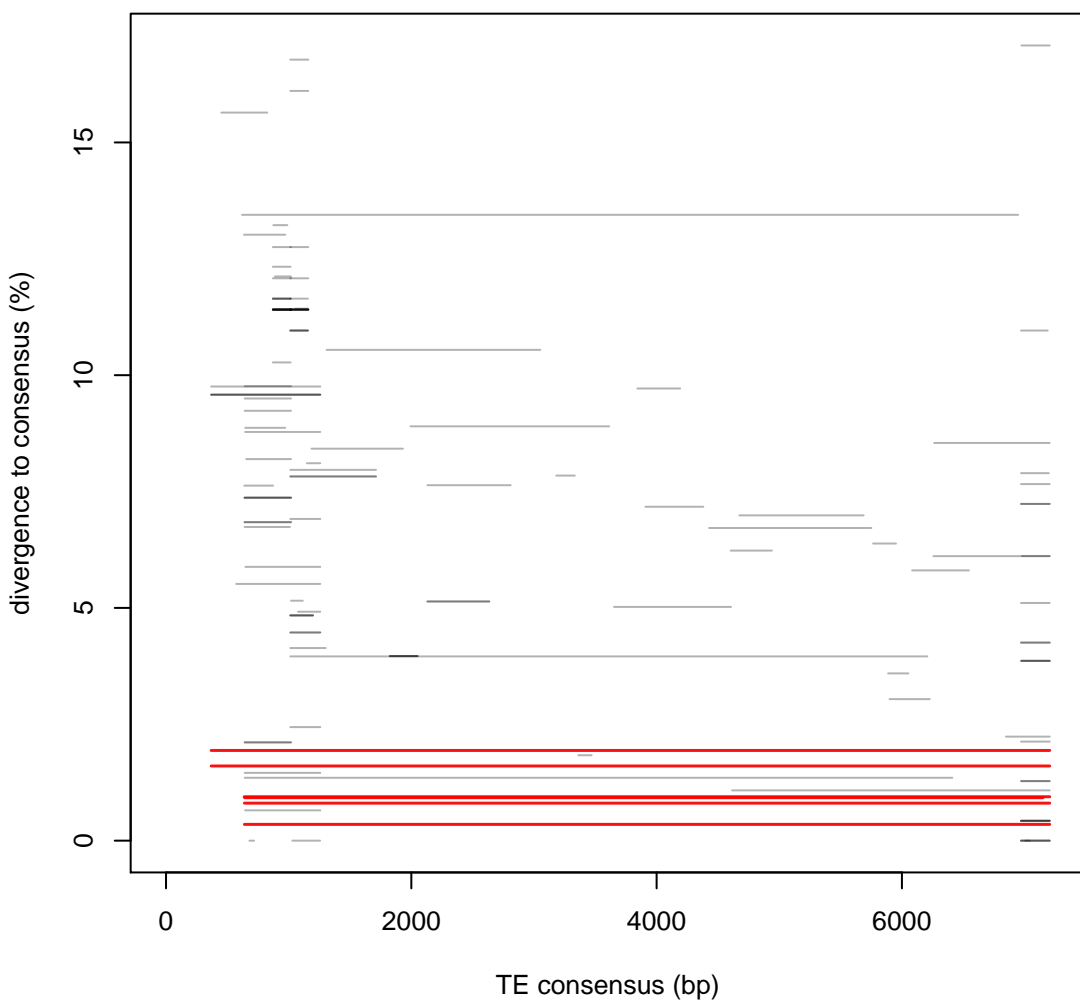
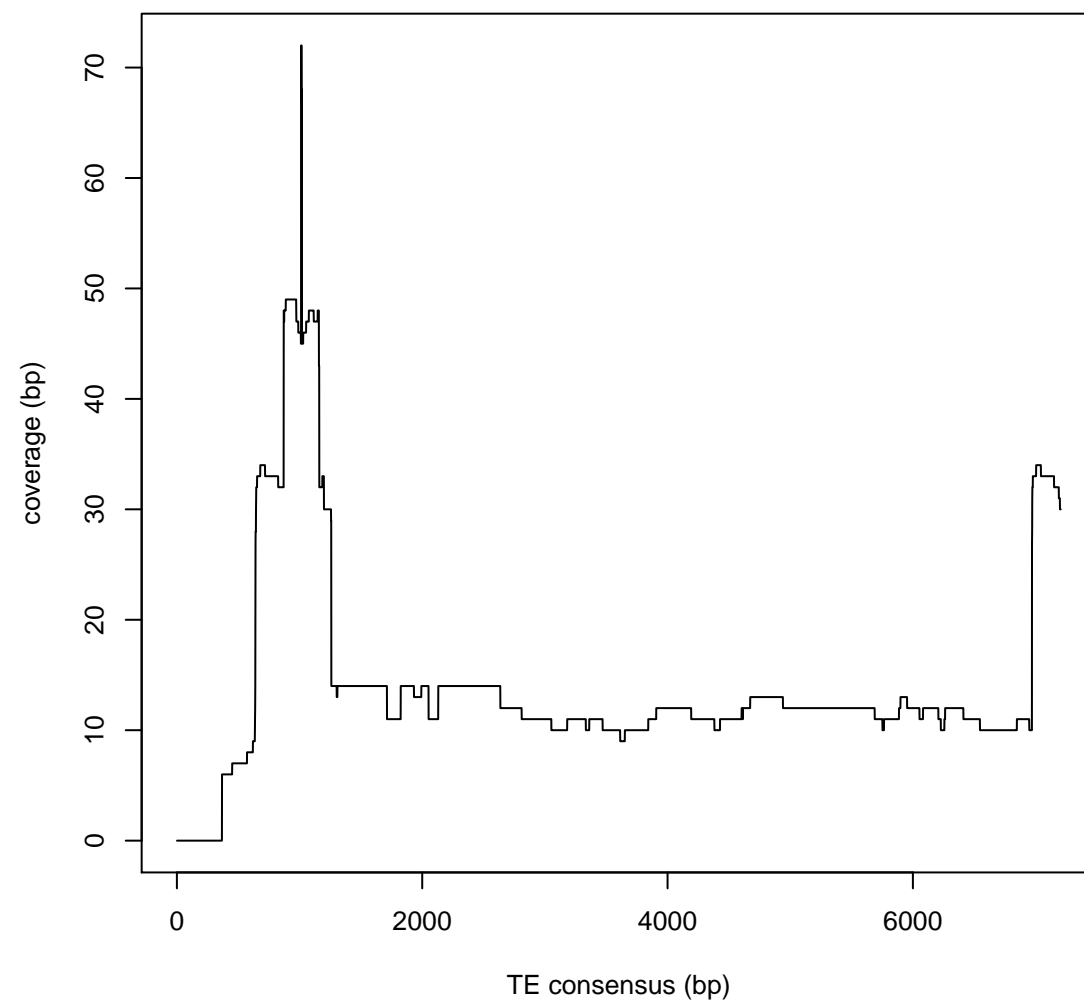


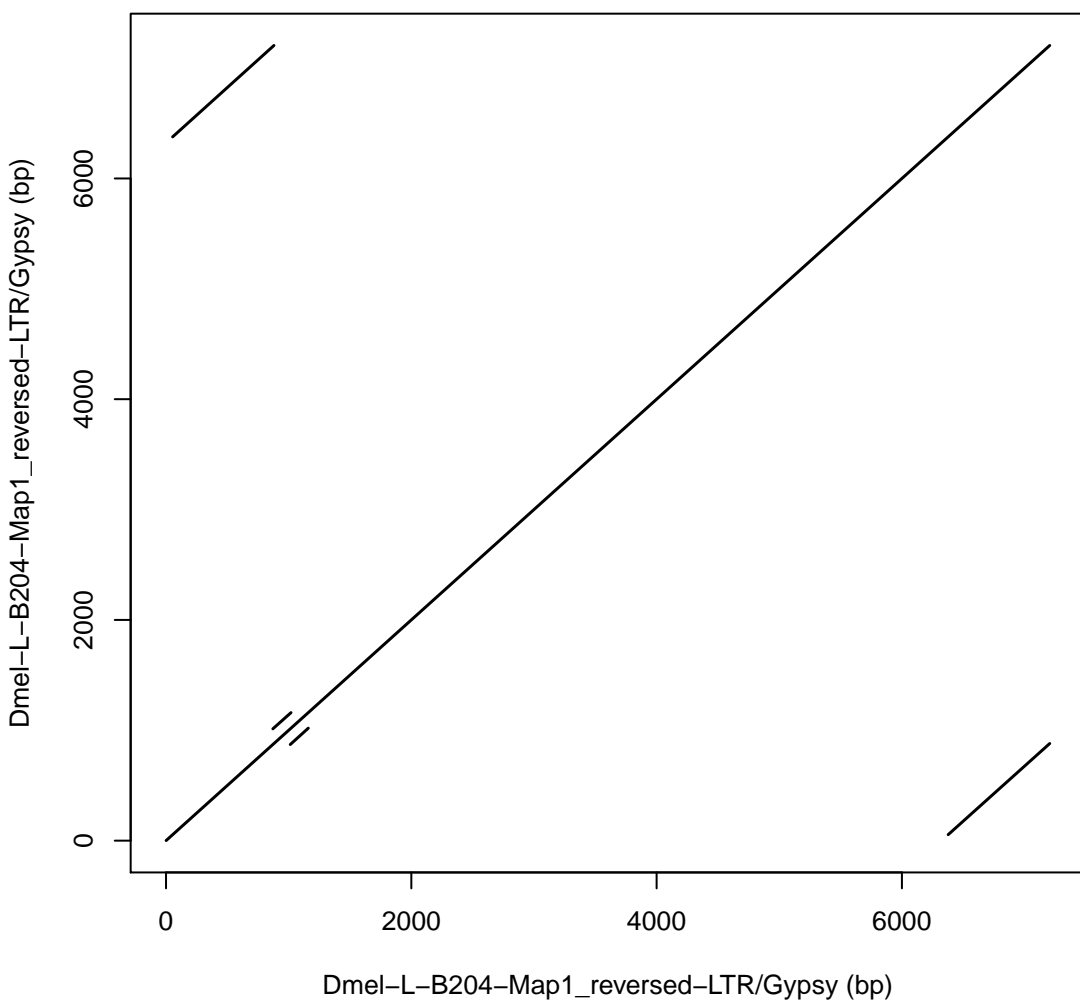
TE: Dmel-L-B204-Map1\_reversed-LTR/Gypsy  
consensus size: 7205bp; fragments: 133; full length: 6 ( $\geq 6484.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

