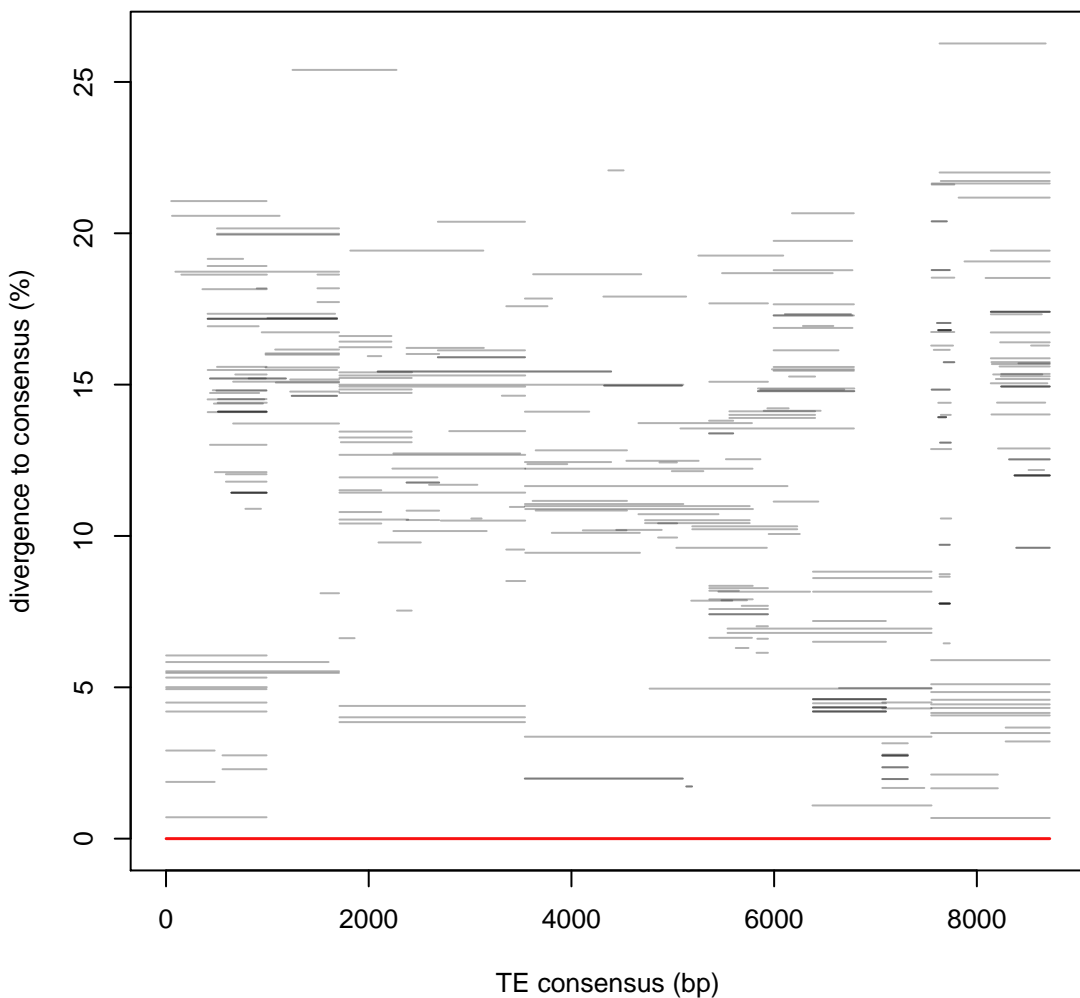
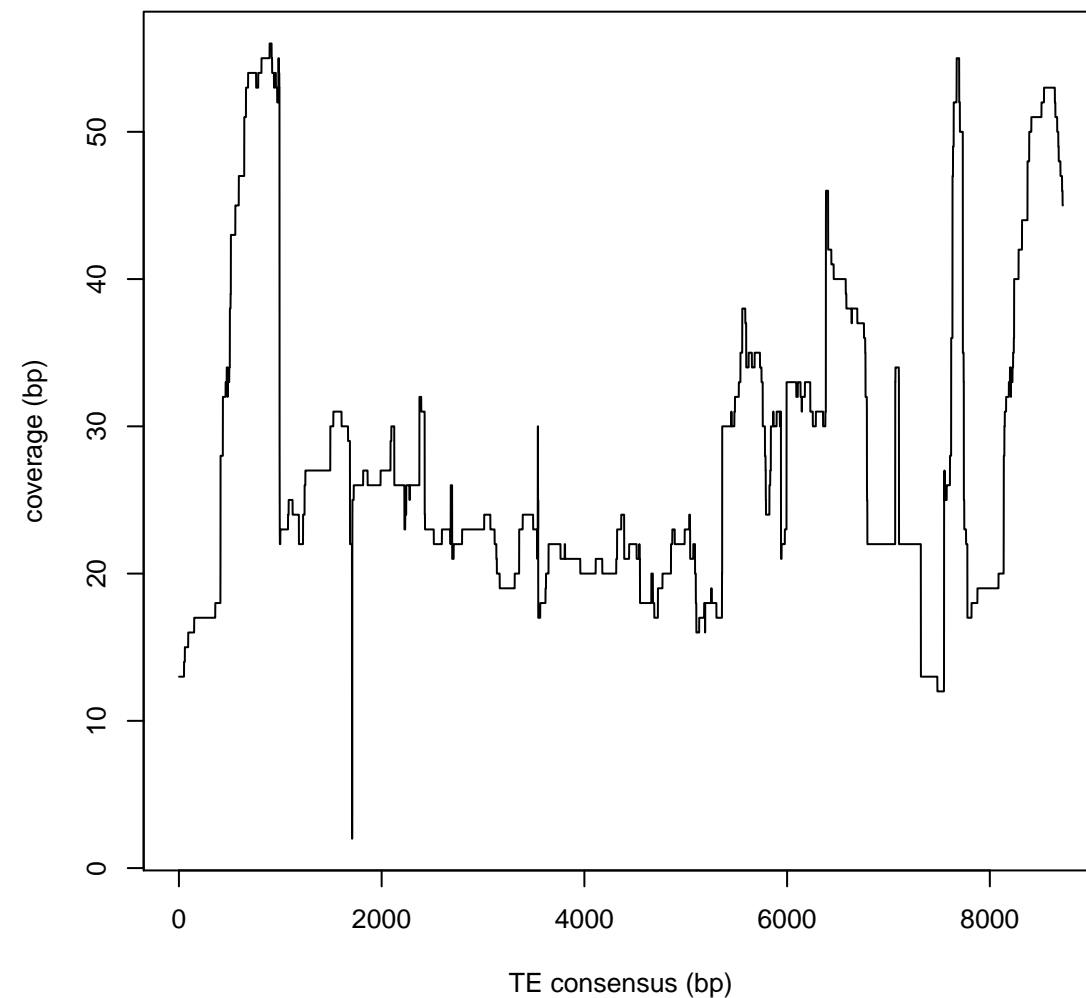


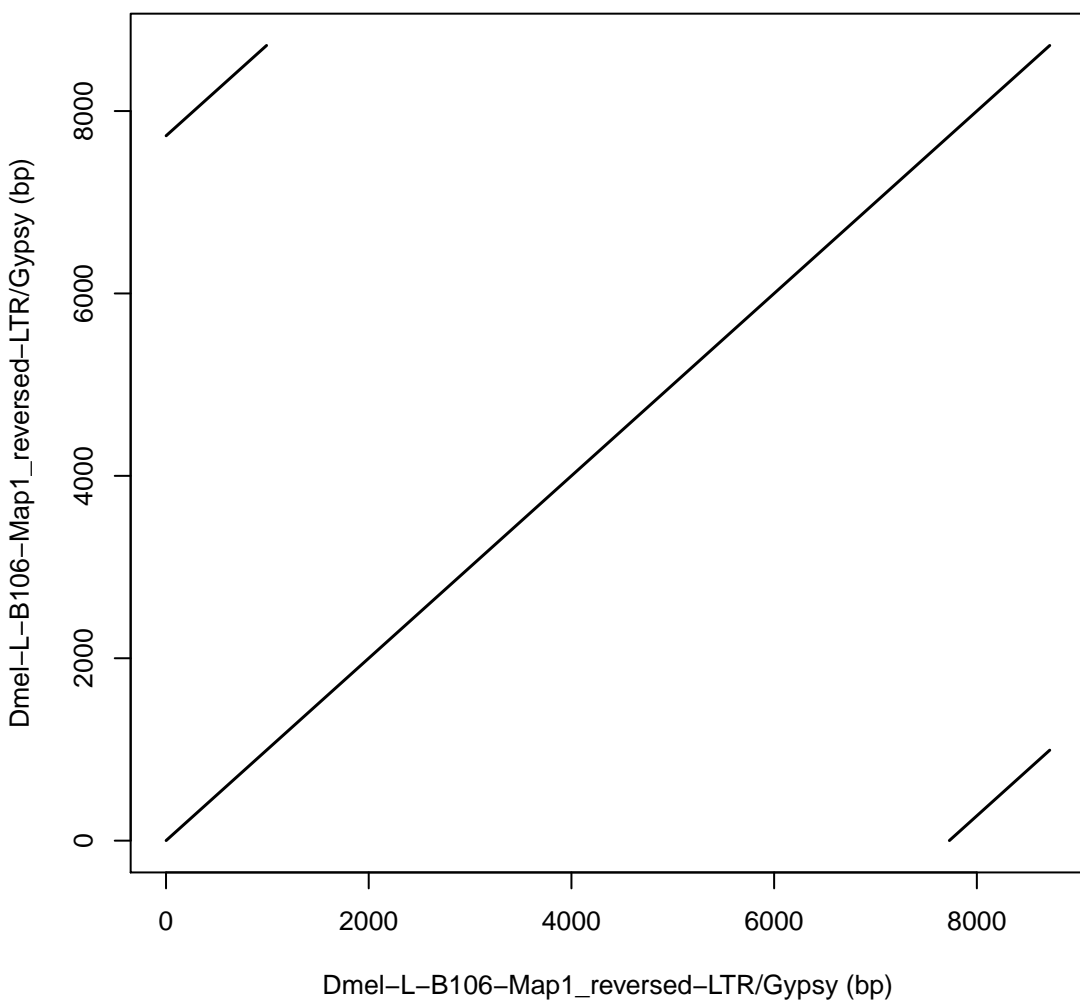
TE: Dmel-L-B106-Map1\_reversed-LTR/Gypsy  
consensus size: 8719bp; fragments: 339; full length: 1 (>=7847.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

