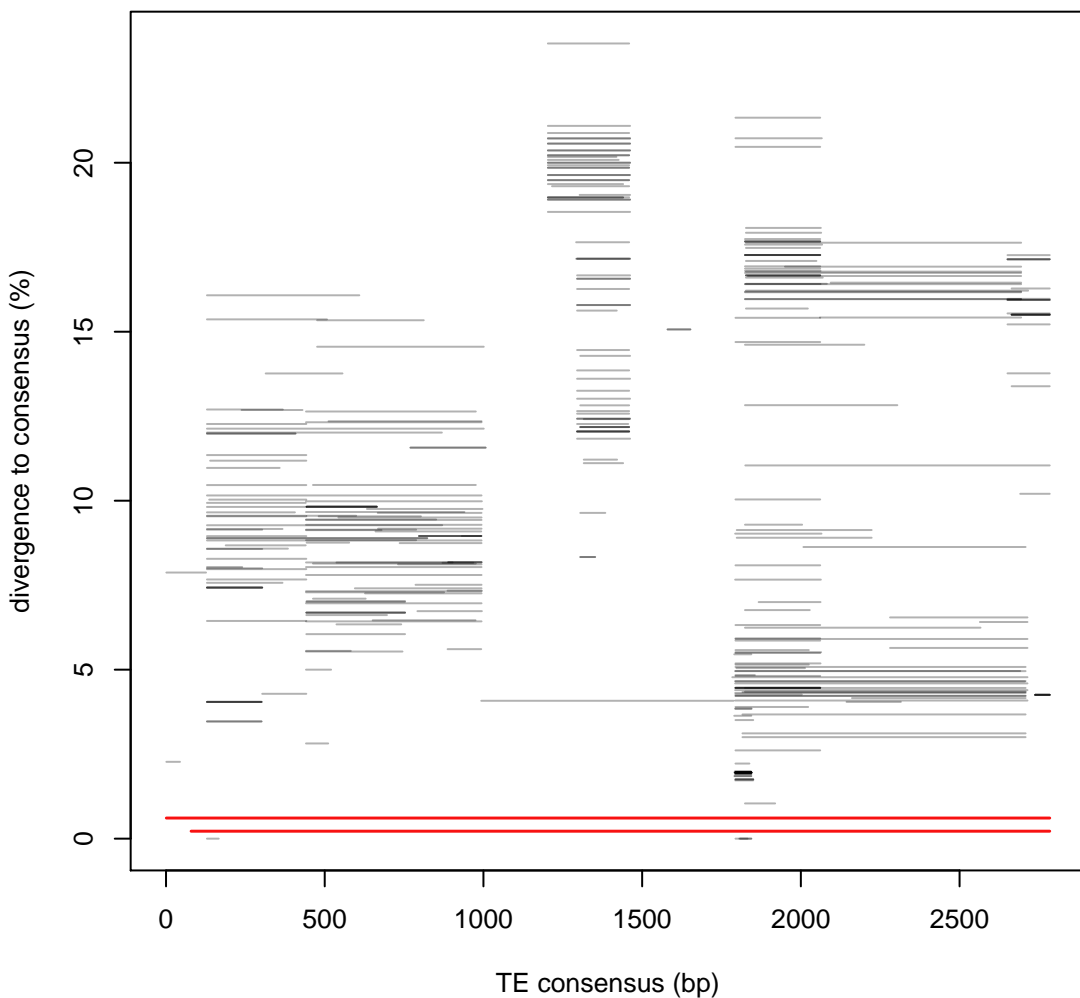
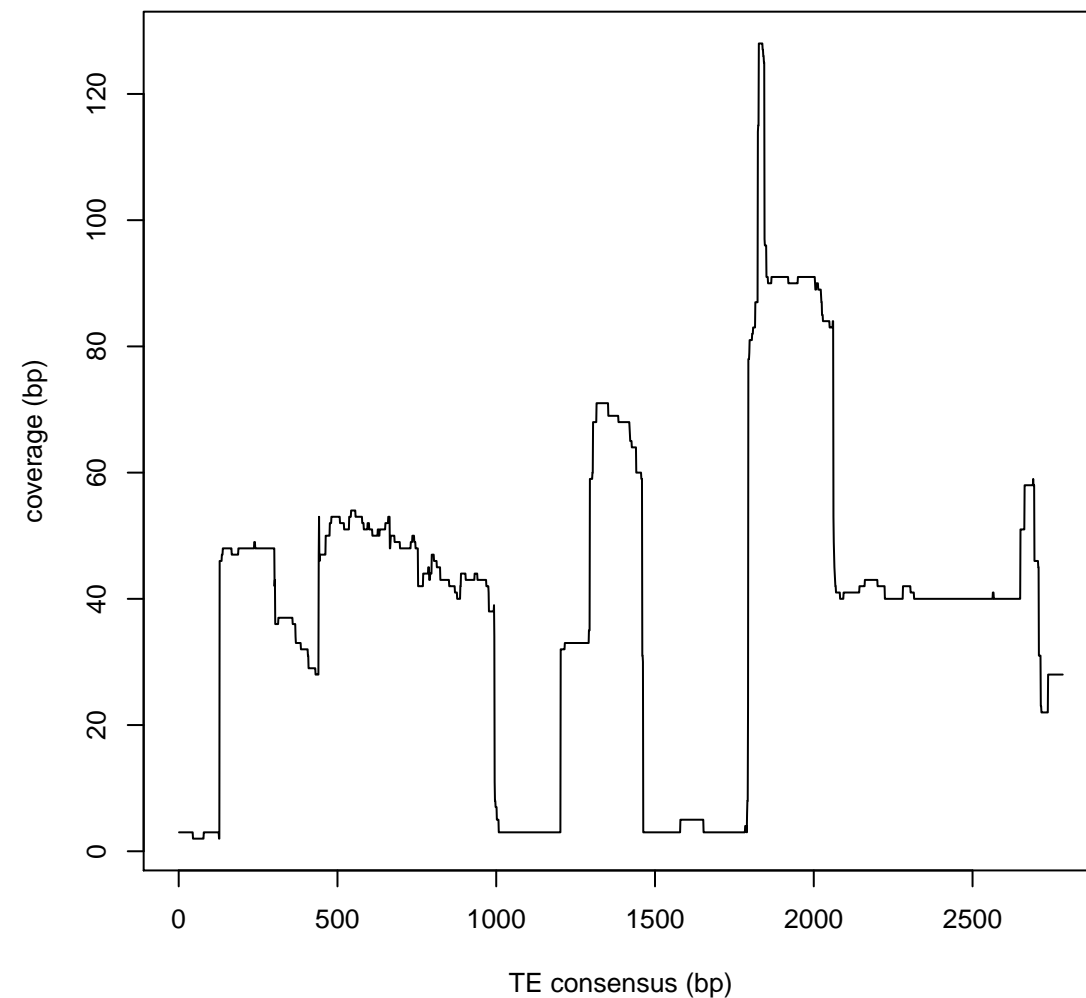


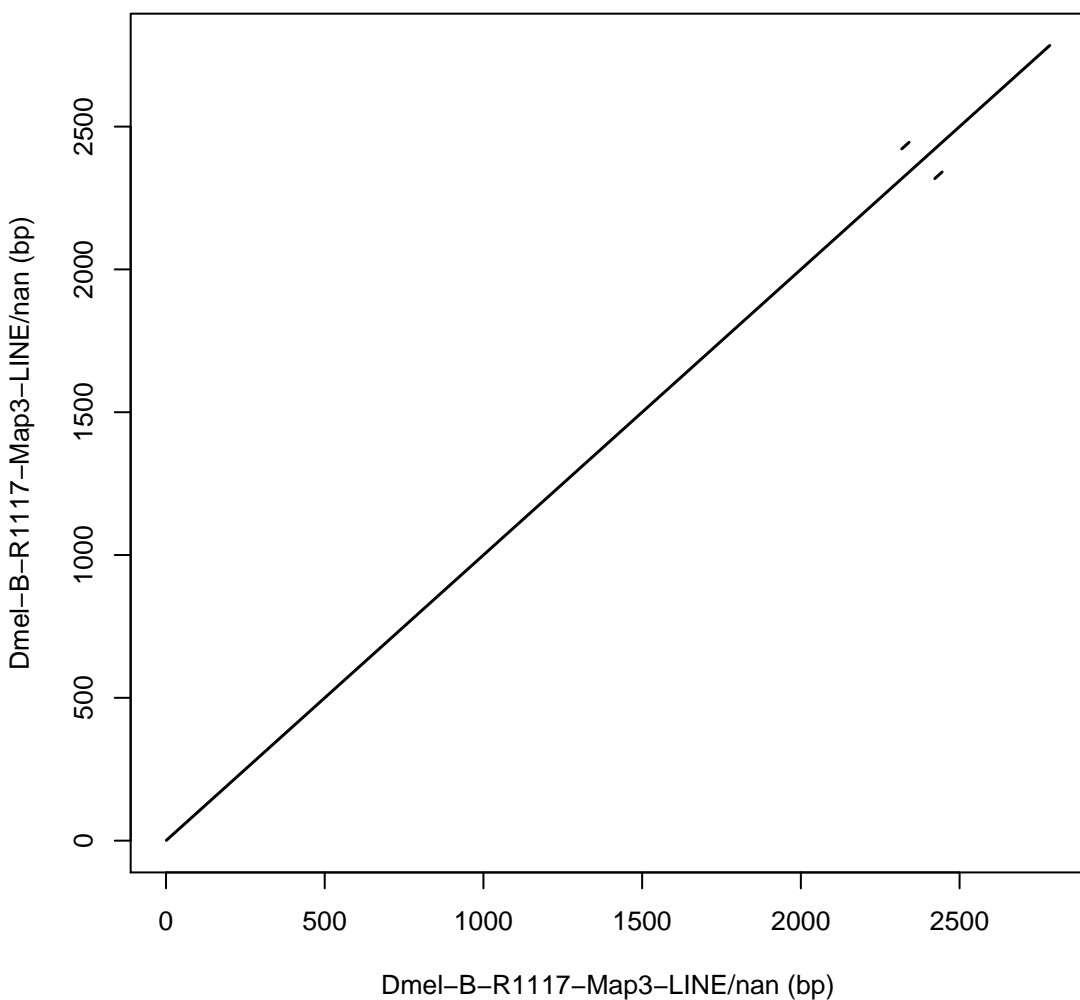
TE: Dmel-B-R1117-Map3-LINE/nan  
consensus size: 2784bp; fragments: 354; full length: 2 ( $\geq 2505.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

