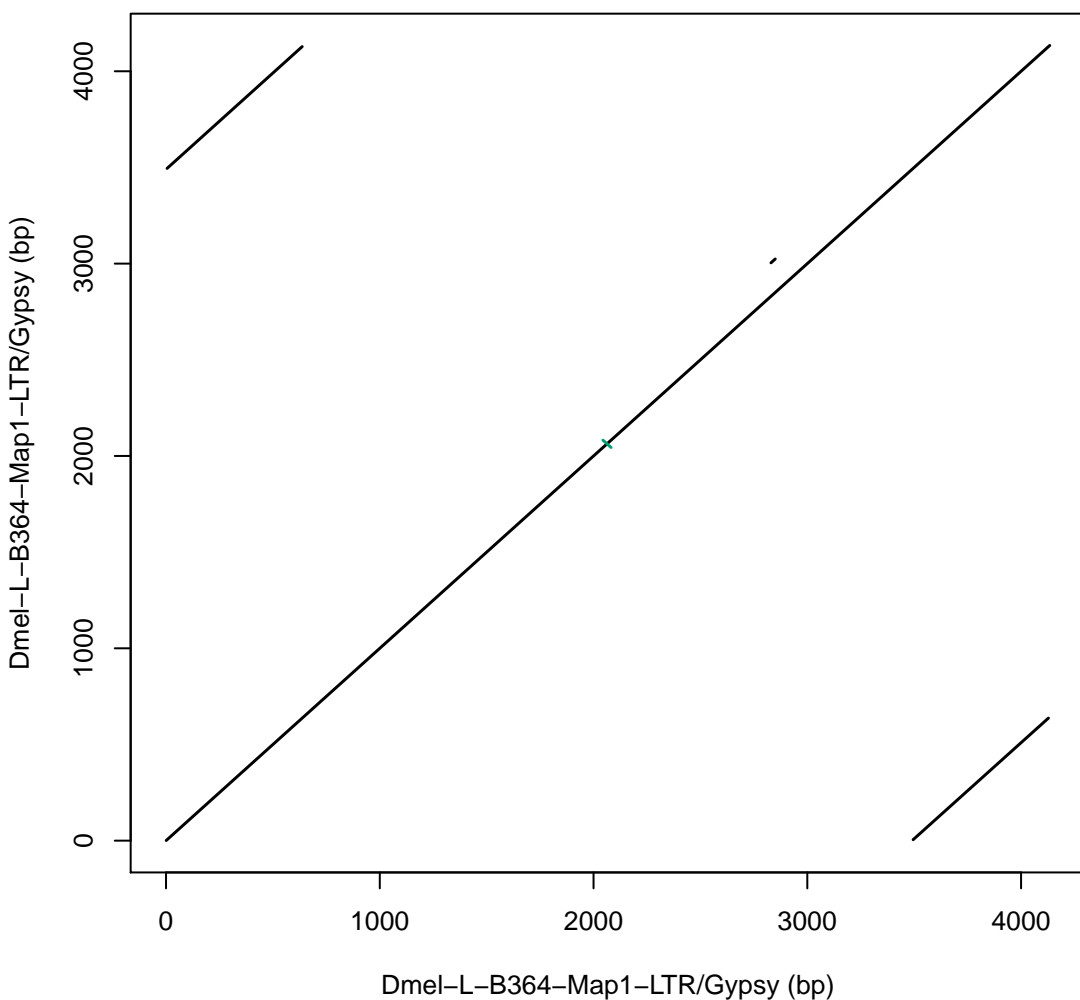
TE: Dmel-L-B364-Map1-LTR/Gypsy  
consensus size: 4134bp; fragments: 28; full length: 1 ( $\geq 3720.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

