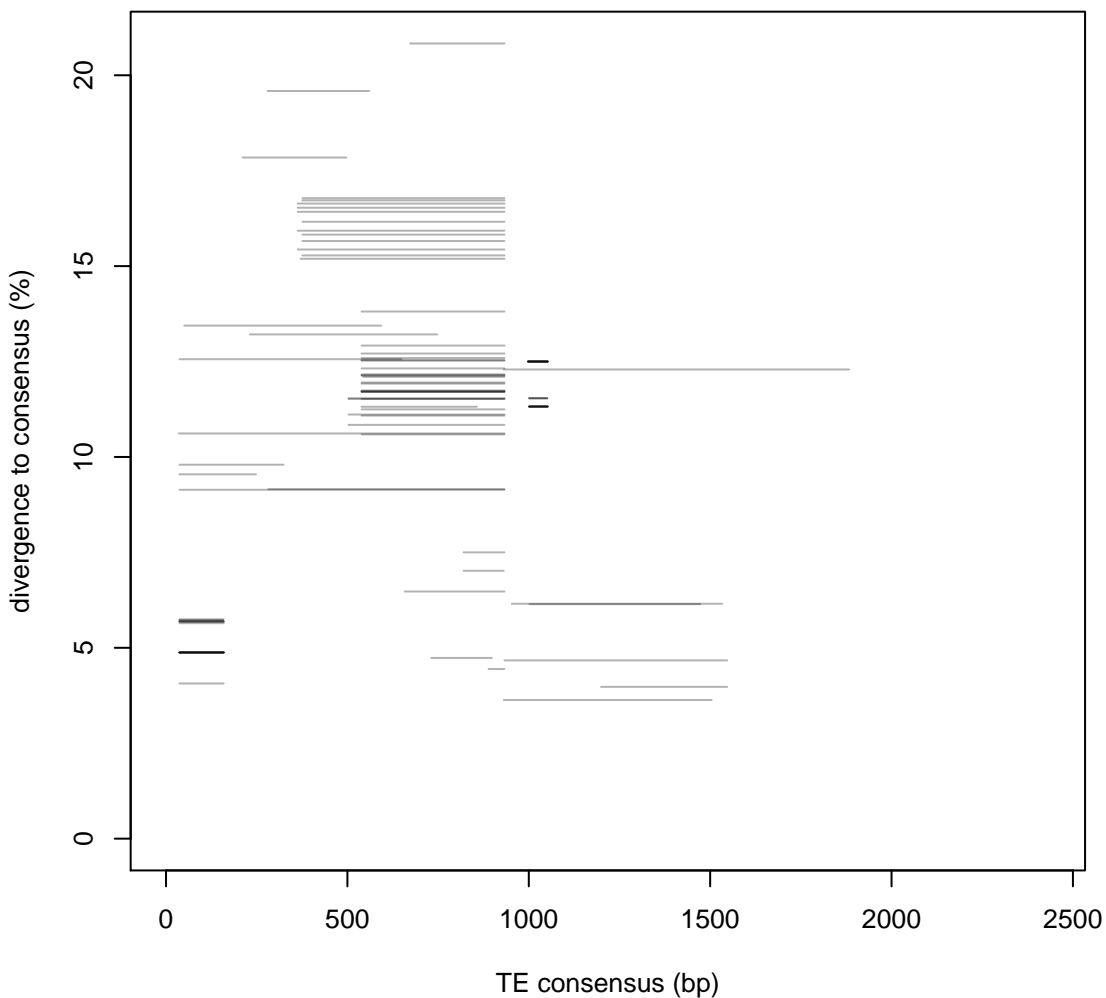
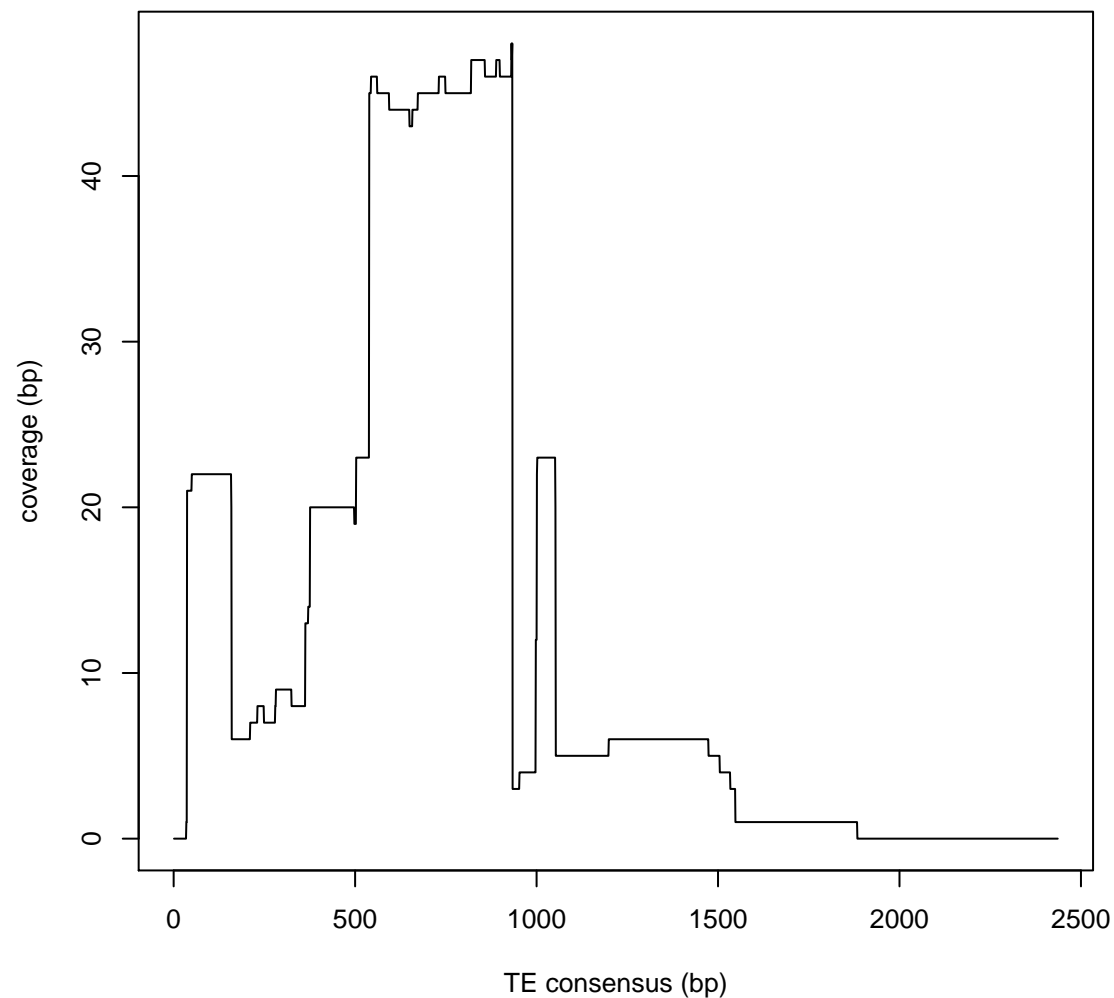


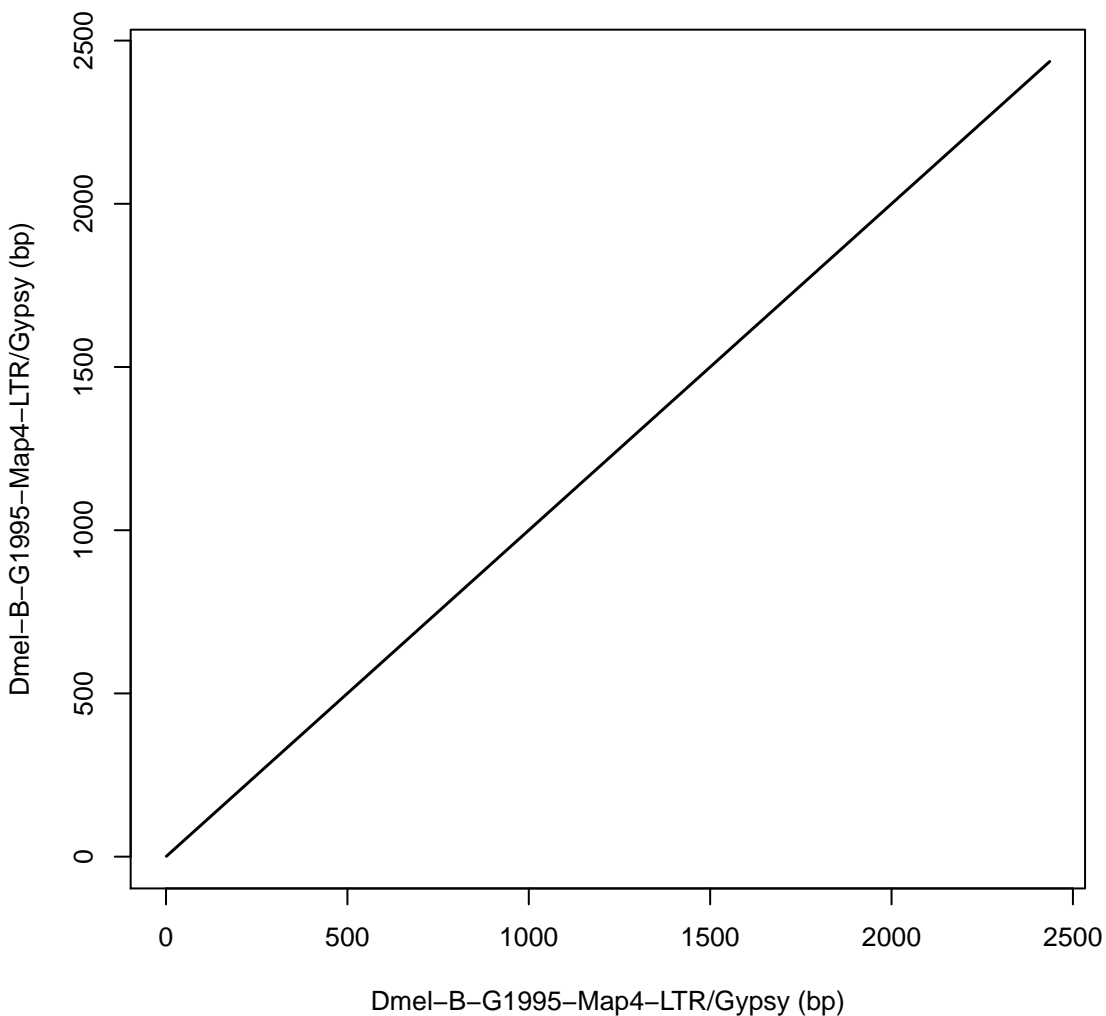
TE: Dmel-B-G1995-Map4-LTR/Gypsy  
consensus size: 2436bp; fragments: 95; full length: 0 (>=2192.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

