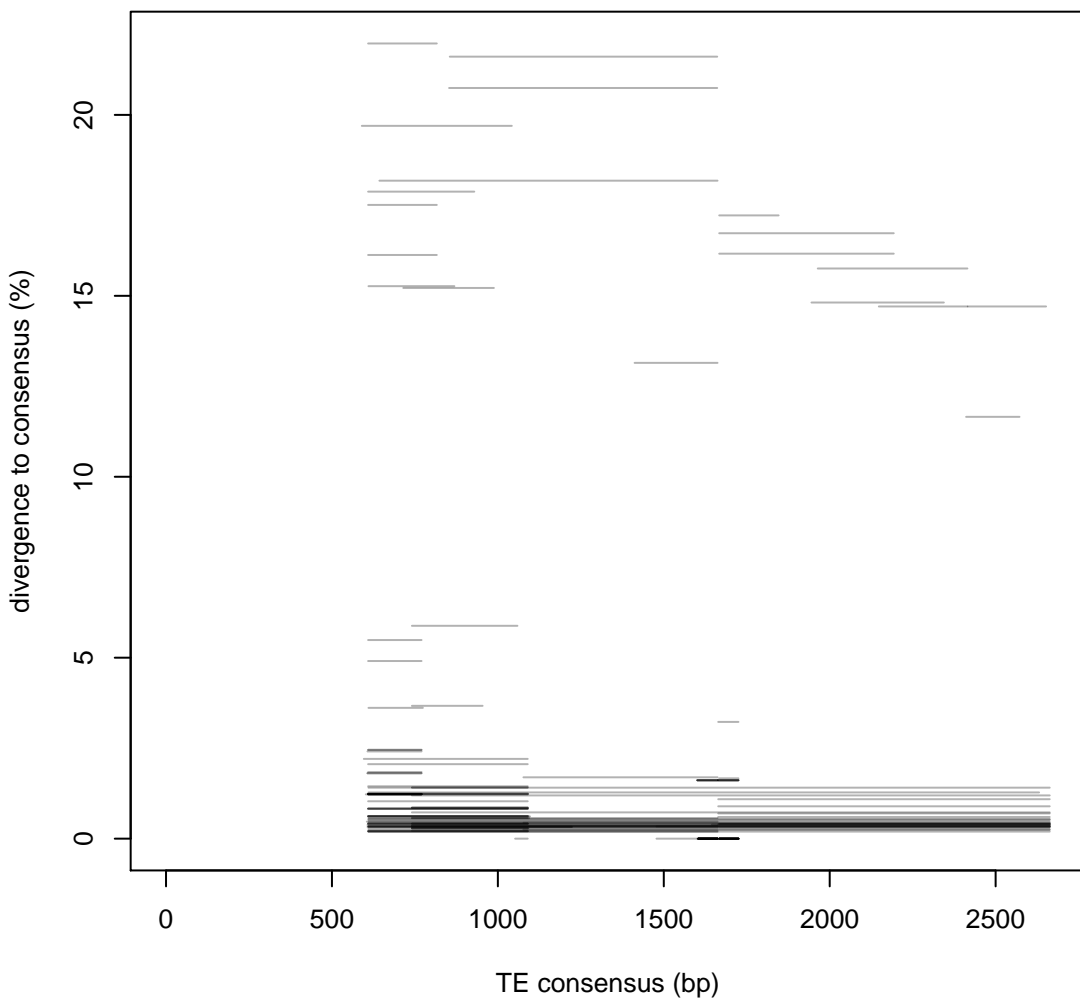
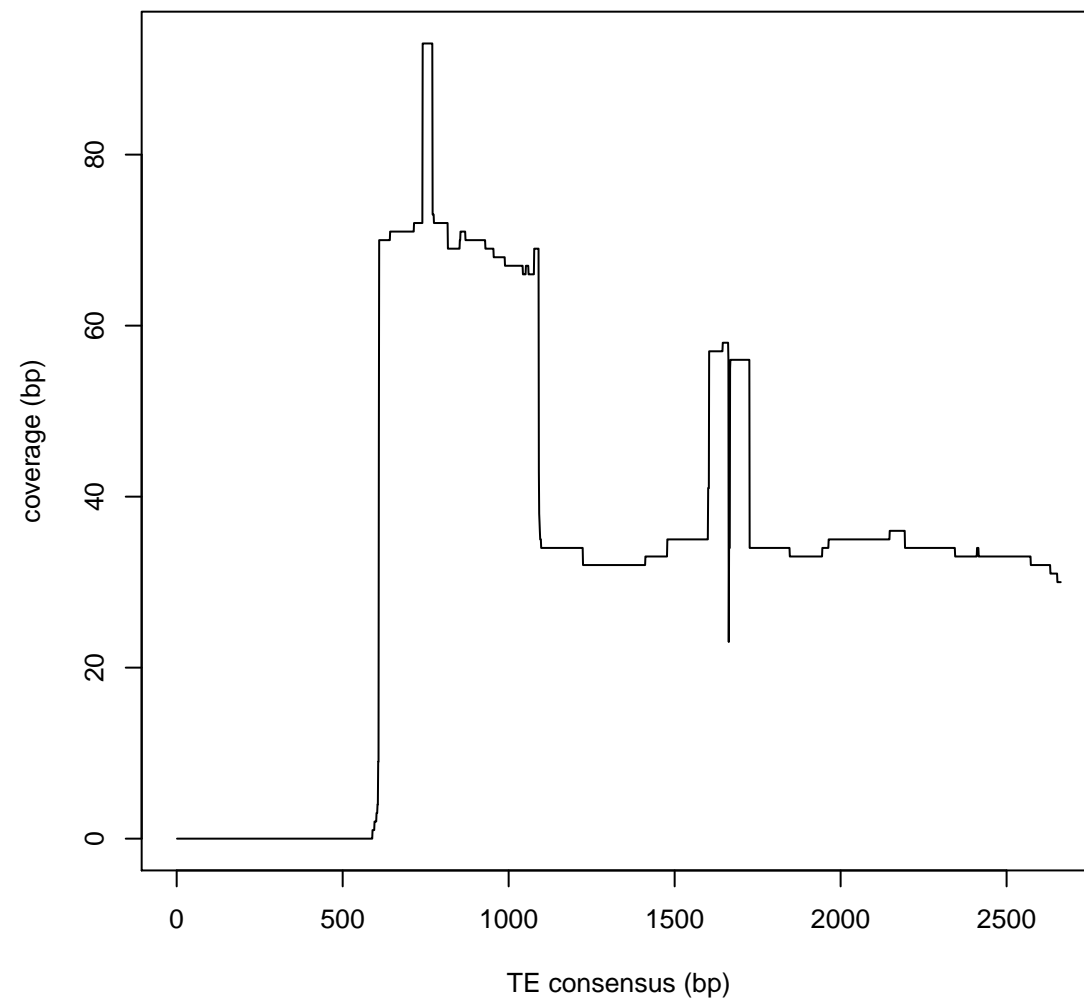


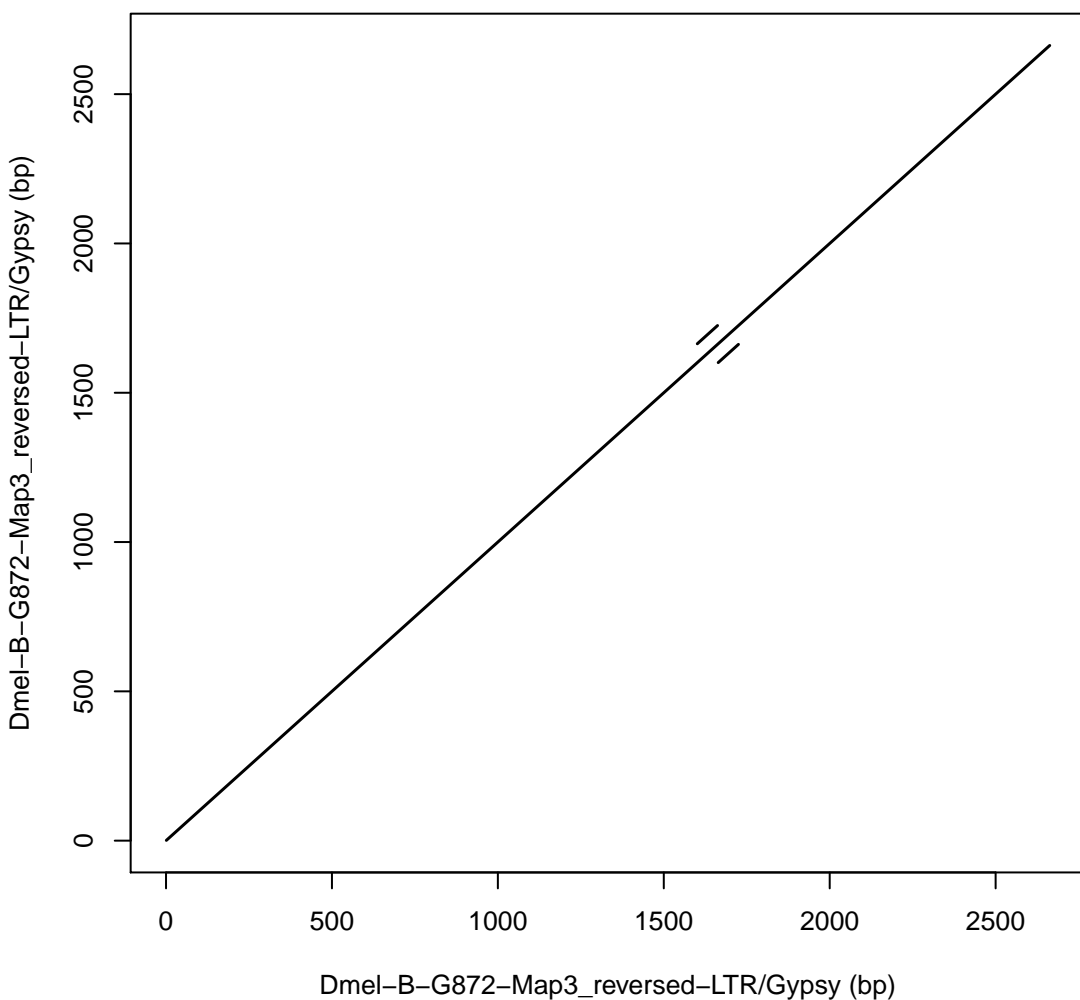
TE: Dmel-B-G872-Map3\_reversed-LTR/Gypsy  
consensus size: 2663bp; fragments: 163; full length: 0 (>=2396.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

