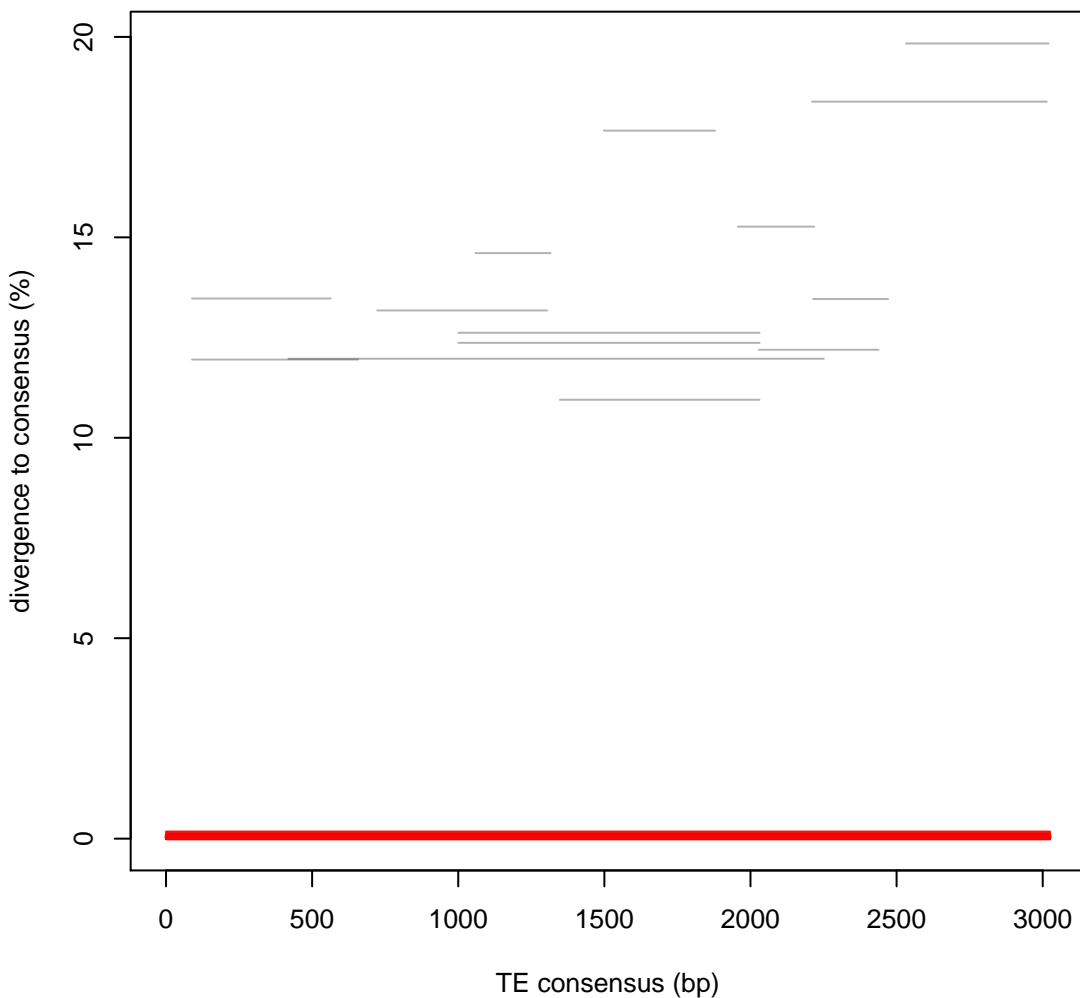
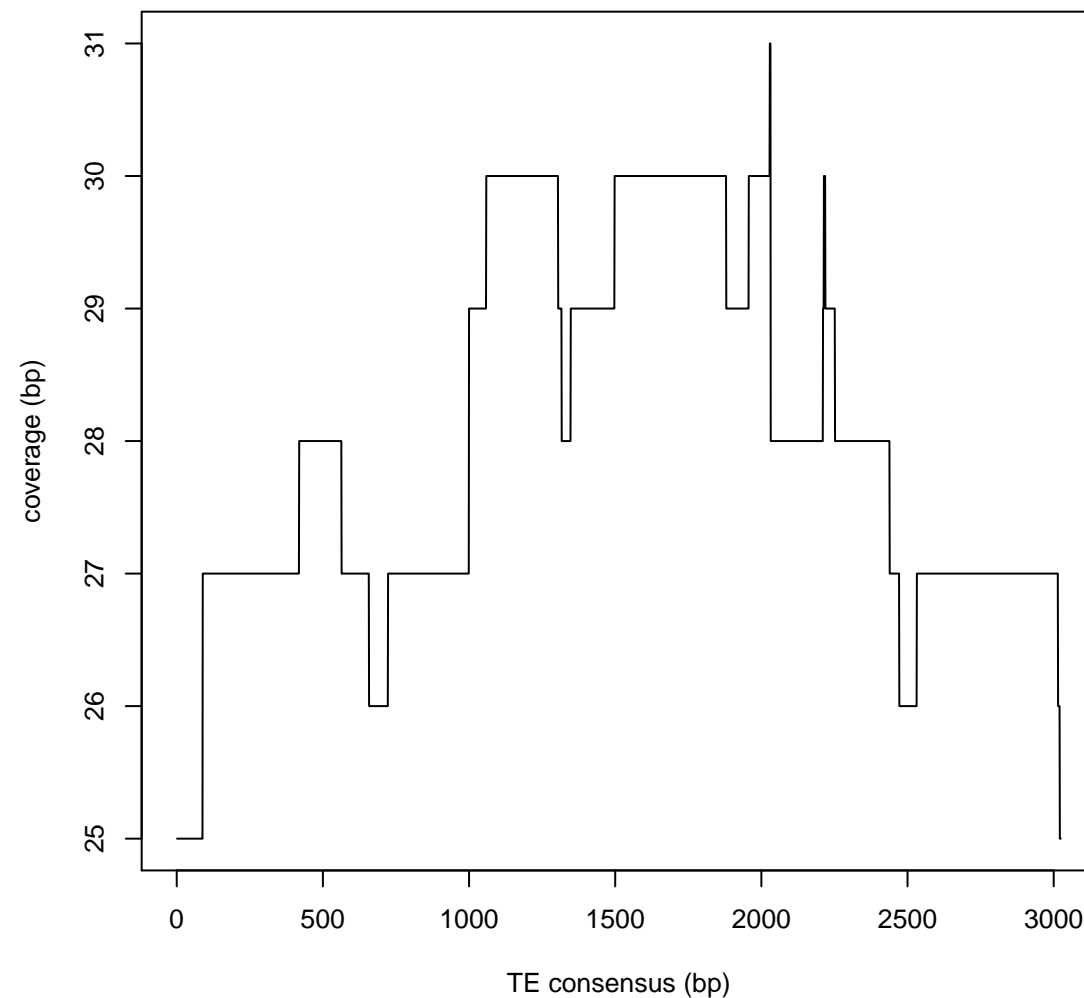


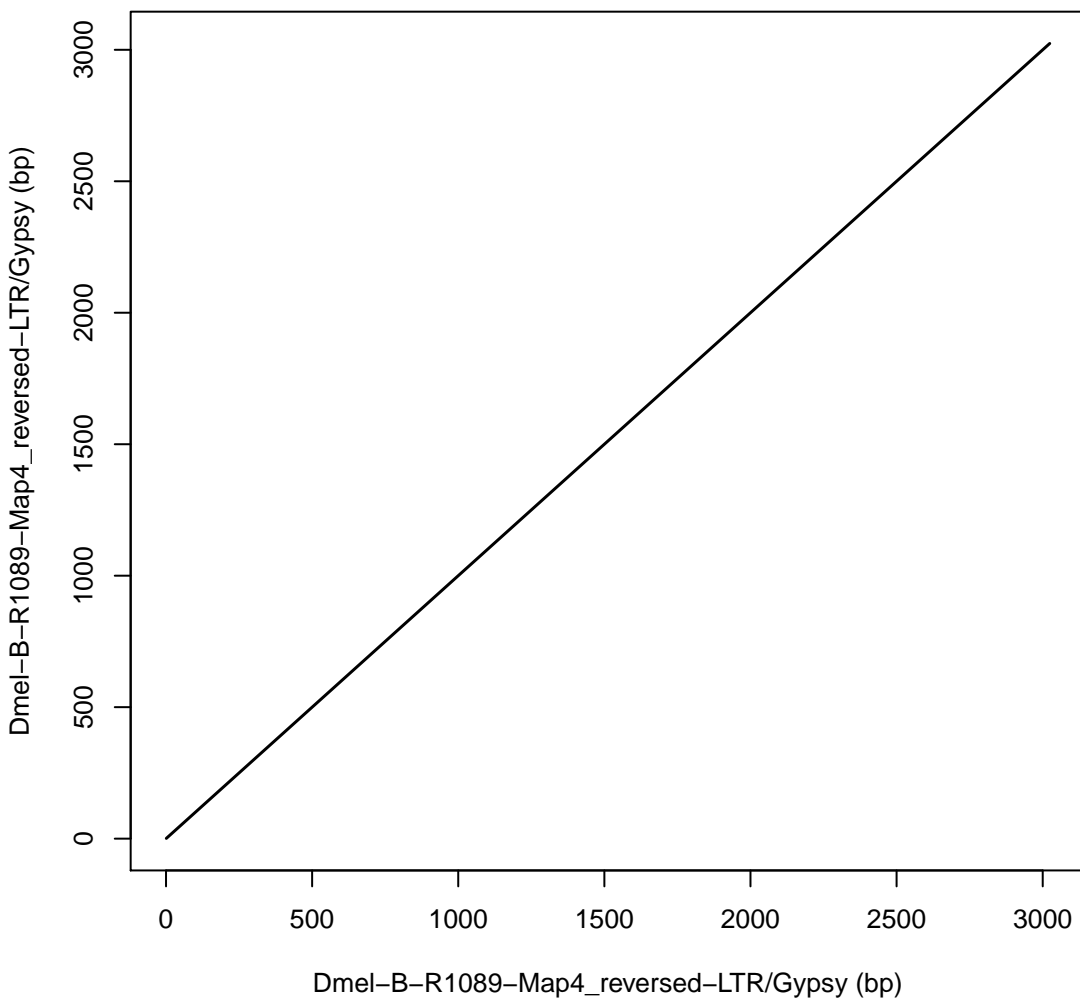
TE: Dmel-B-R1089-Map4\_reversed-LTR/Gypsy  
consensus size: 3024bp; fragments: 39; full length: 25 ( $\geq 2721.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

no orf >400 bp detected