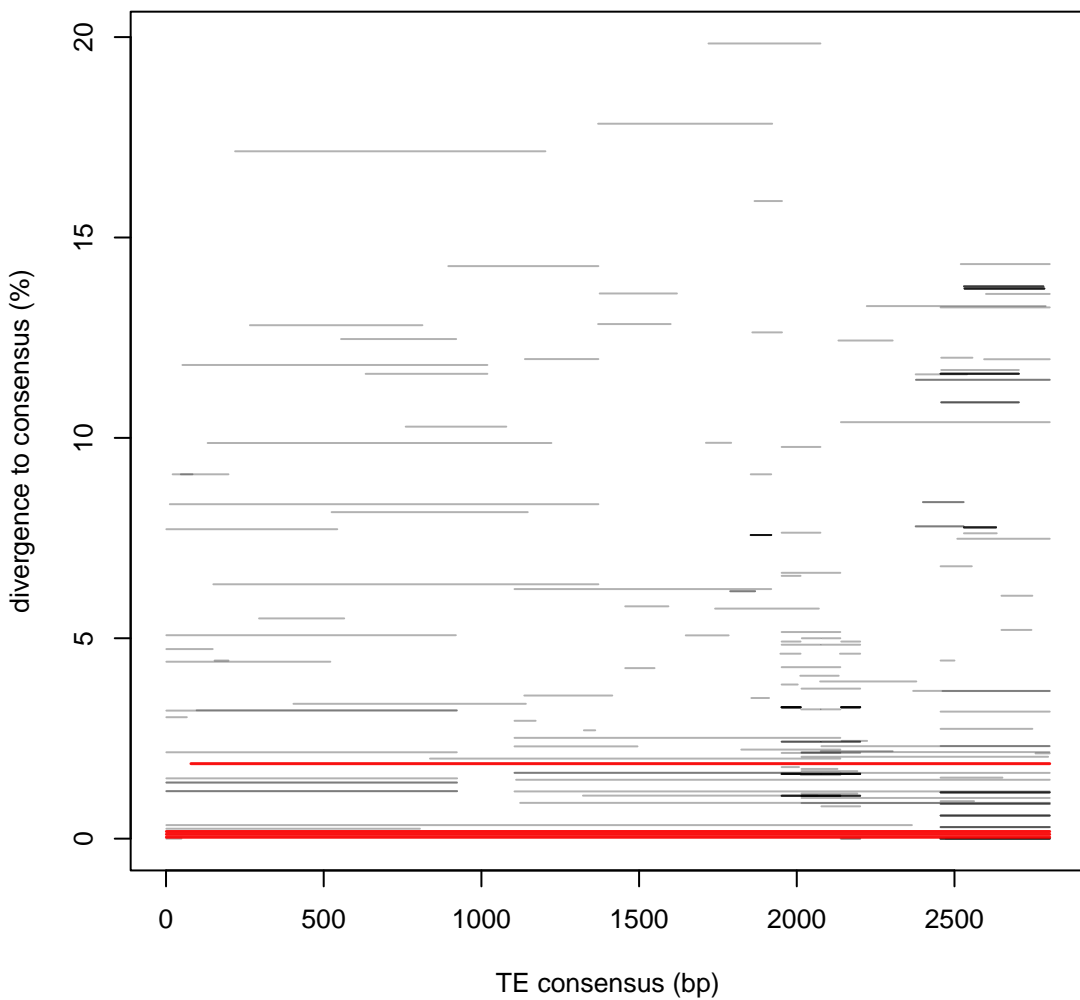
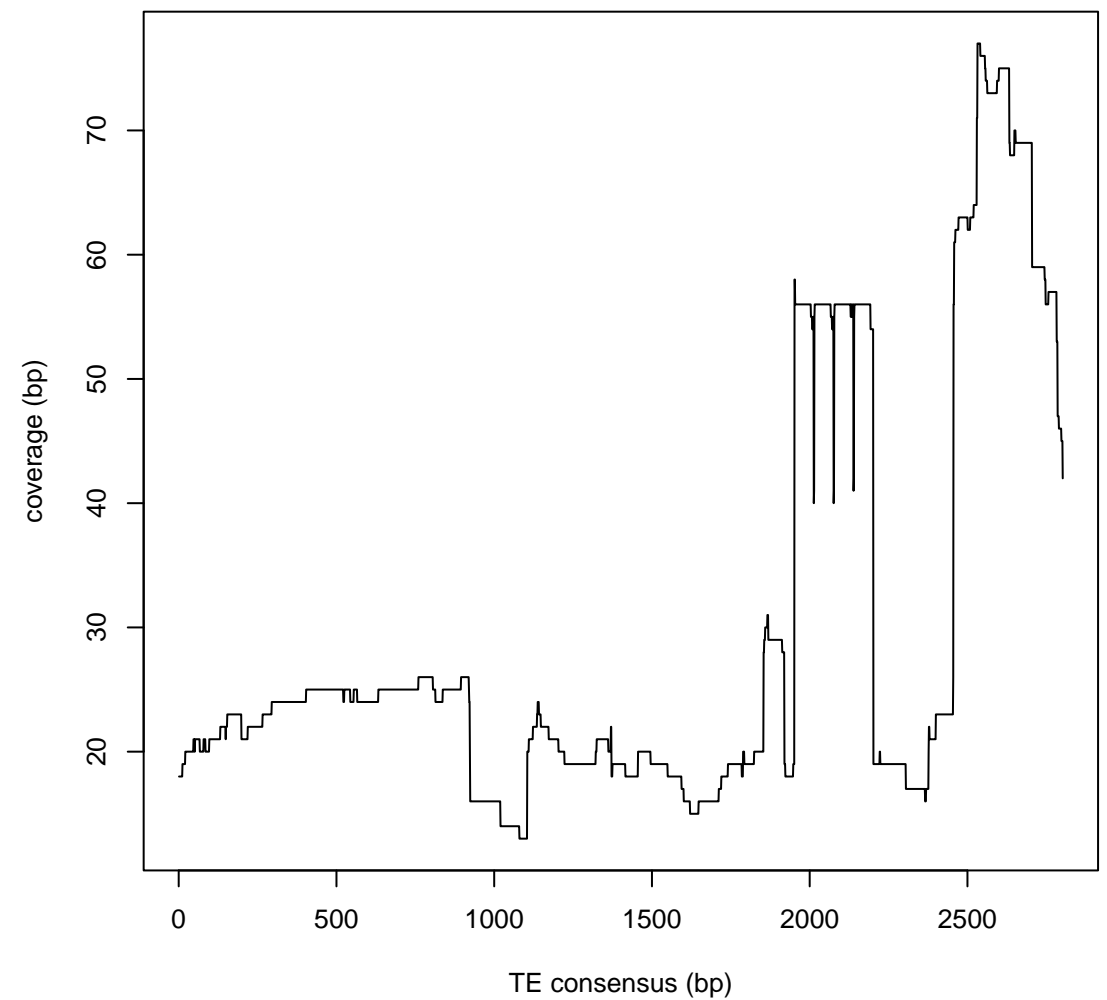


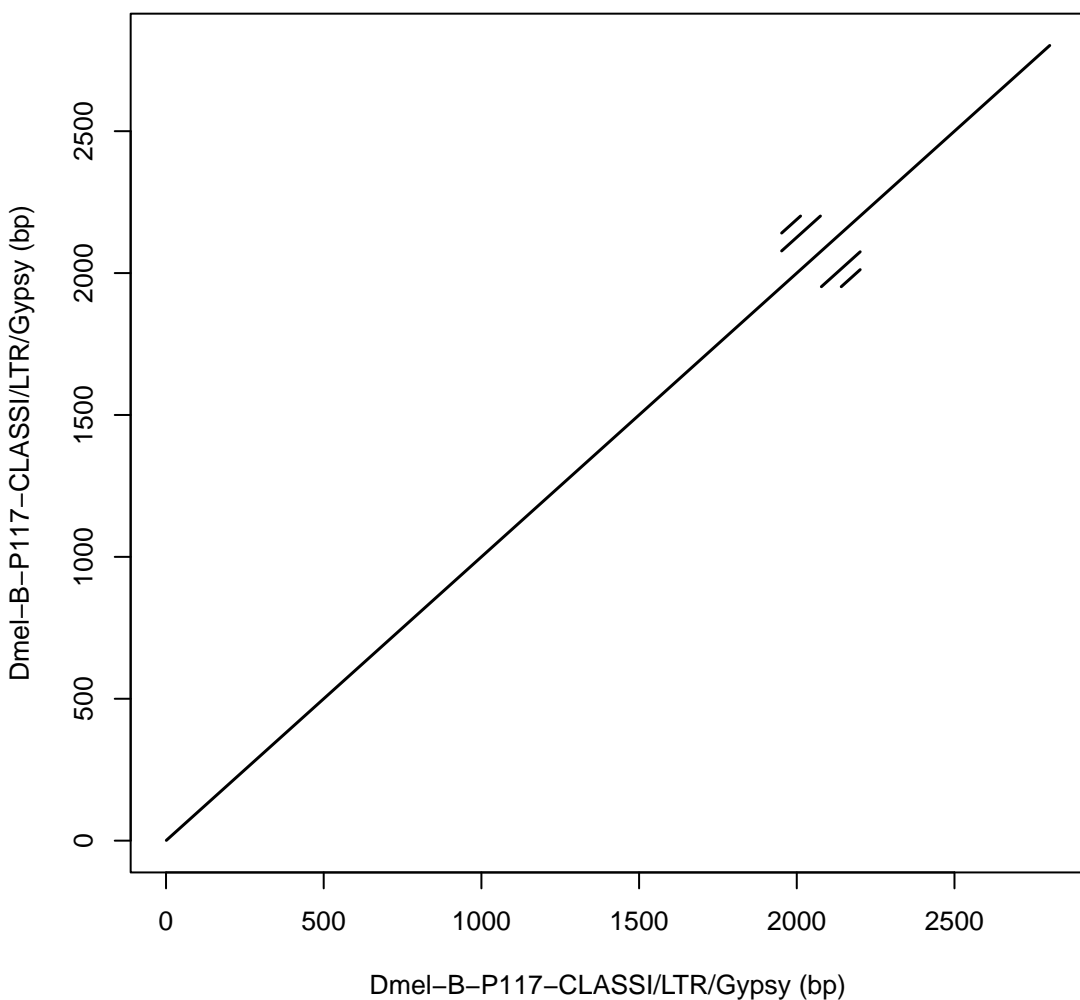
TE: Dmel-B-P117-CLASSI/LTR/Gypsy  
consensus size: 2802bp; fragments: 240; full length: 4 (>=2521.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

