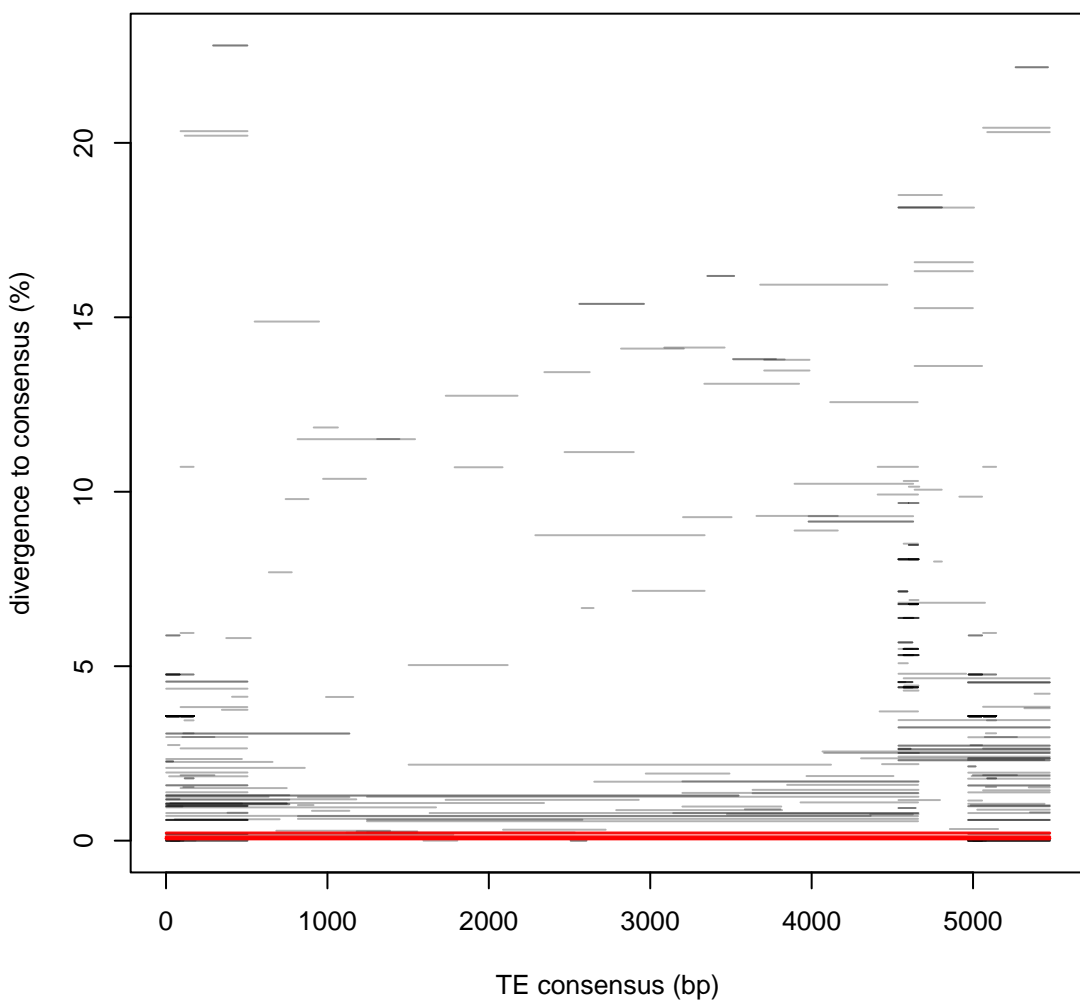
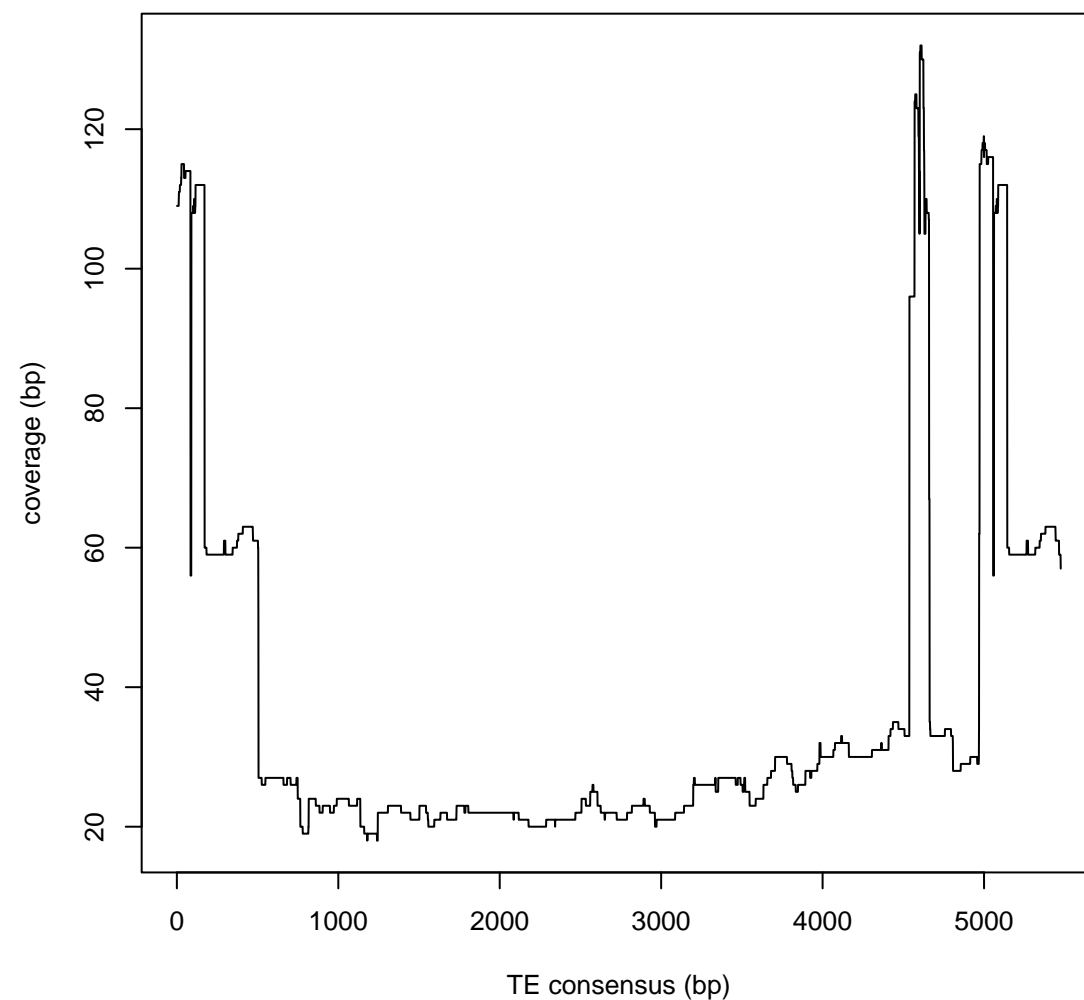


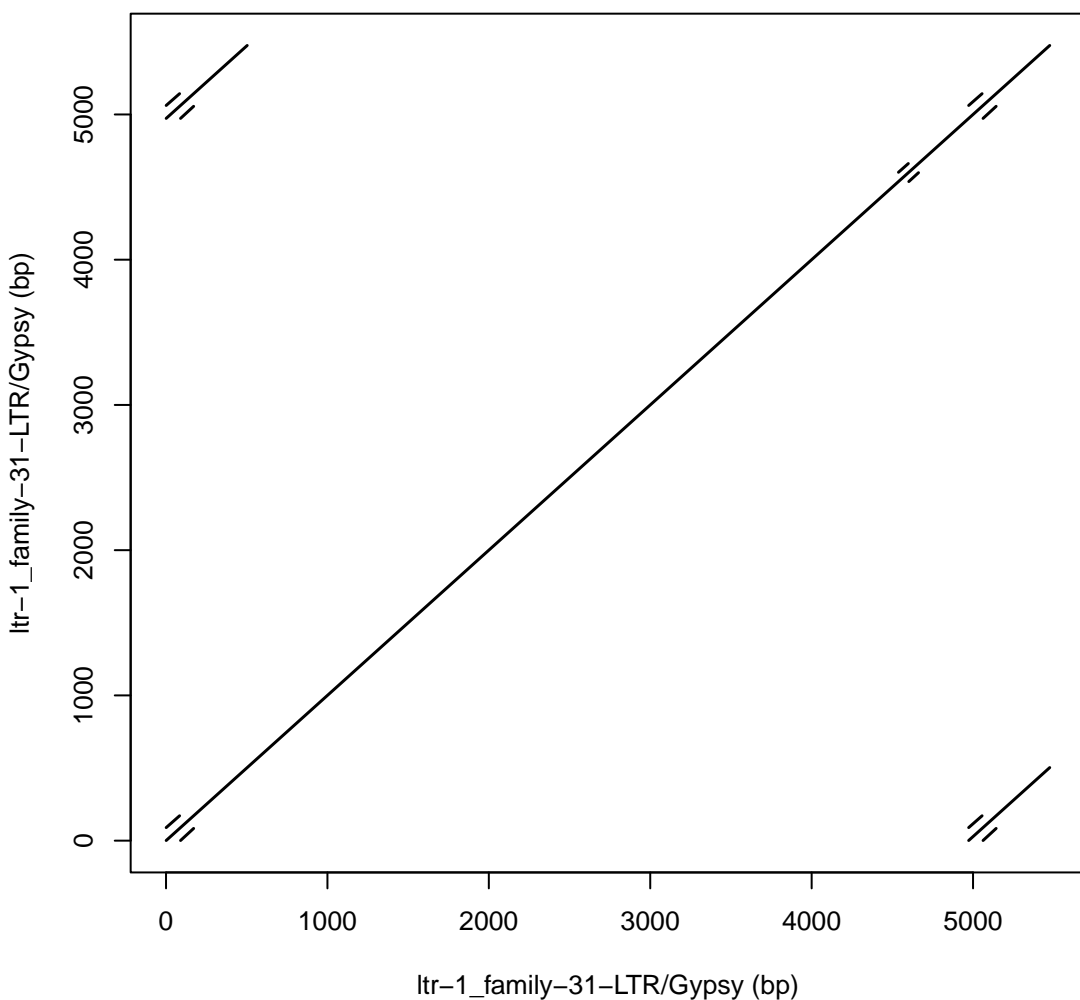
TE: ltr-1\_family-31-LTR/Gypsy  
consensus size: 5475bp; fragments: 543; full length: 6 ( $\geq 4927.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

