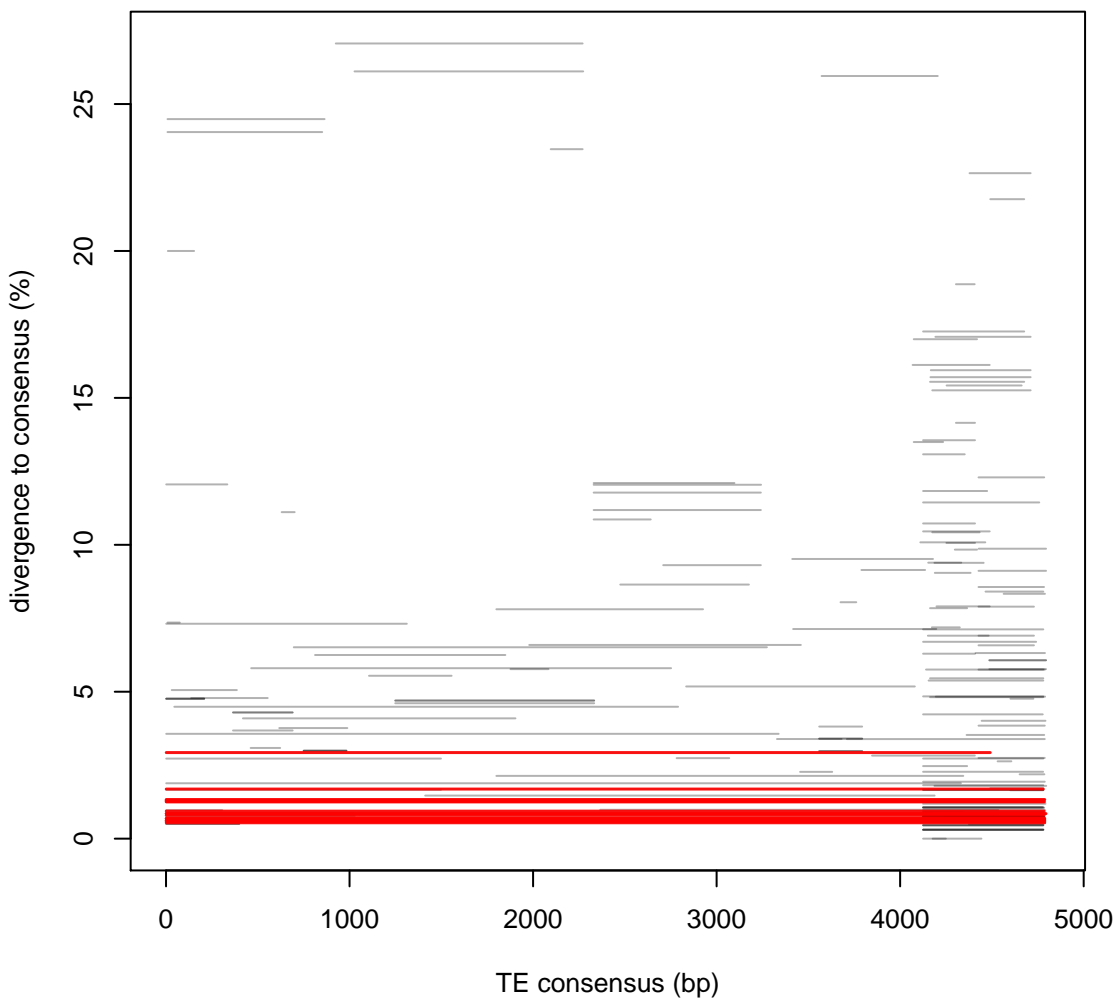
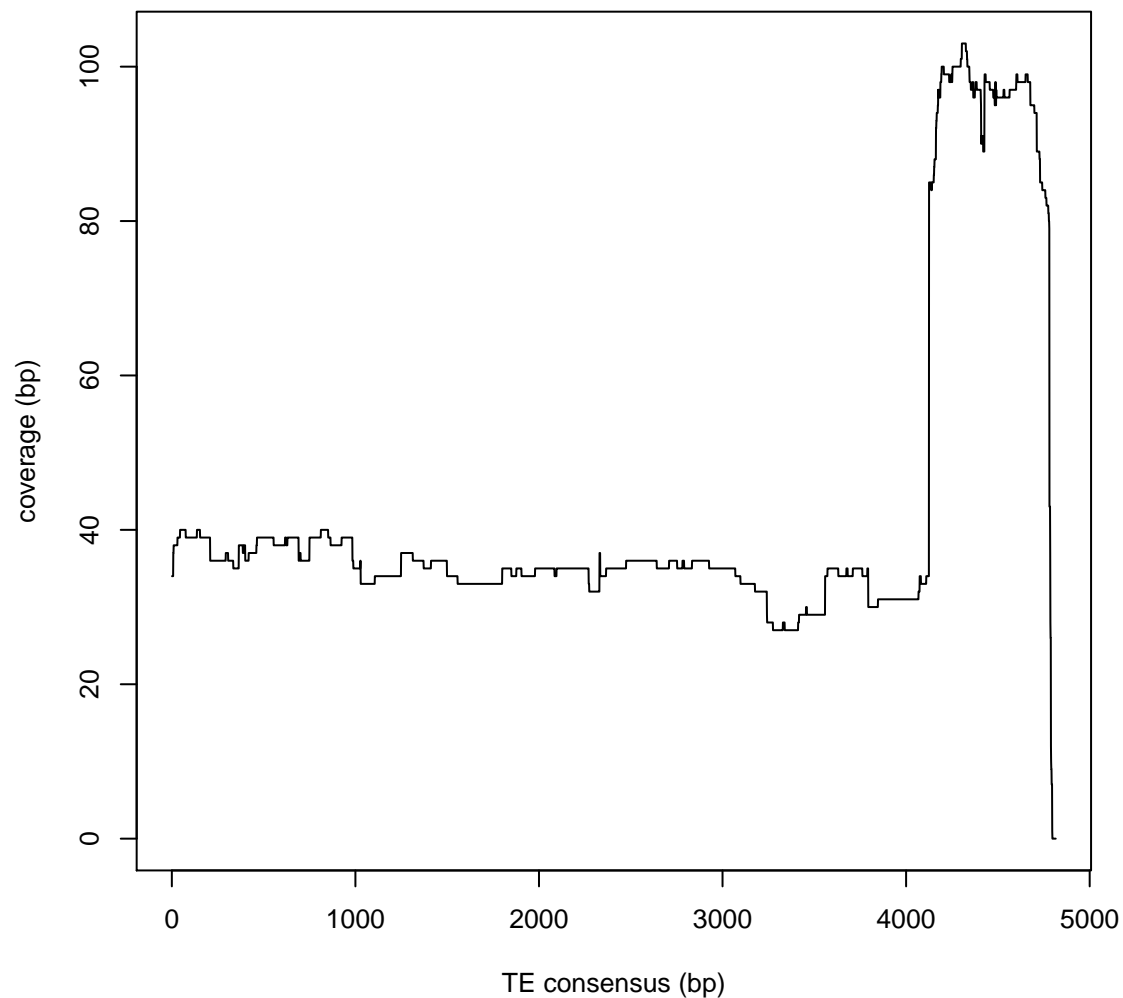


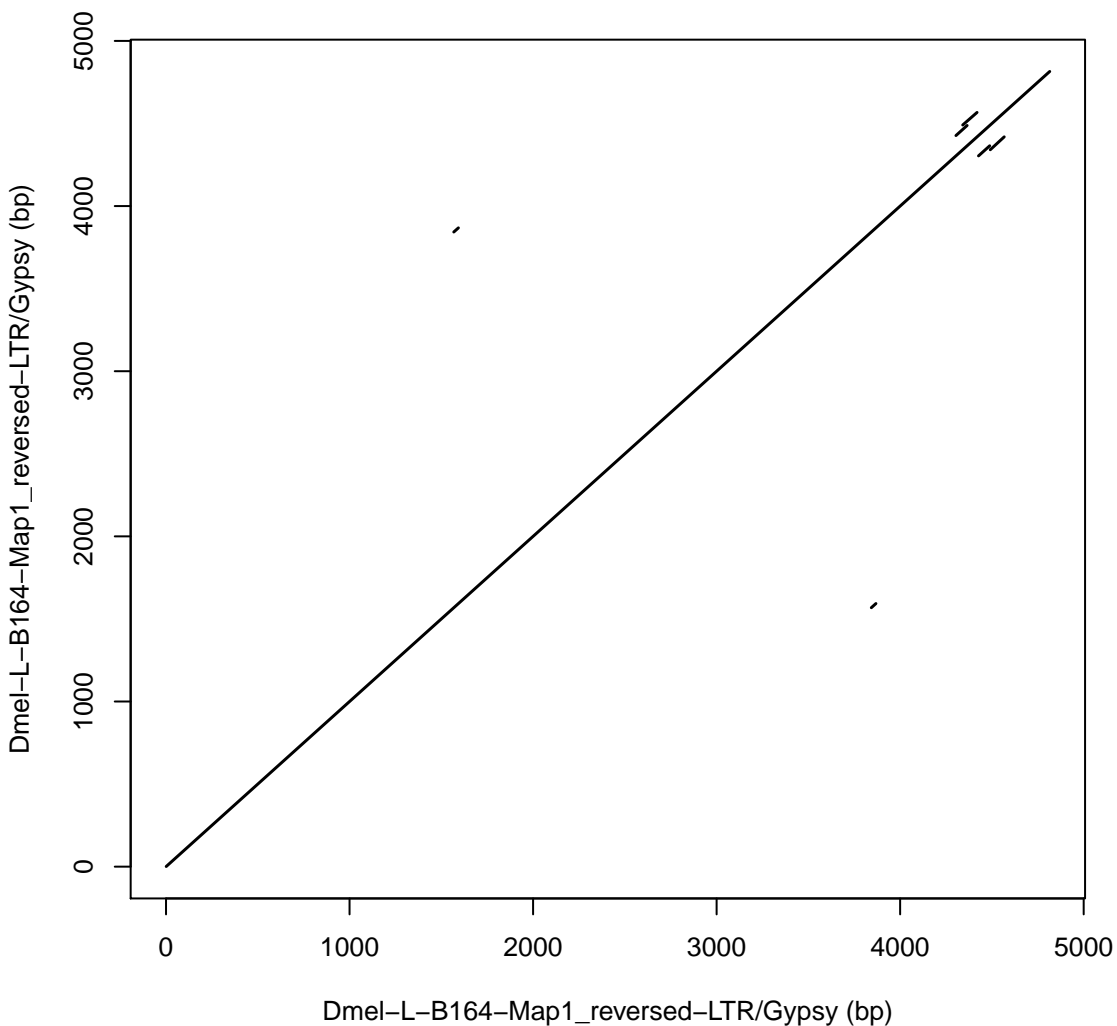
TE: Dmel-L-B164-Map1\_reversed-LTR/Gypsy  
consensus size: 4815bp; fragments: 199; full length: 19 ( $\geq 4333.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

