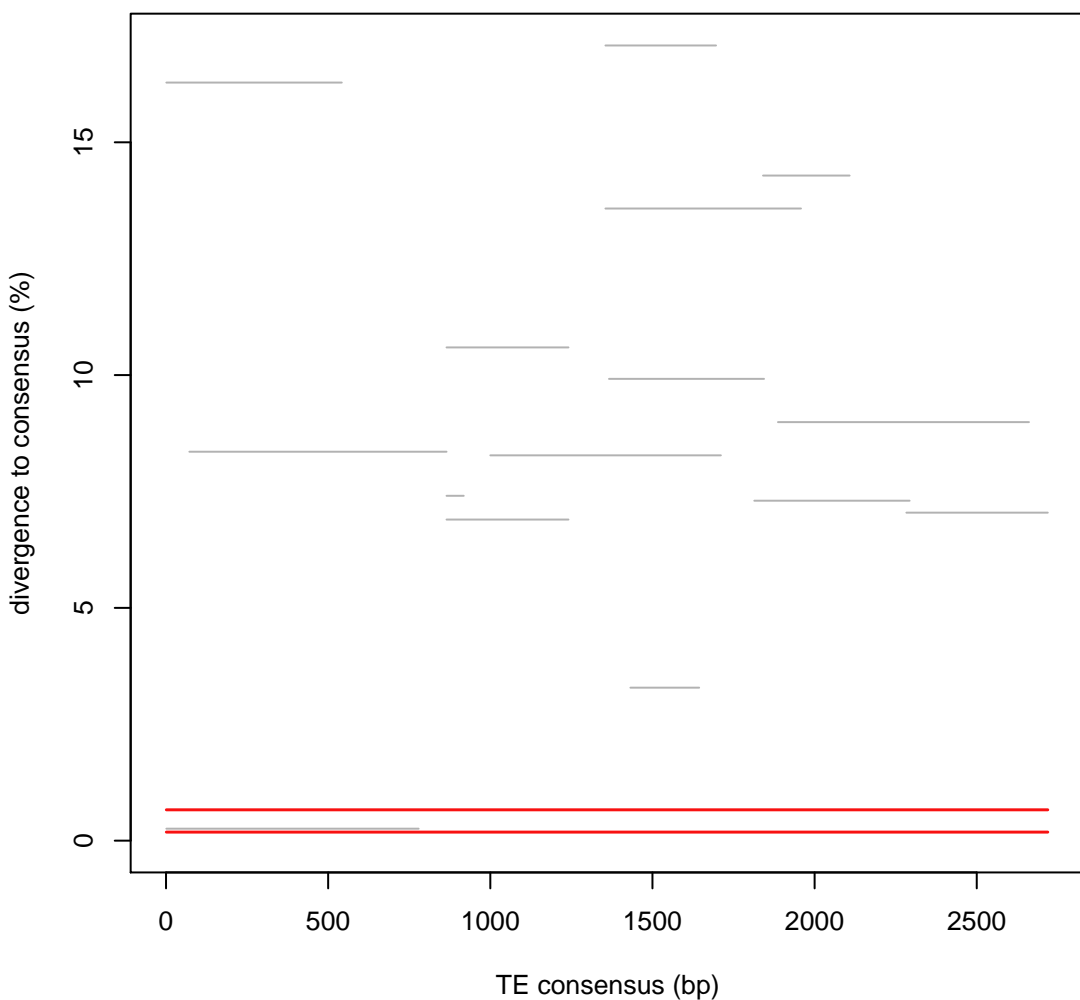
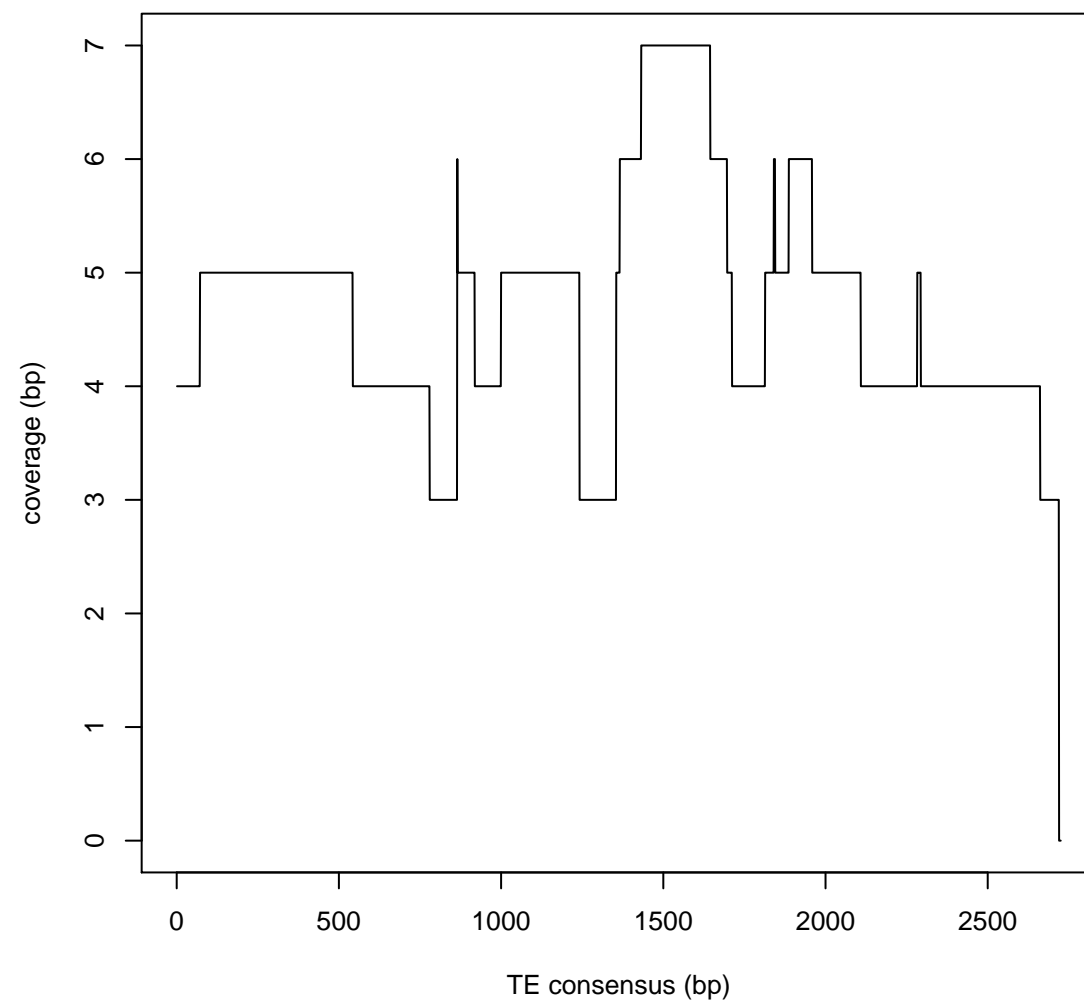


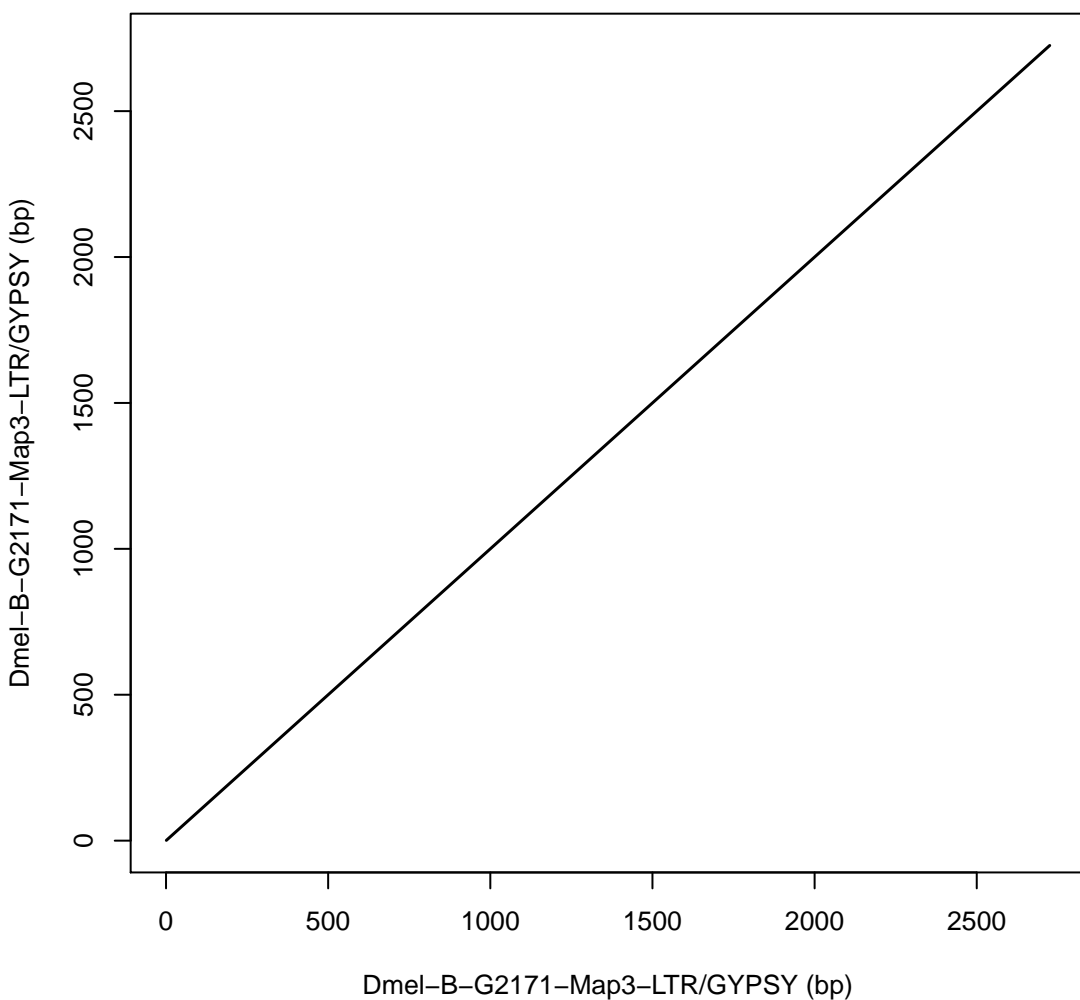
TE: Dmel-B-G2171-Map3-LTR/GYPSY  
consensus size: 2725bp; fragments: 17; full length: 2 ( $\geq 2452.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

