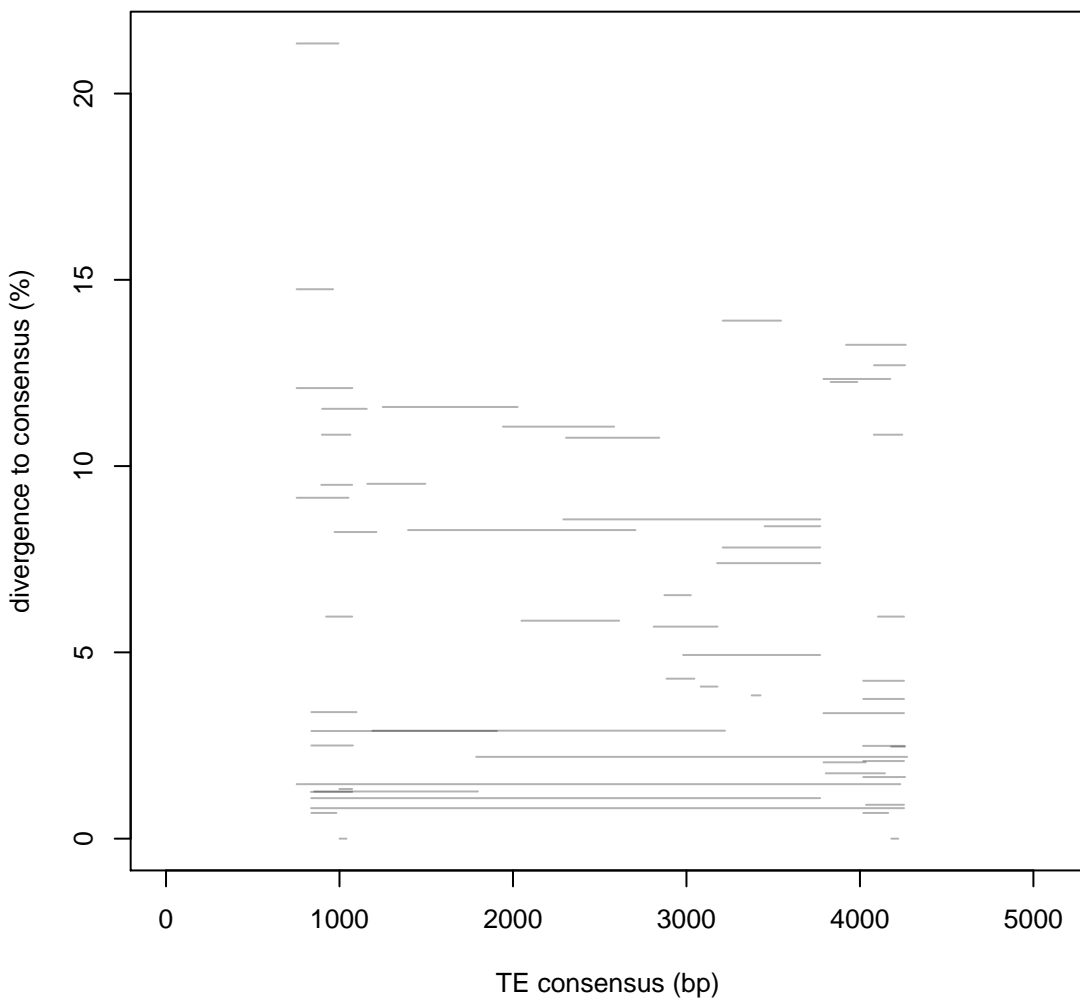
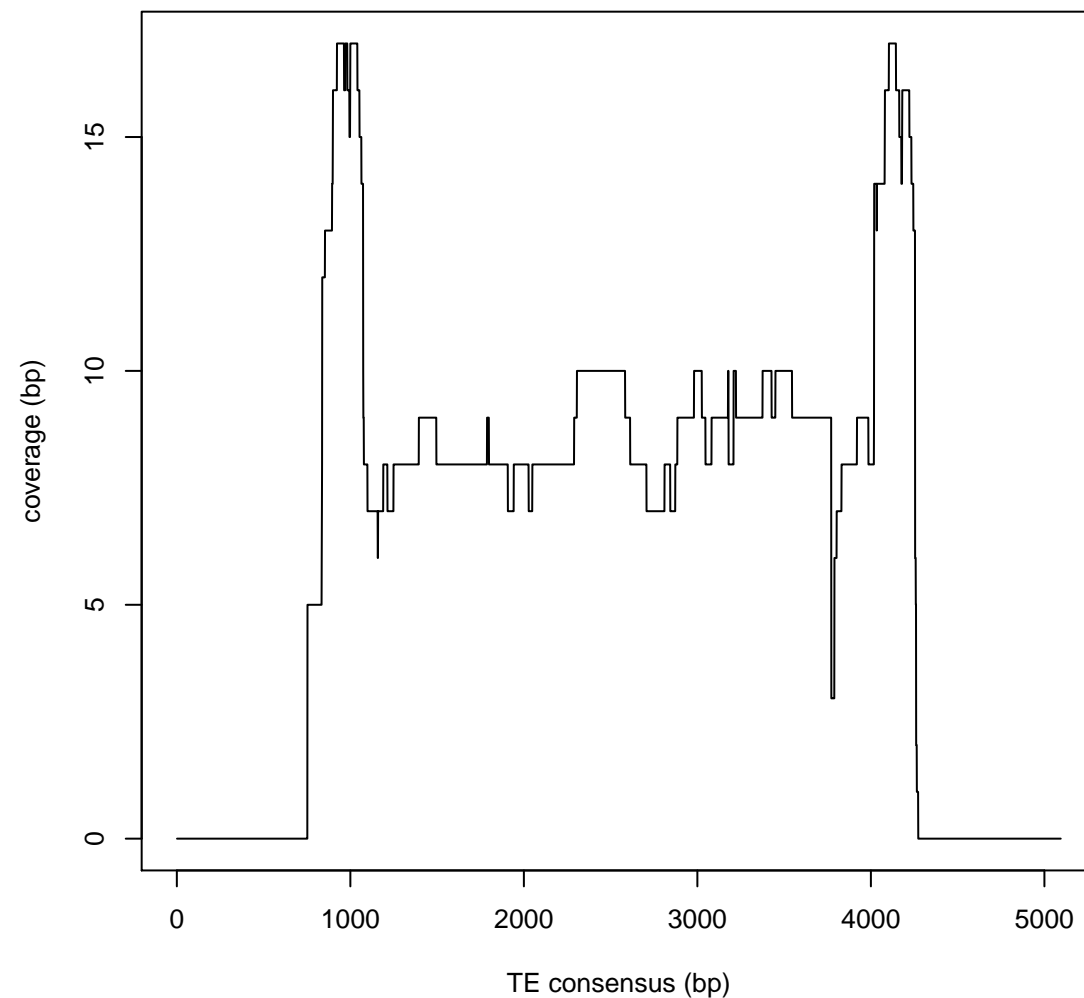


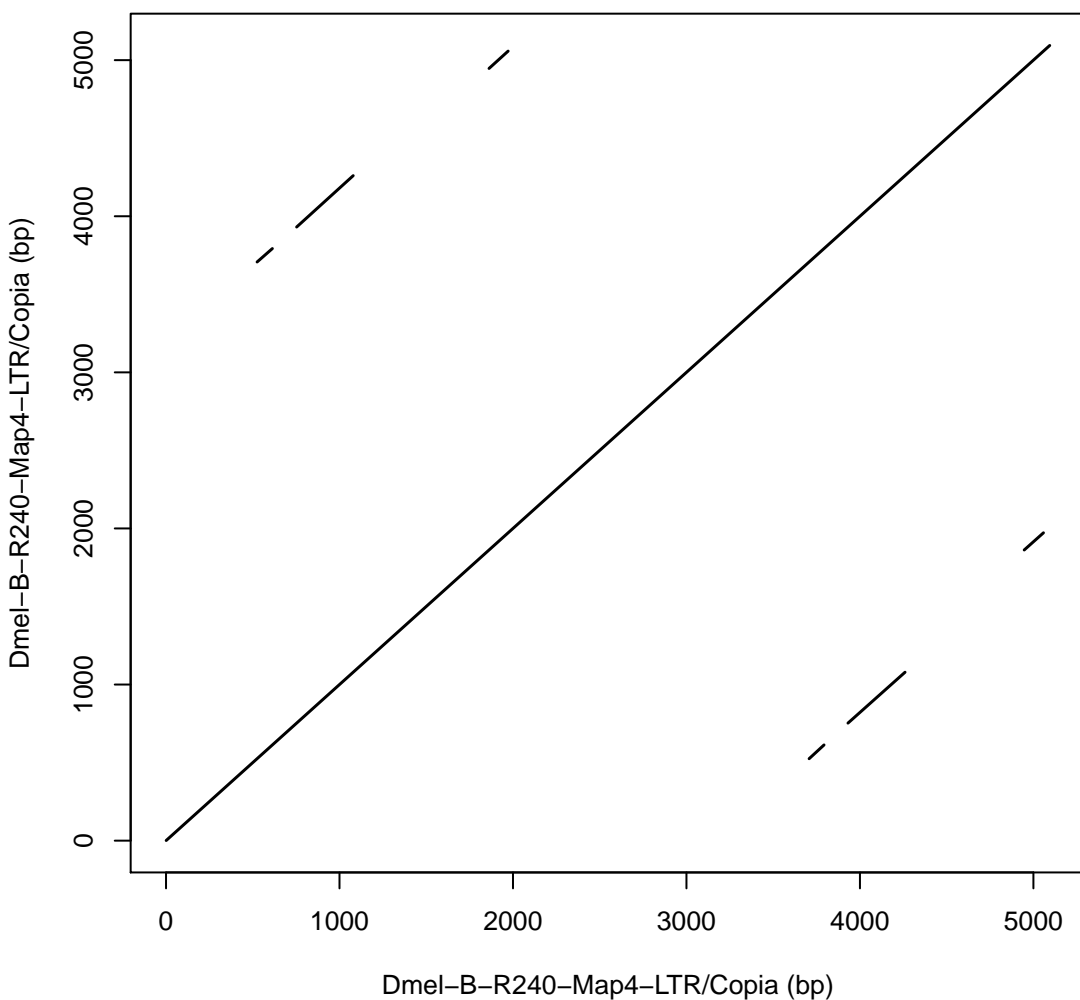
TE: Dmel-B-R240-Map4-LTR/Copia  
consensus size: 5094bp; fragments: 57; full length: 0 ( $\geq 4584.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

