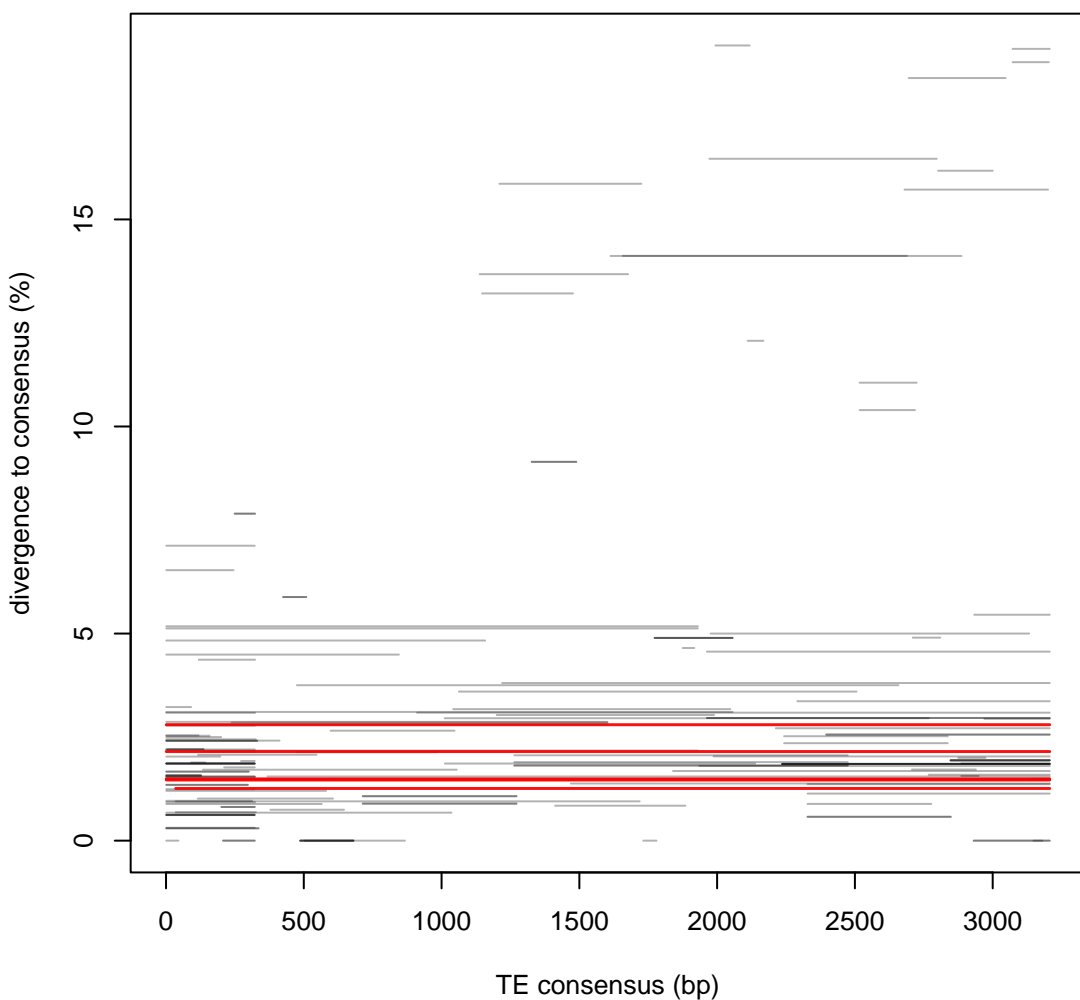
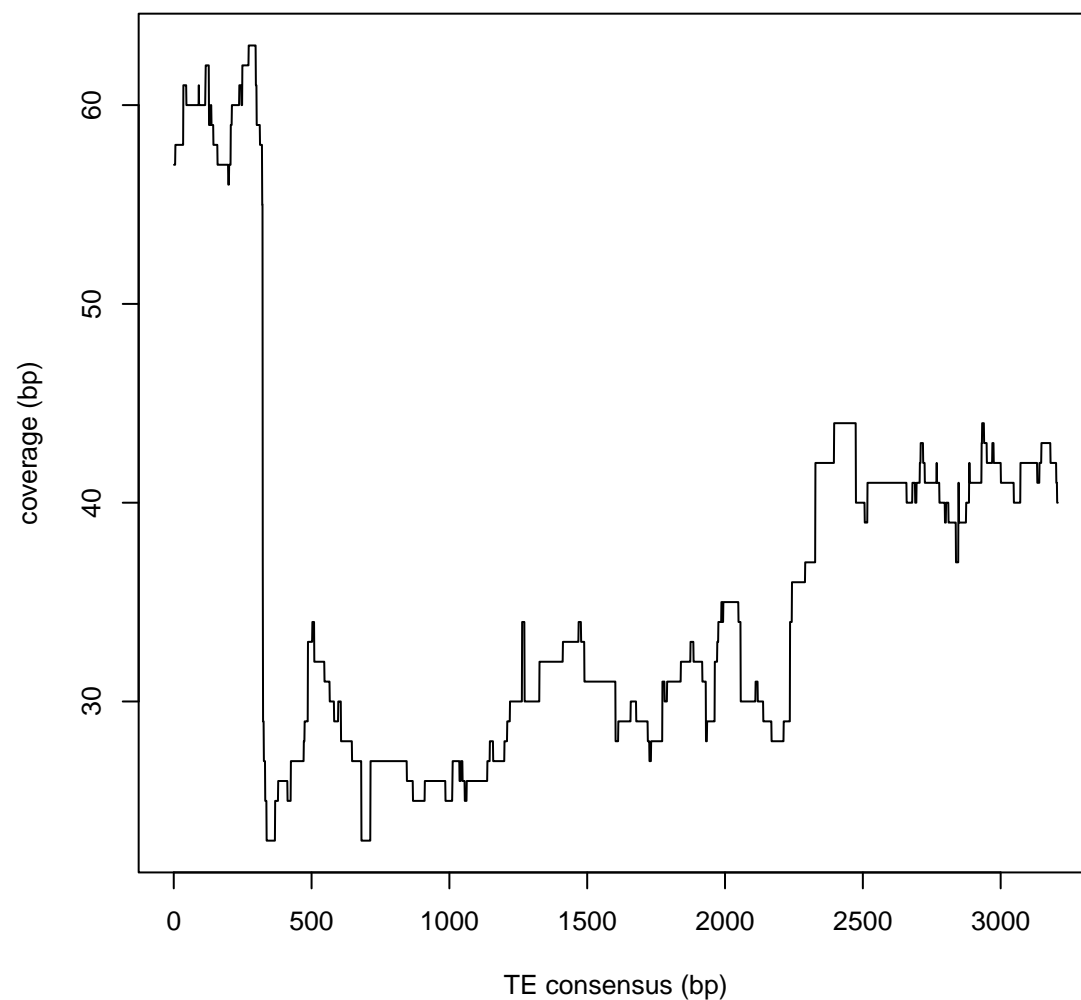


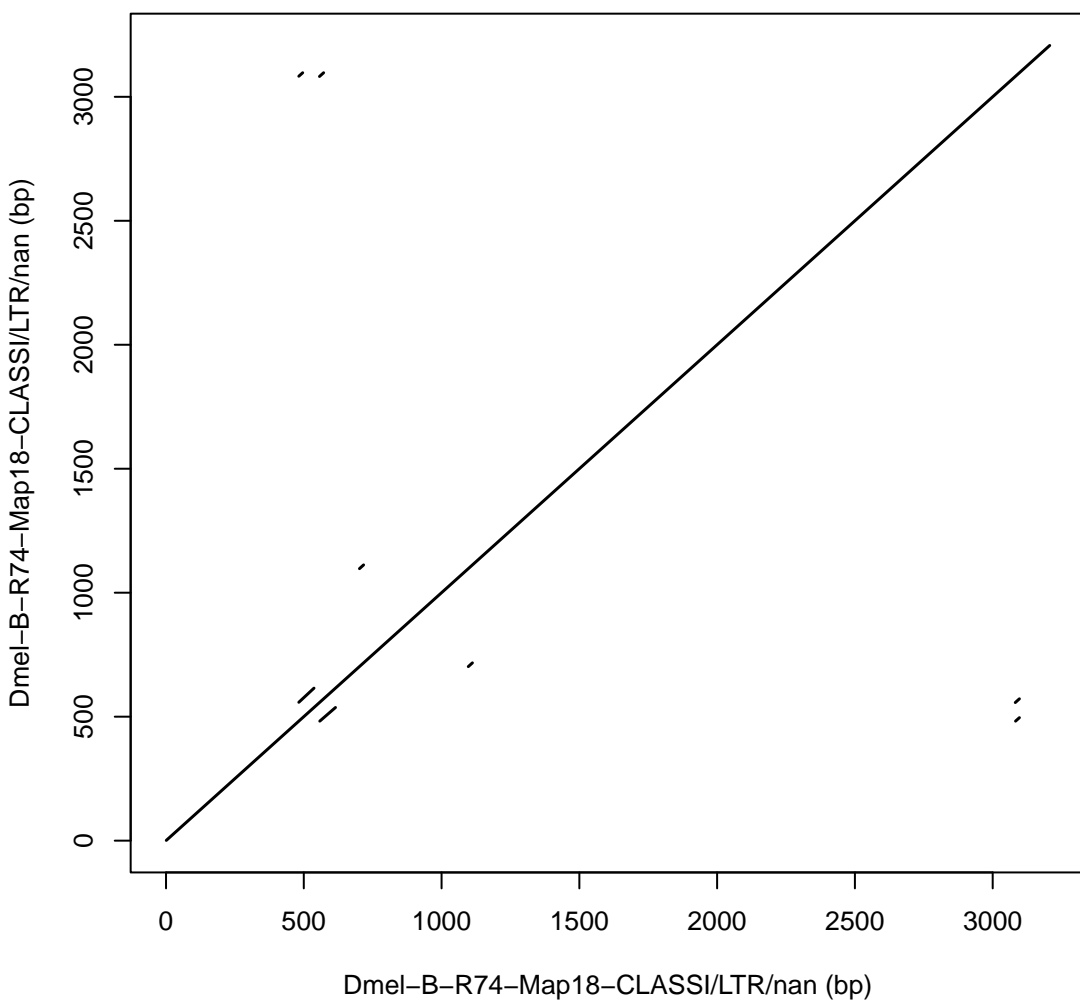
TE: Dmel-B-R74-Map18-CLASSI/LTR/nan  
consensus size: 3207bp; fragments: 168; full length: 5 ( $\geq 2886.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

