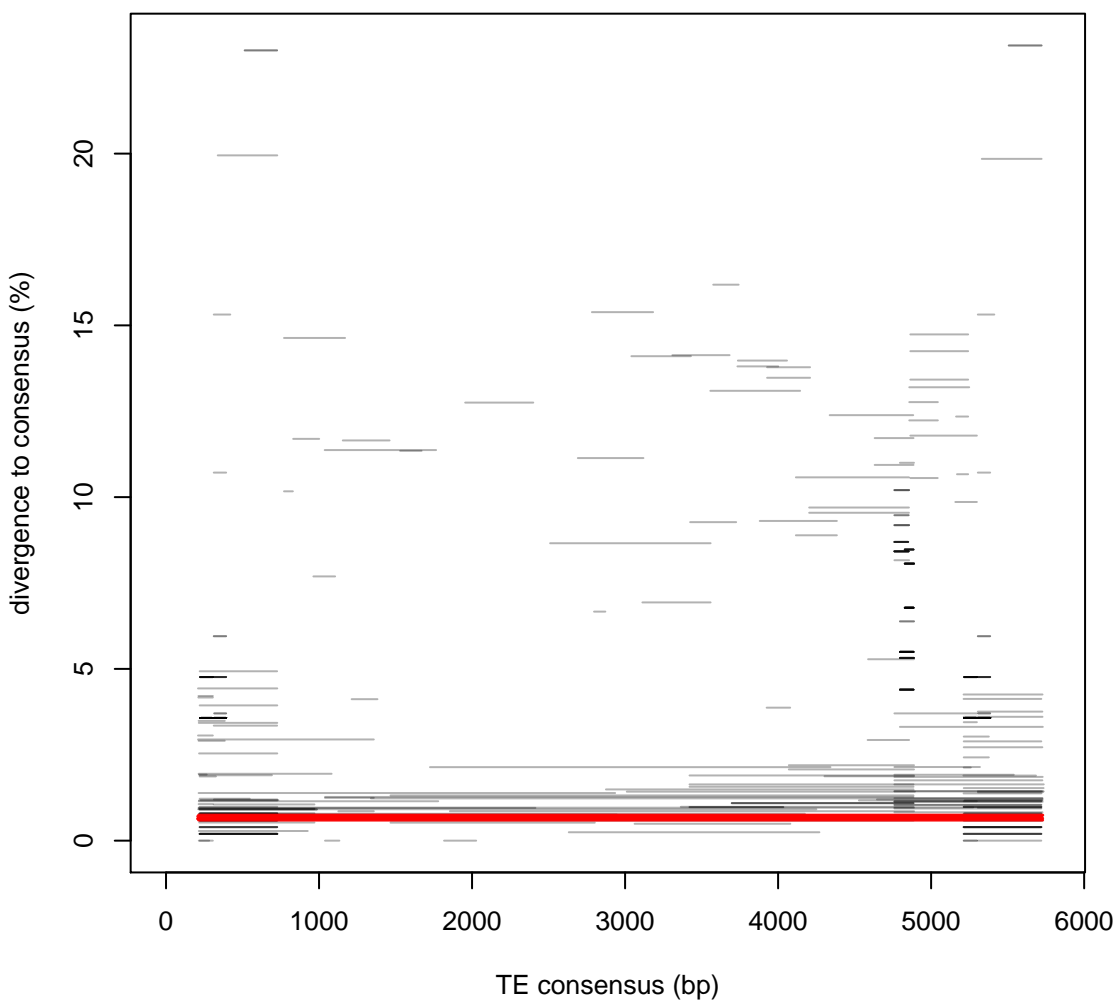
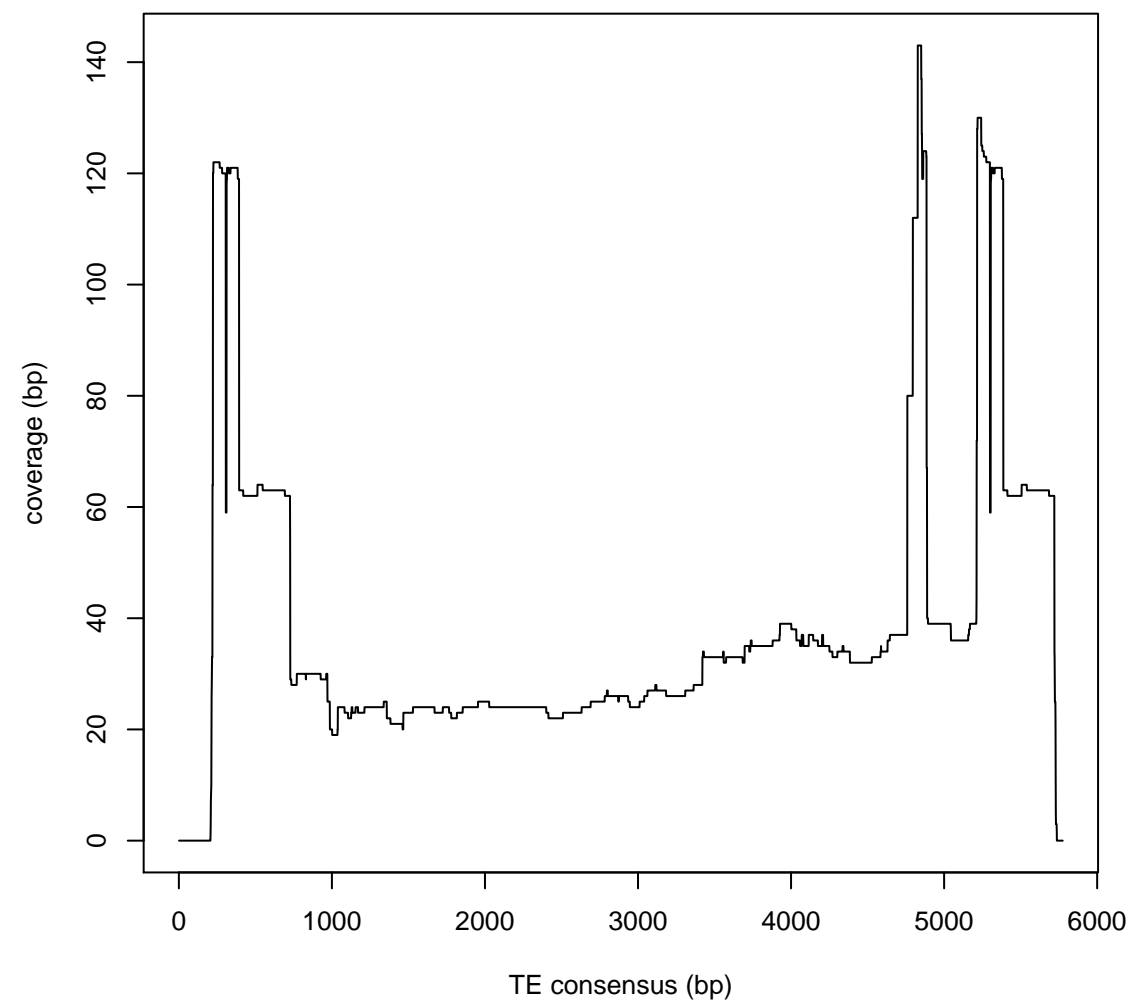


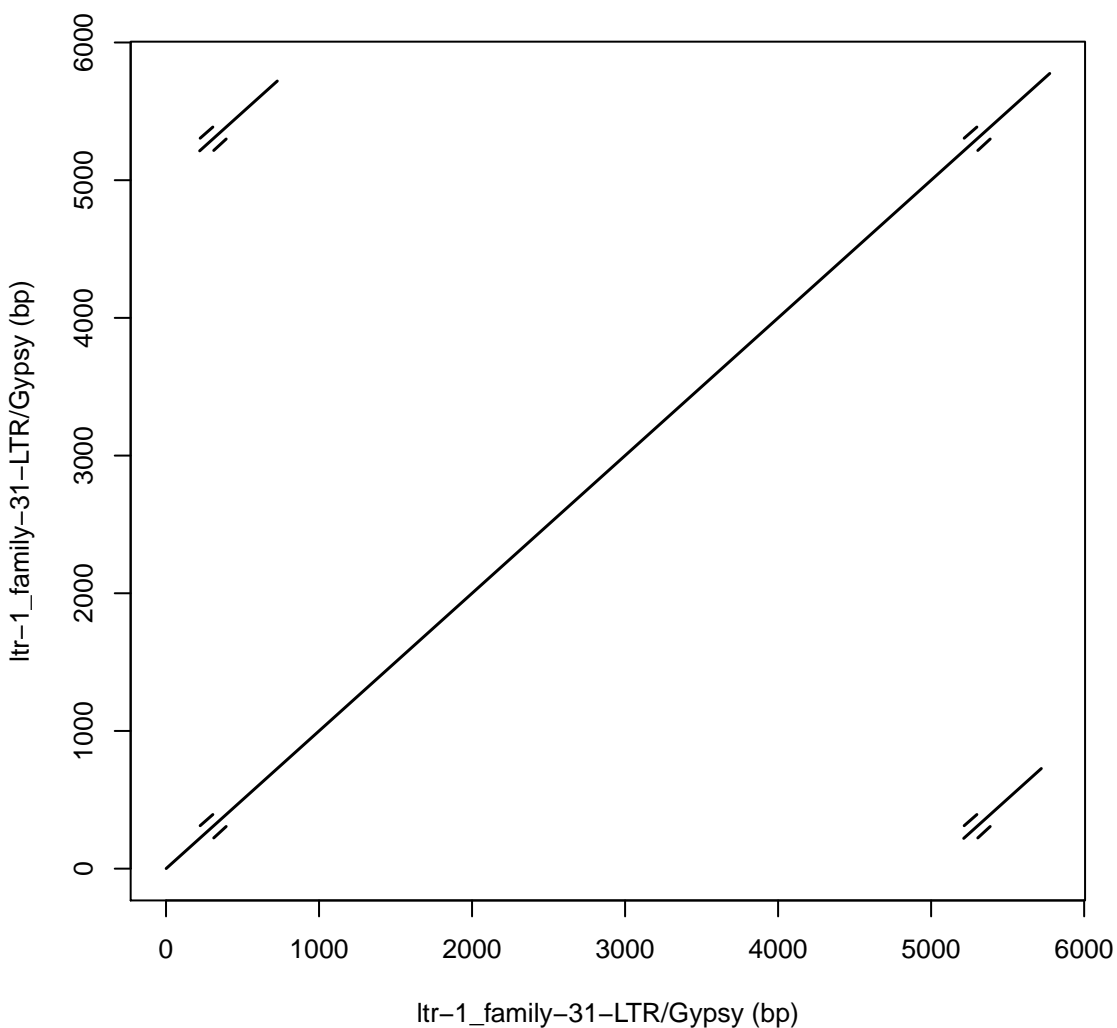
TE: ltr-1_family-31-LTR/Gypsy
consensus size: 5775bp; fragments: 525; full length: 7 (≥ 5197.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

