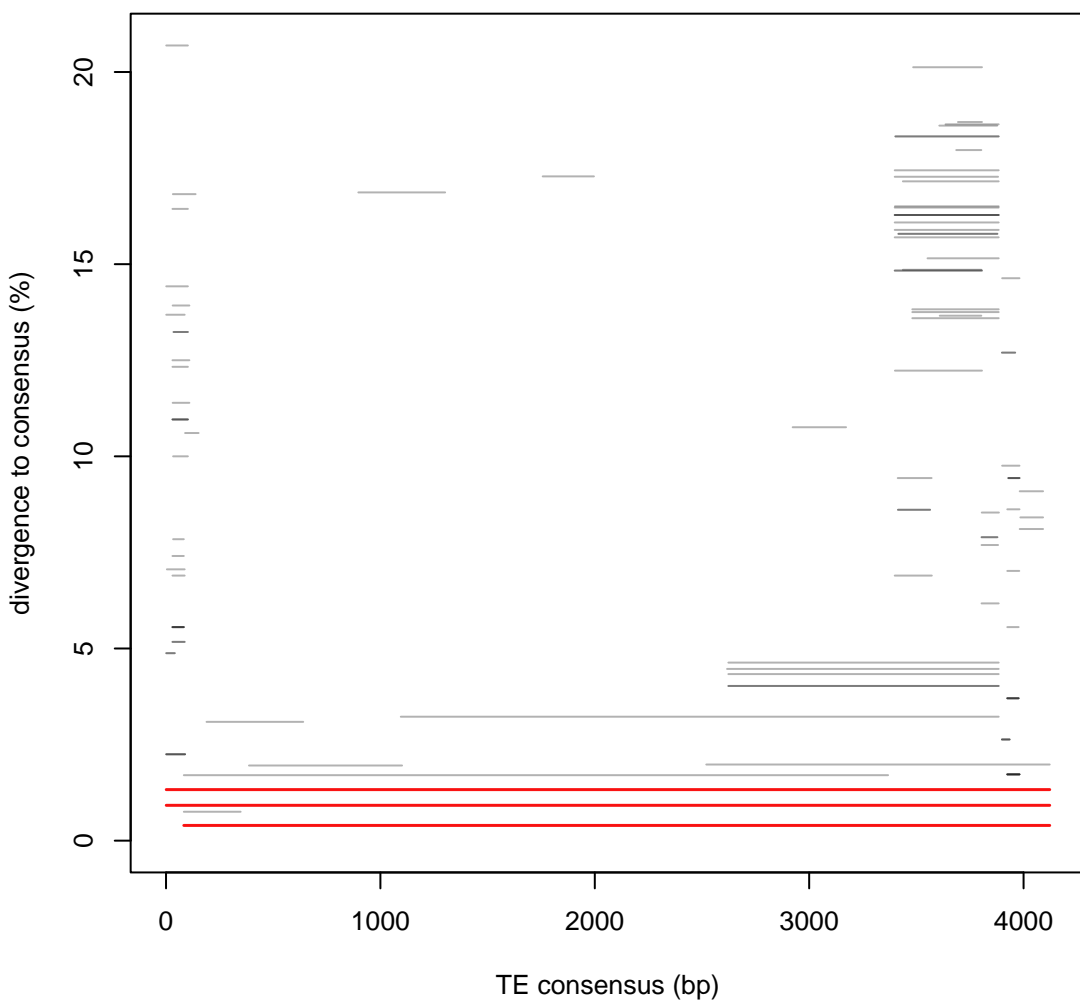
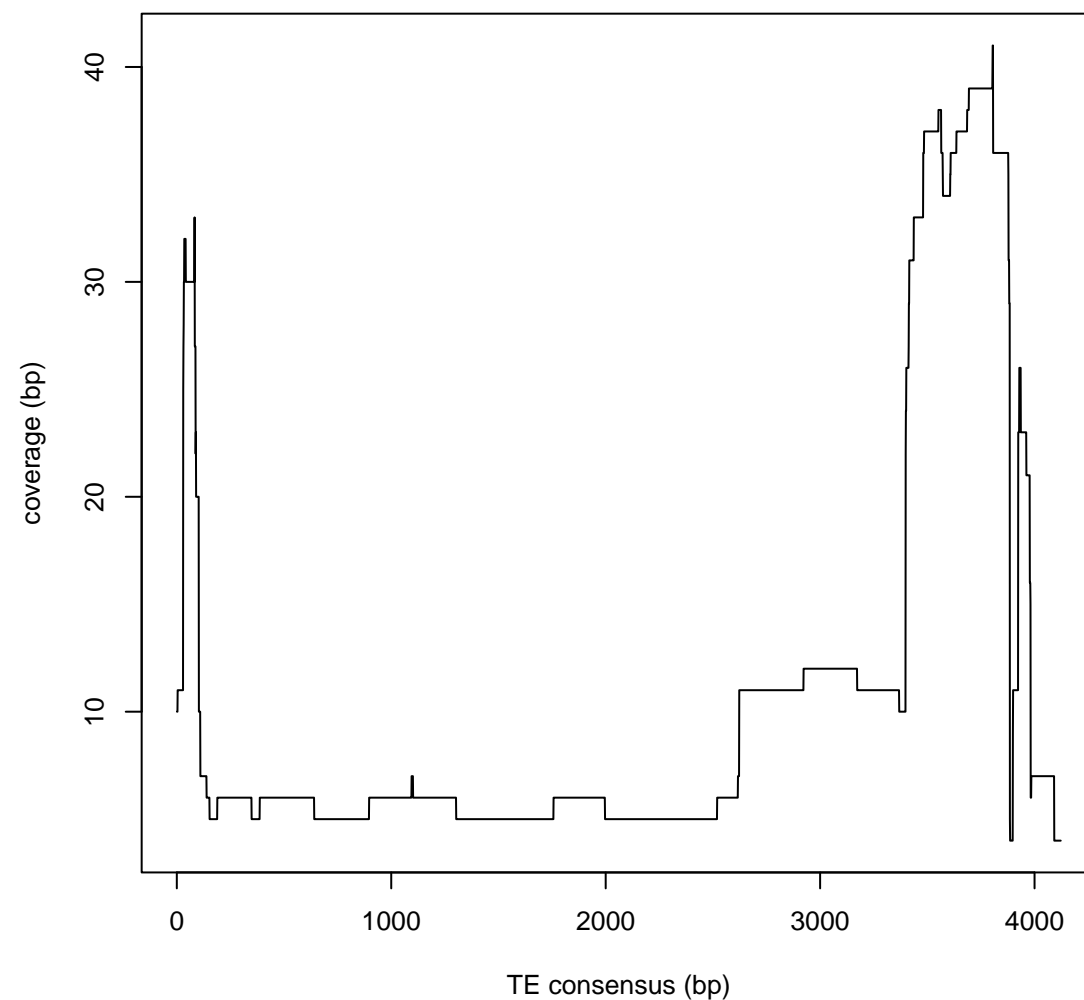


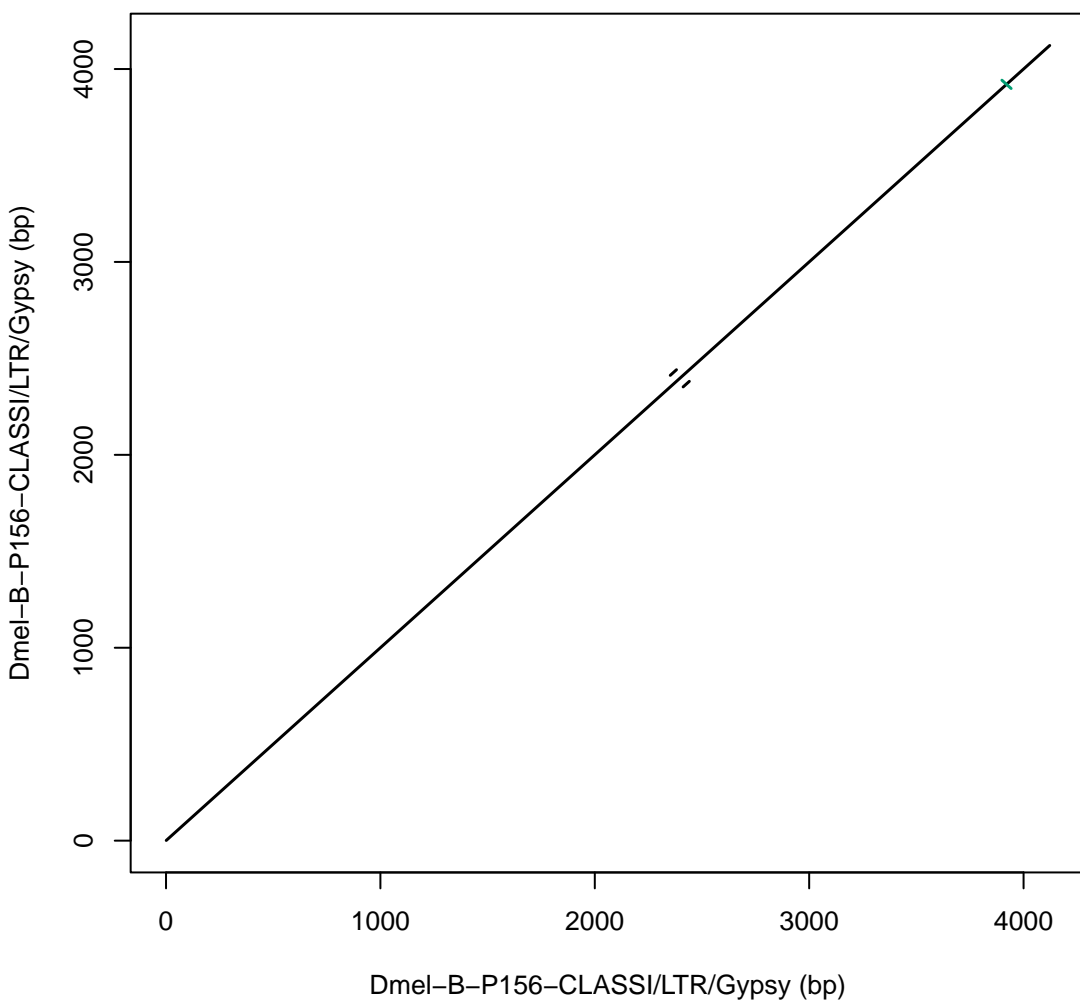
TE: Dmel-B-P156-CLASSI/LTR/Gypsy  
consensus size: 4122bp; fragments: 111; full length: 3 ( $\geq 3709.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

