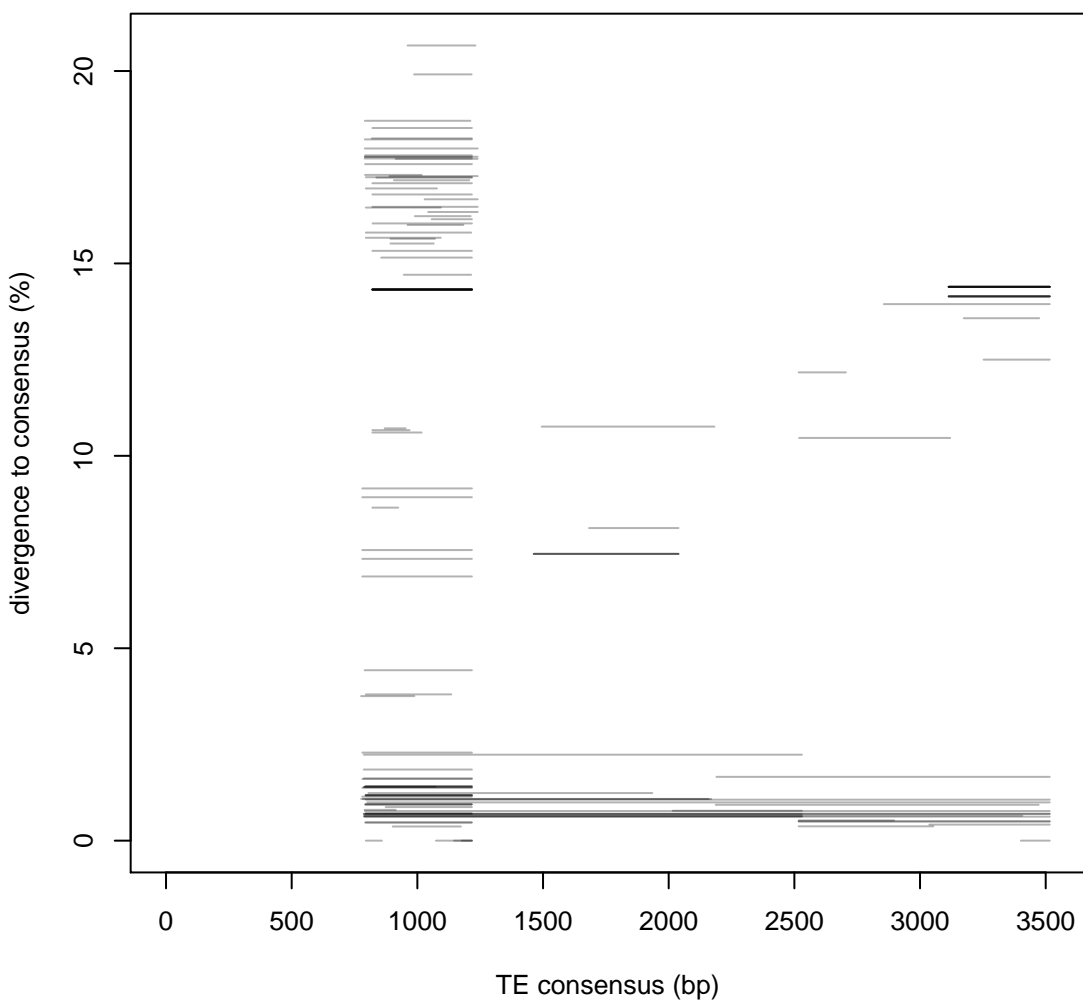
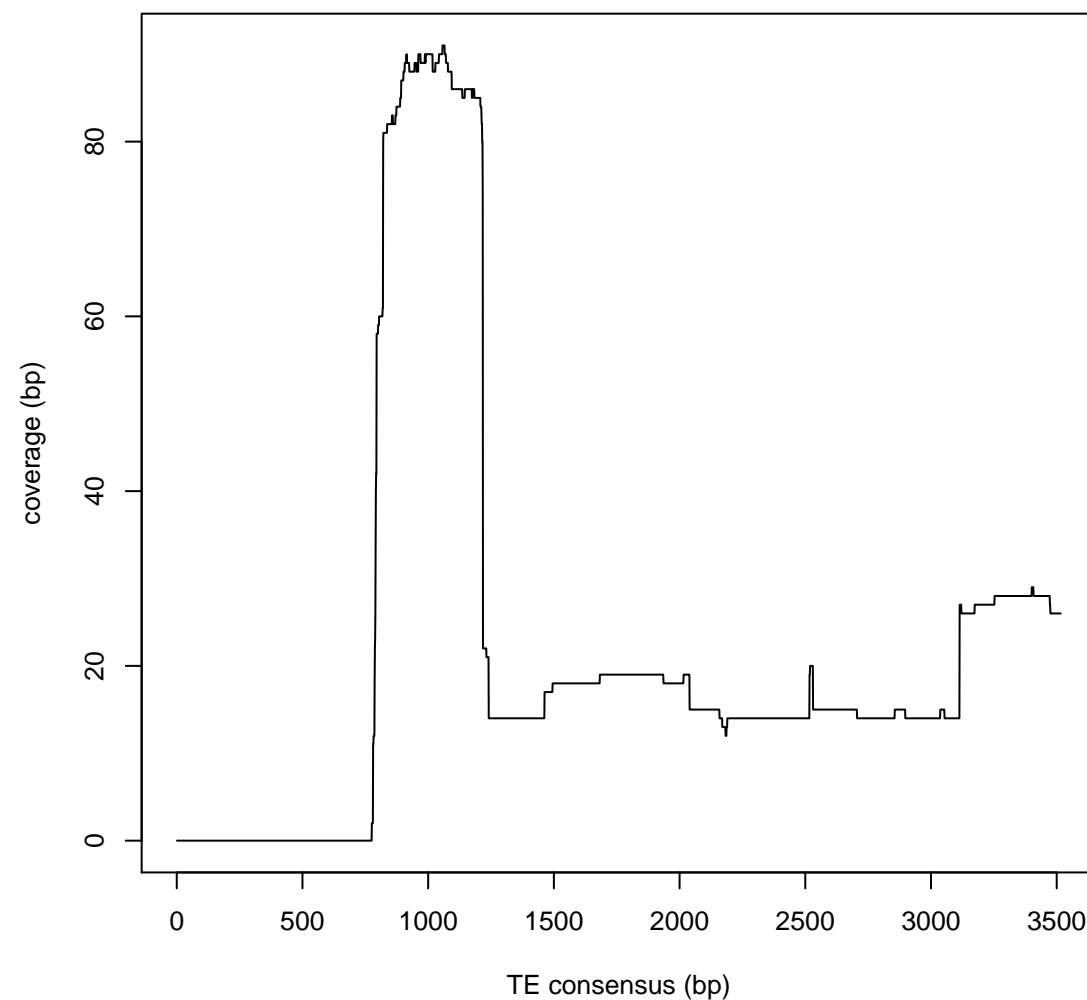


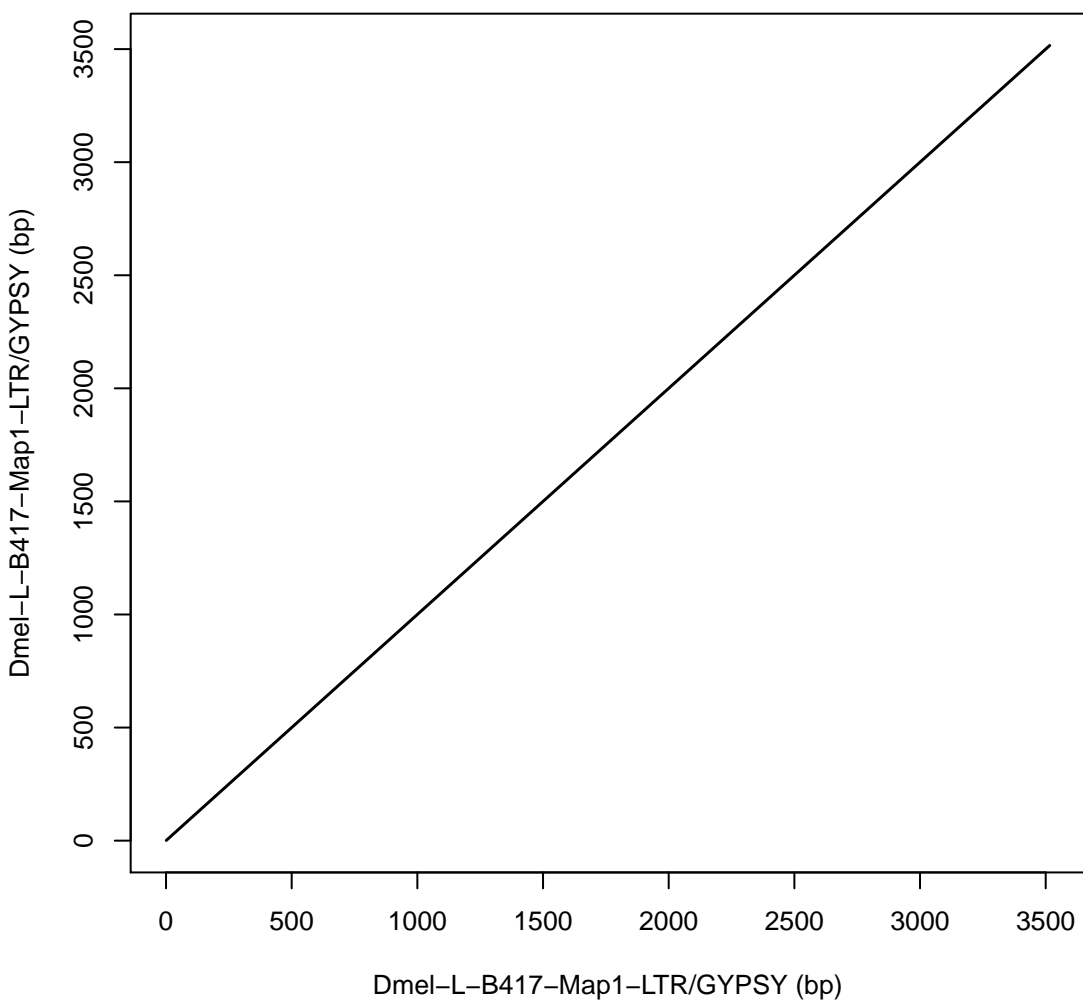
TE: Dmel-L-B417-Map1-LTR/GYPsy  
consensus size: 3516bp; fragments: 134; full length: 0 ( $\geq 3164.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

