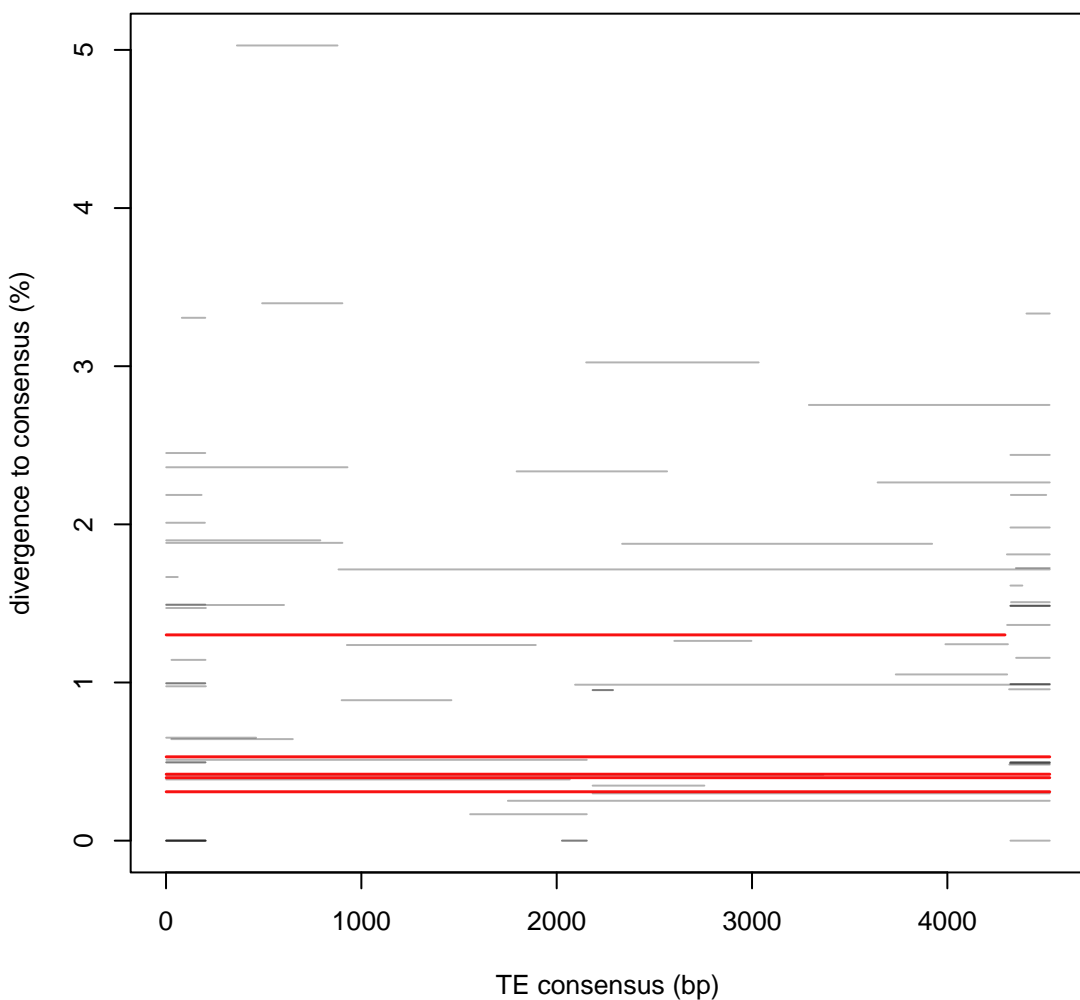
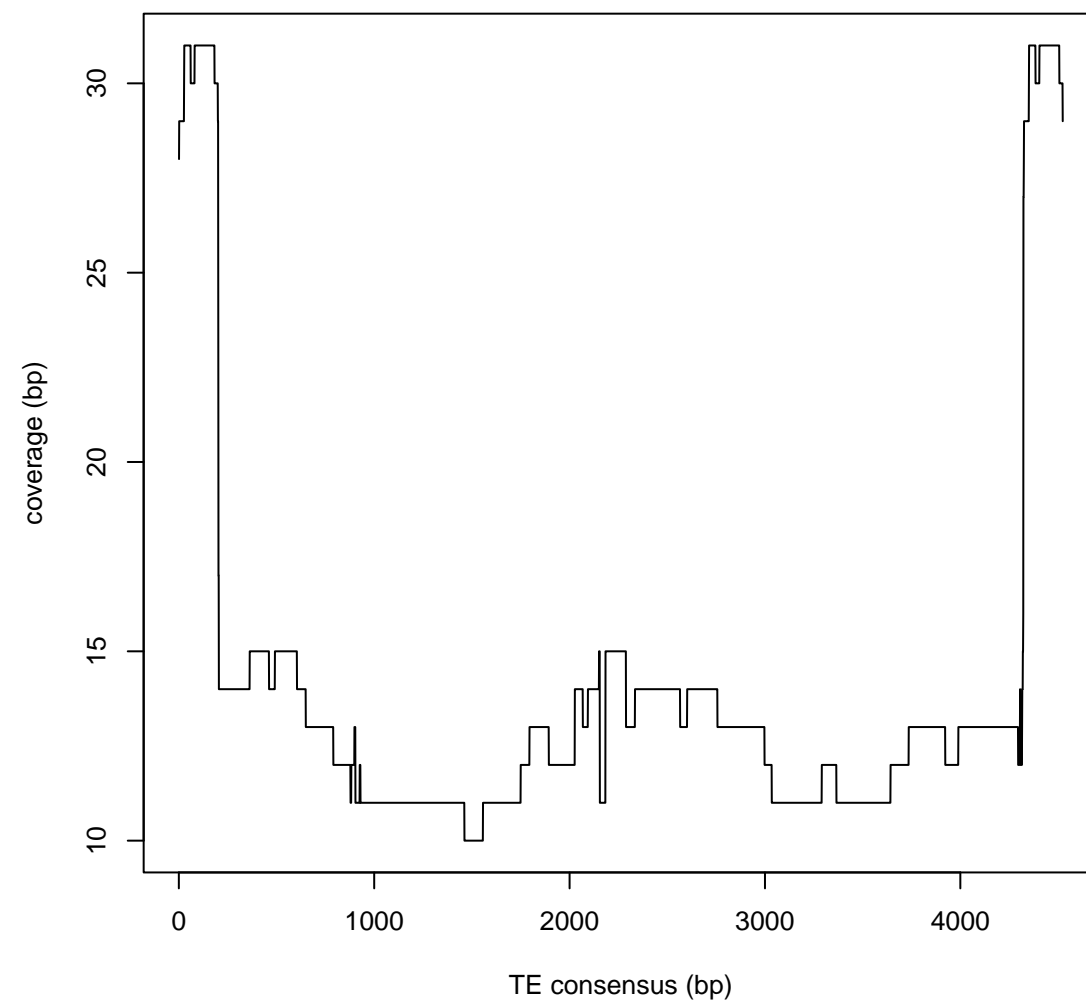


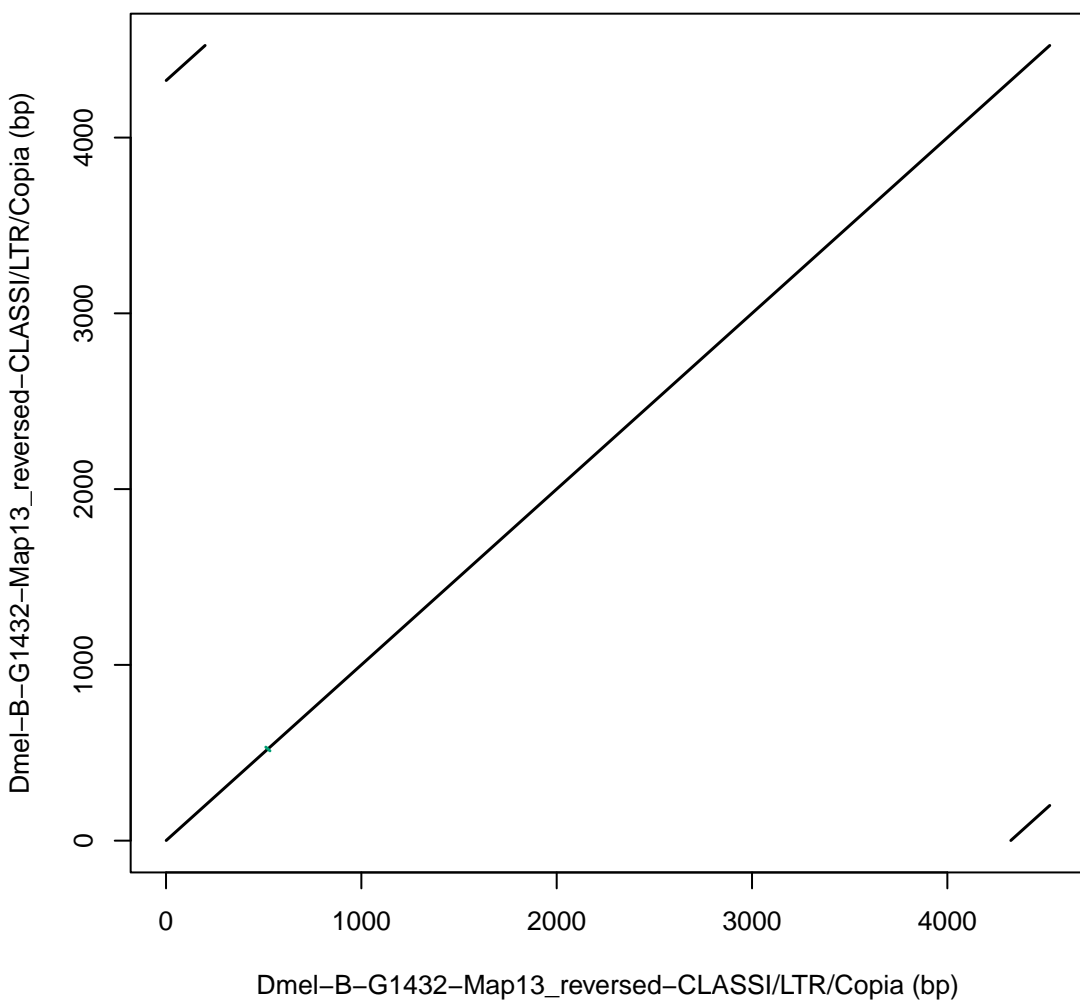
TE: Dmel-B-G1432-Map13\_reversed-CLASSI/LTR/Copia  
consensus size: 4524bp; fragments: 76; full length: 5 ( $\geq 4071.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

