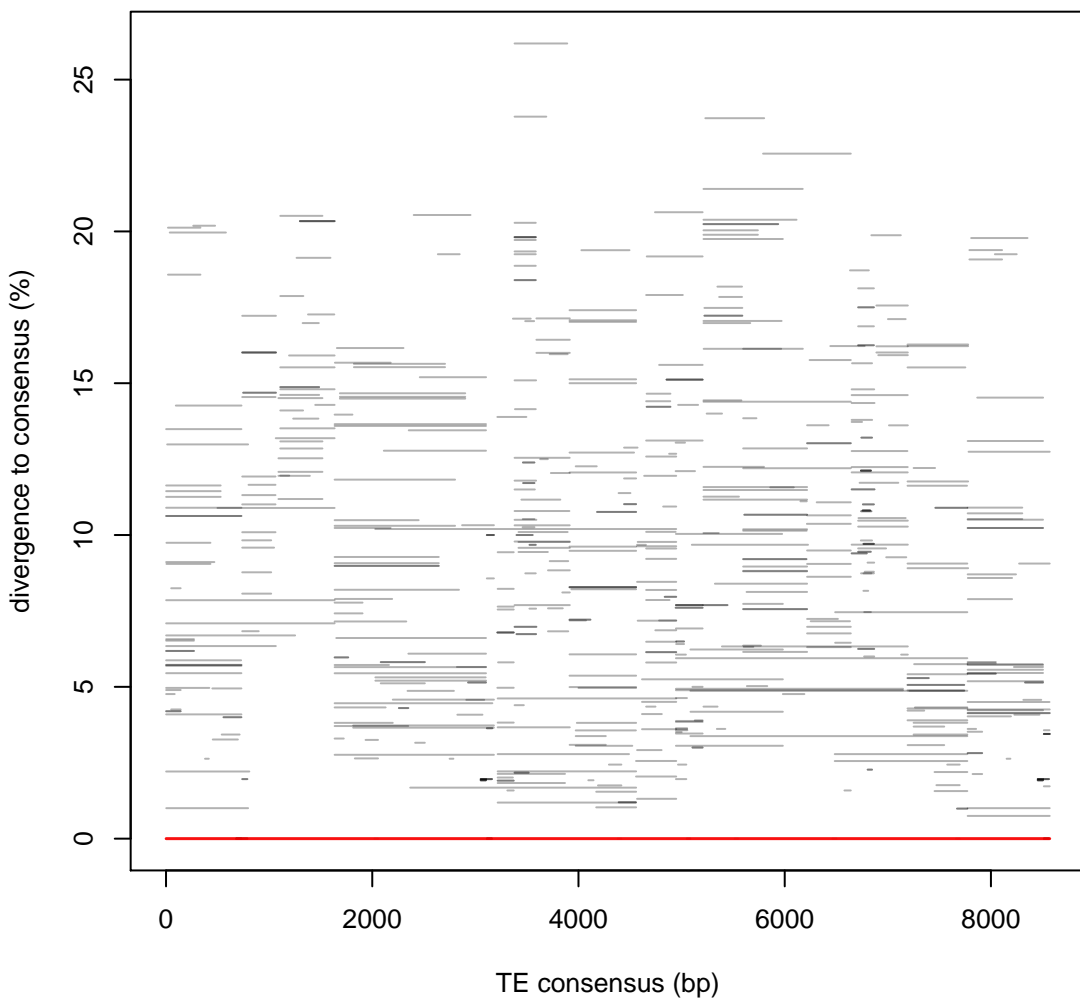
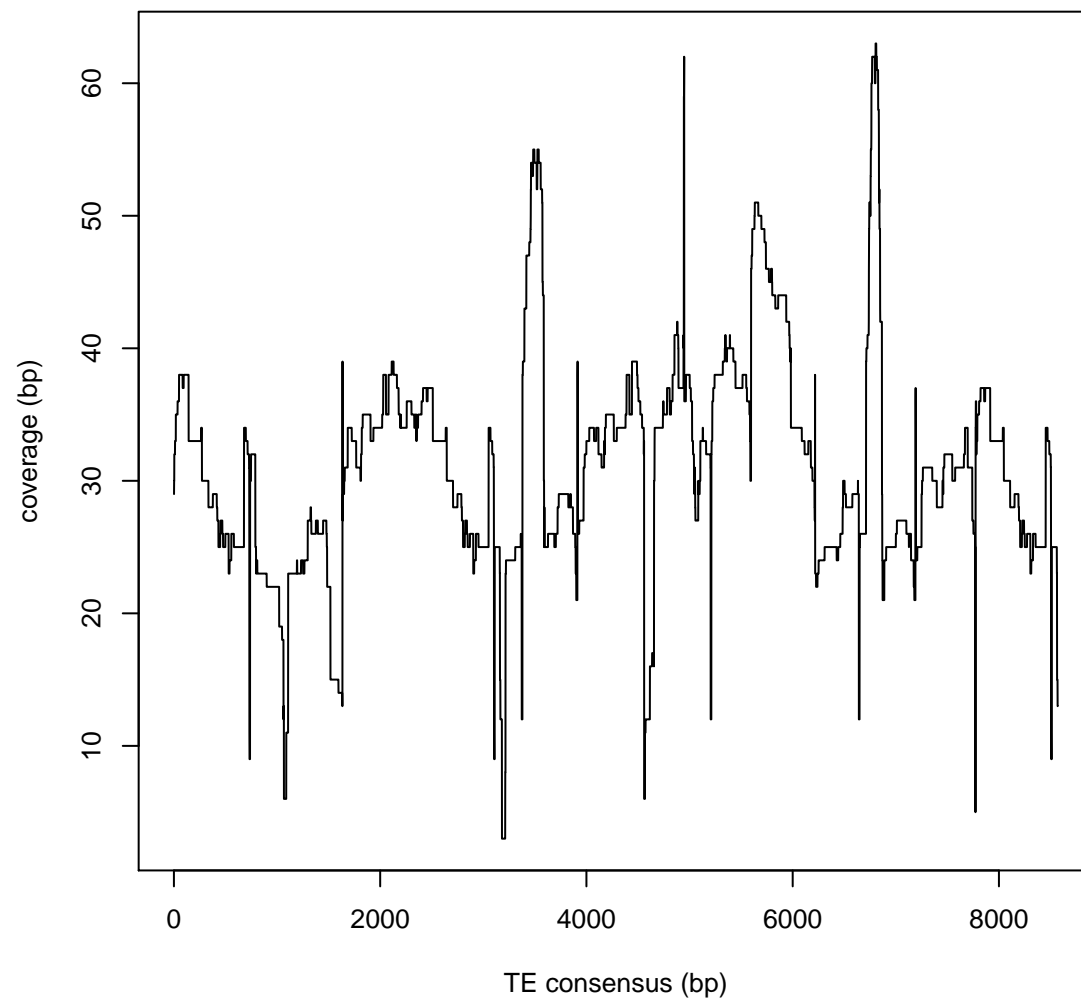


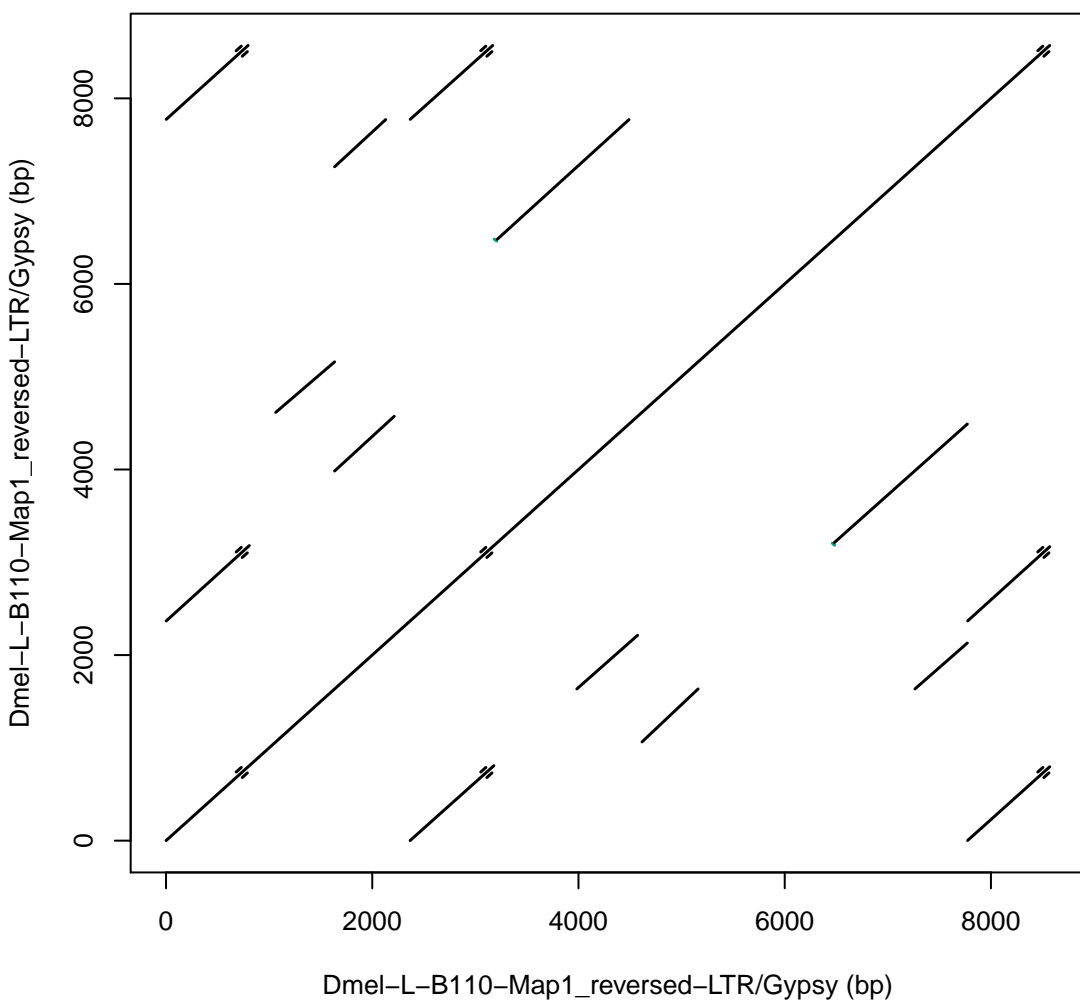
TE: Dmel-L-B110-Map1\_reversed-LTR/Gypsy  
consensus size: 8570bp; fragments: 649; full length: 1 (>=7713bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

