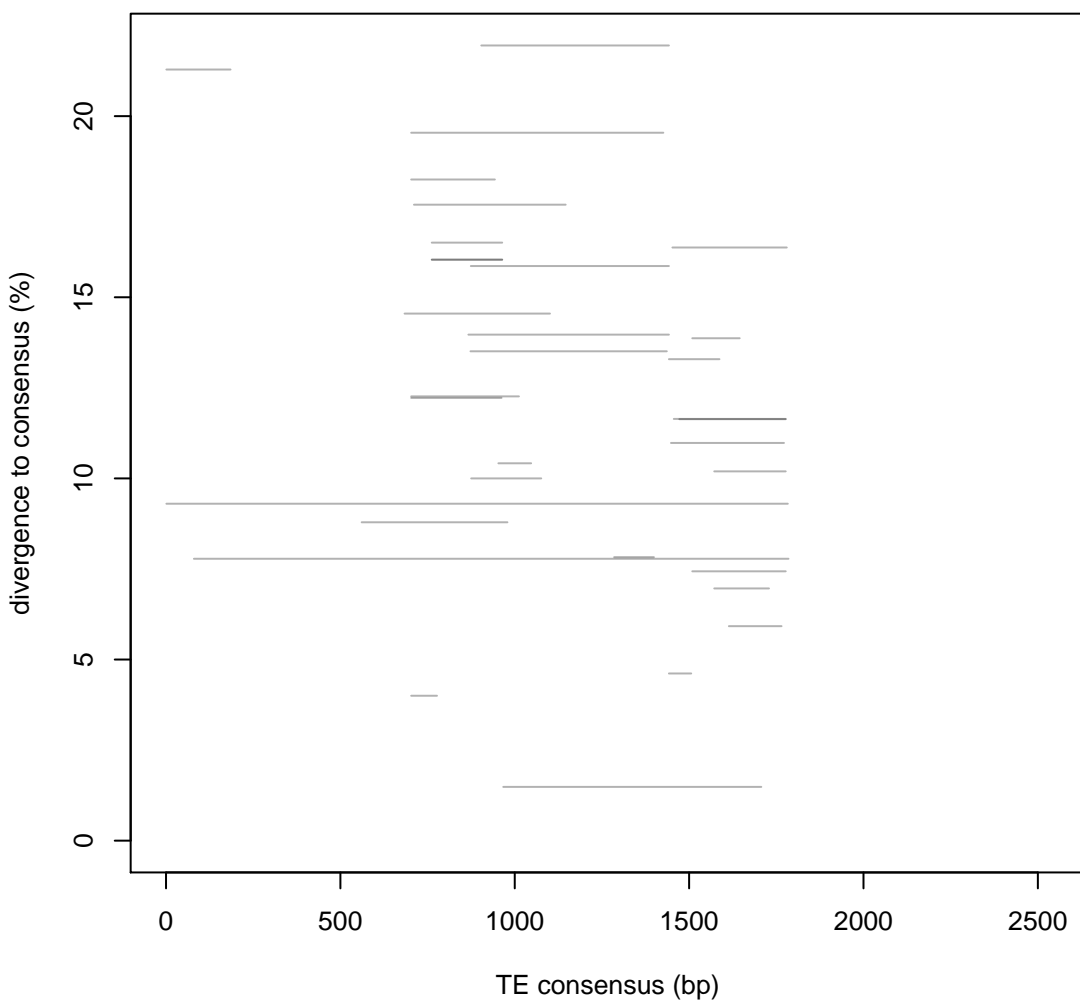
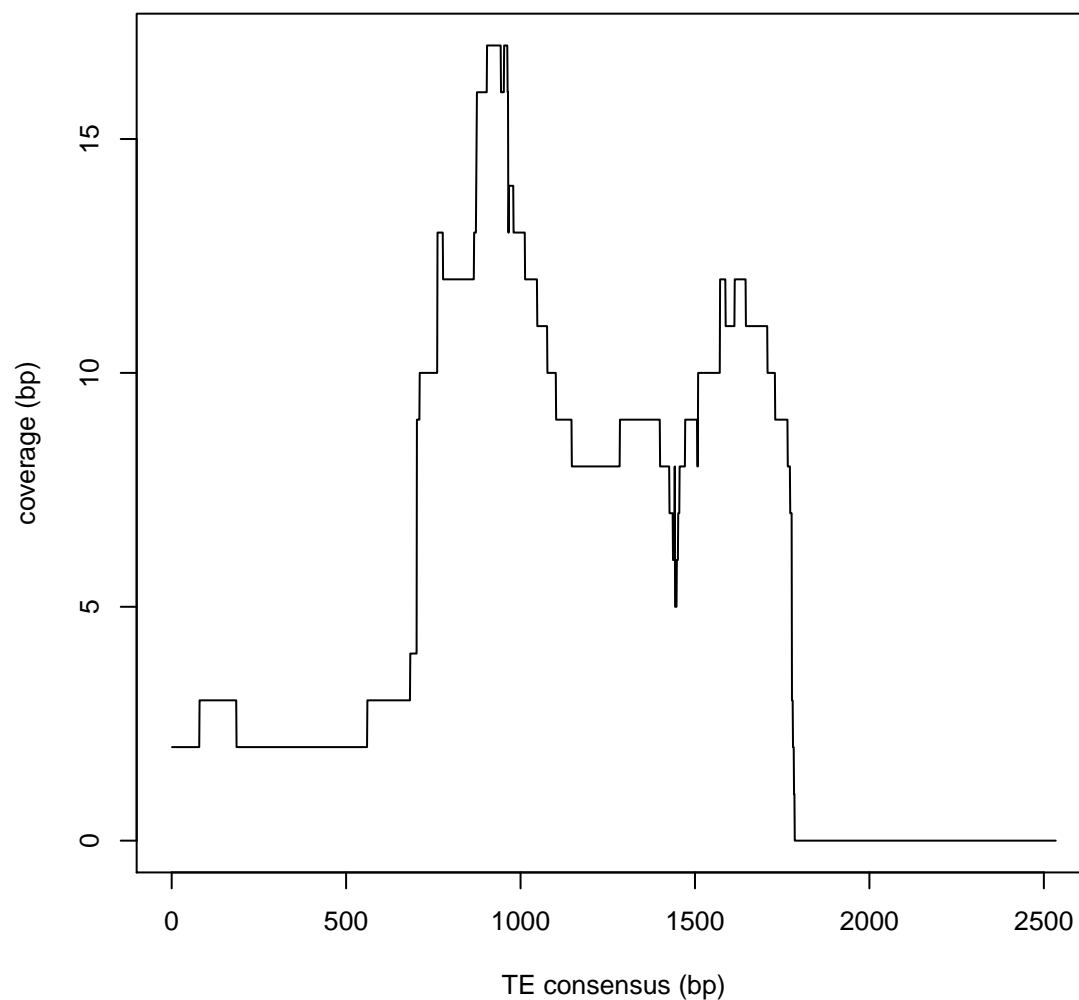


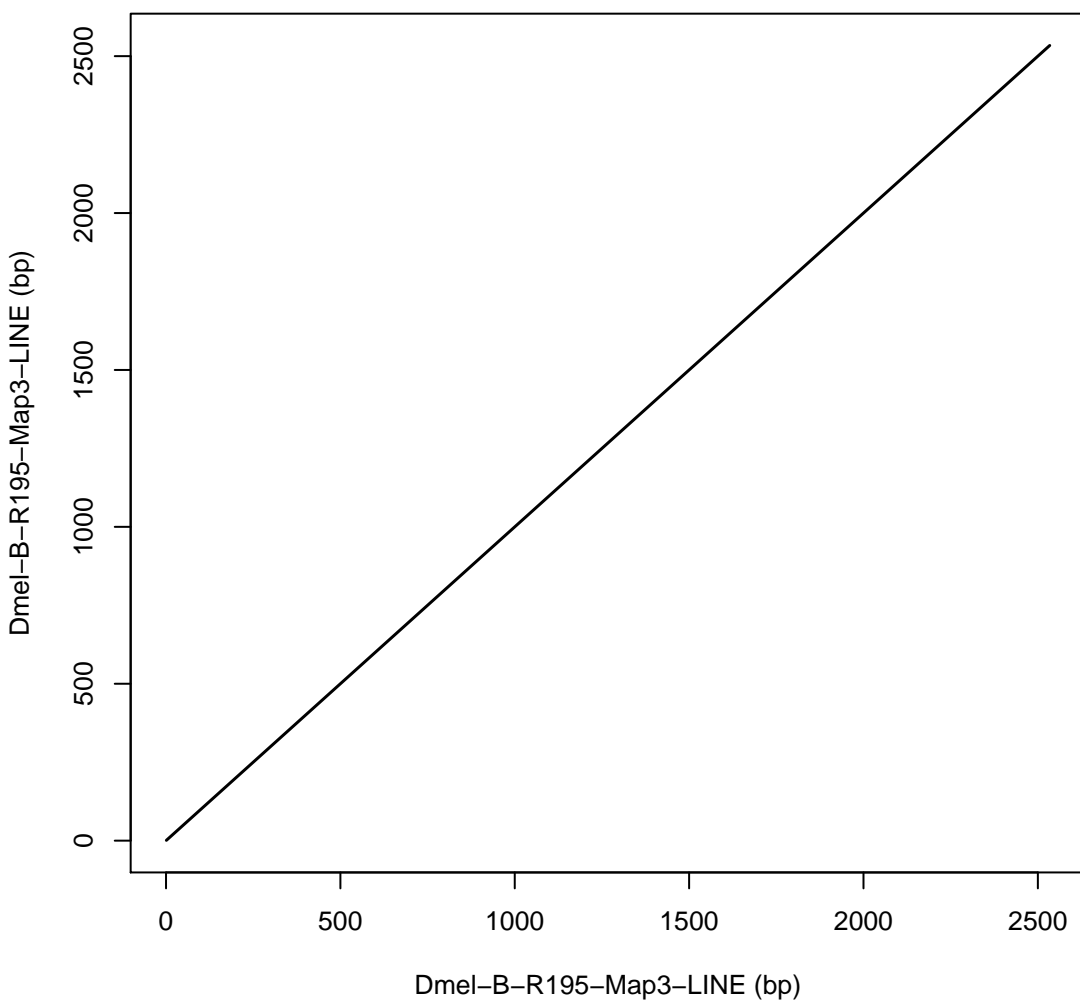
TE: Dmel-B-R195-Map3-LINE  
consensus size: 2534bp; fragments: 33; full length: 0 ( $\geq 2280.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

