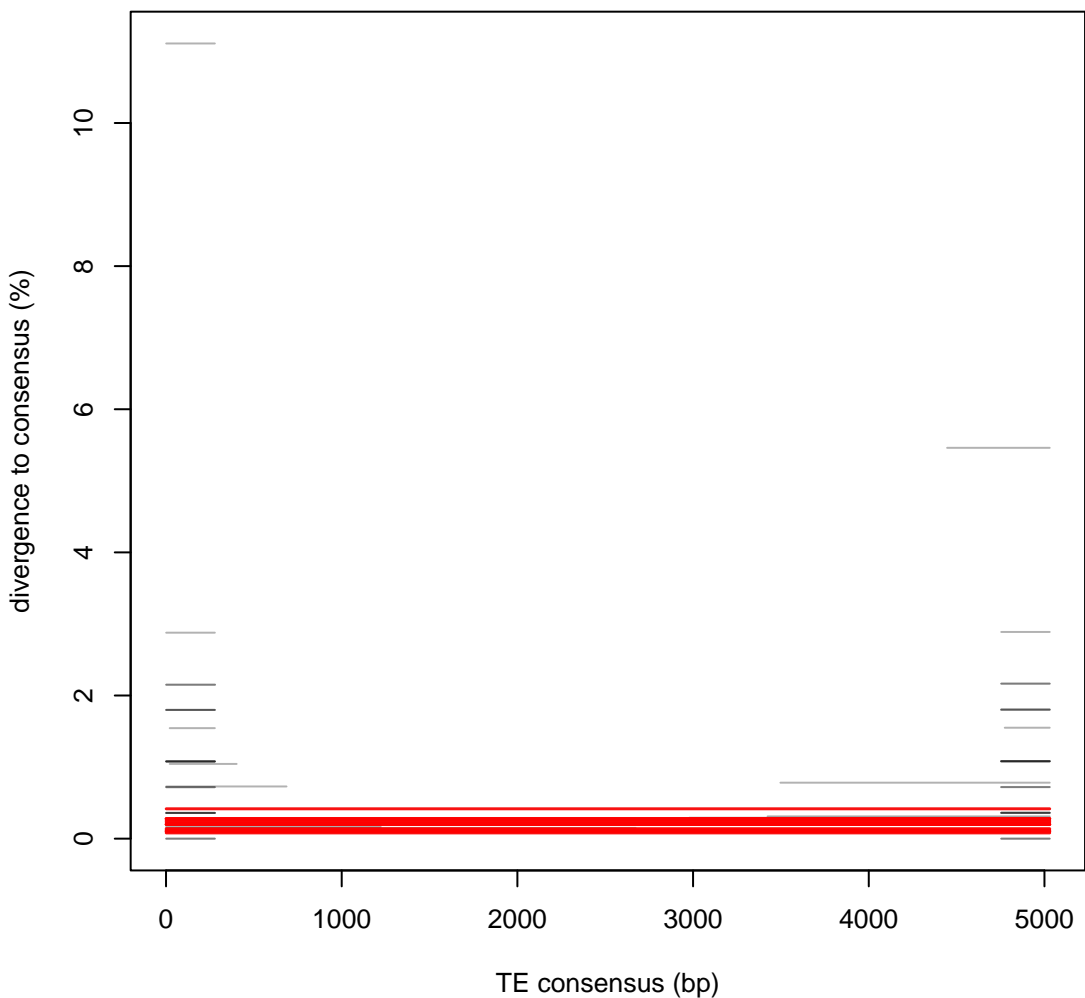
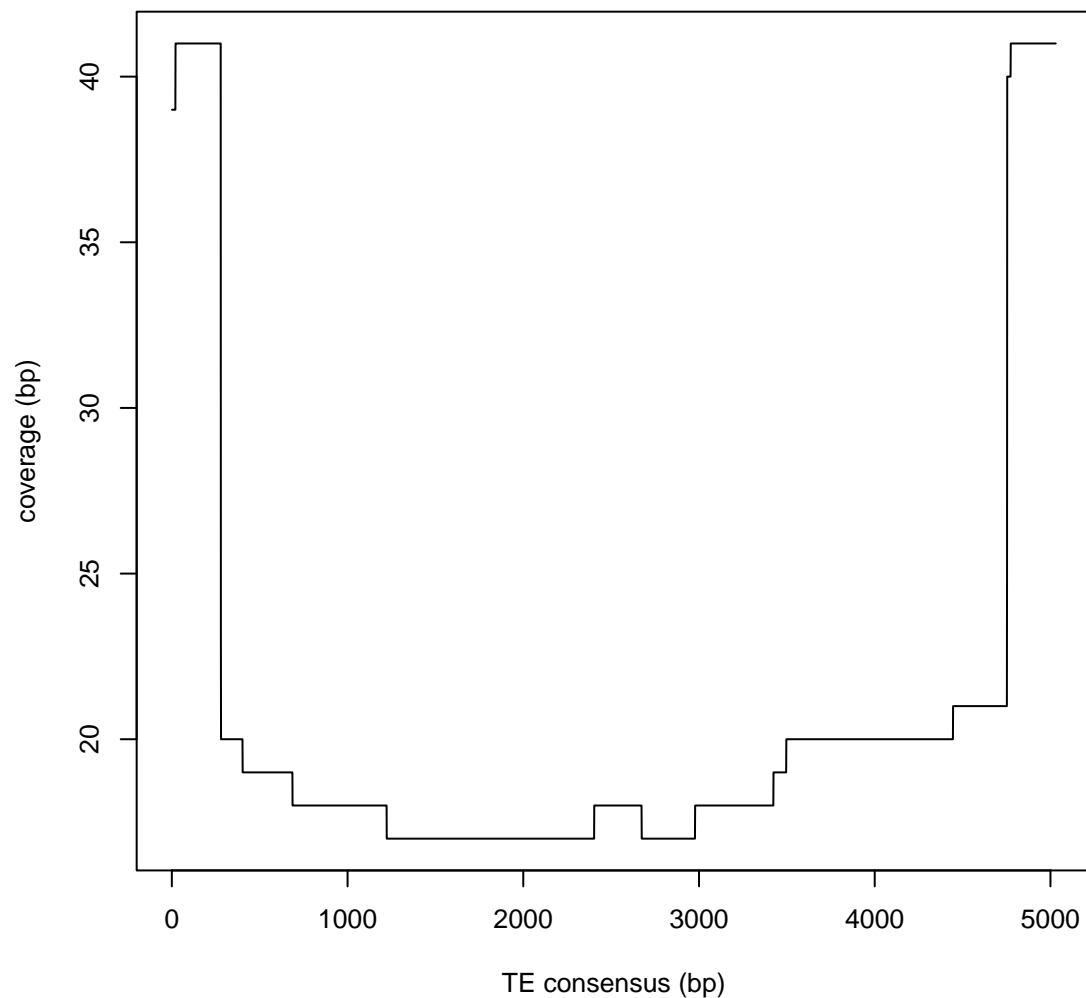


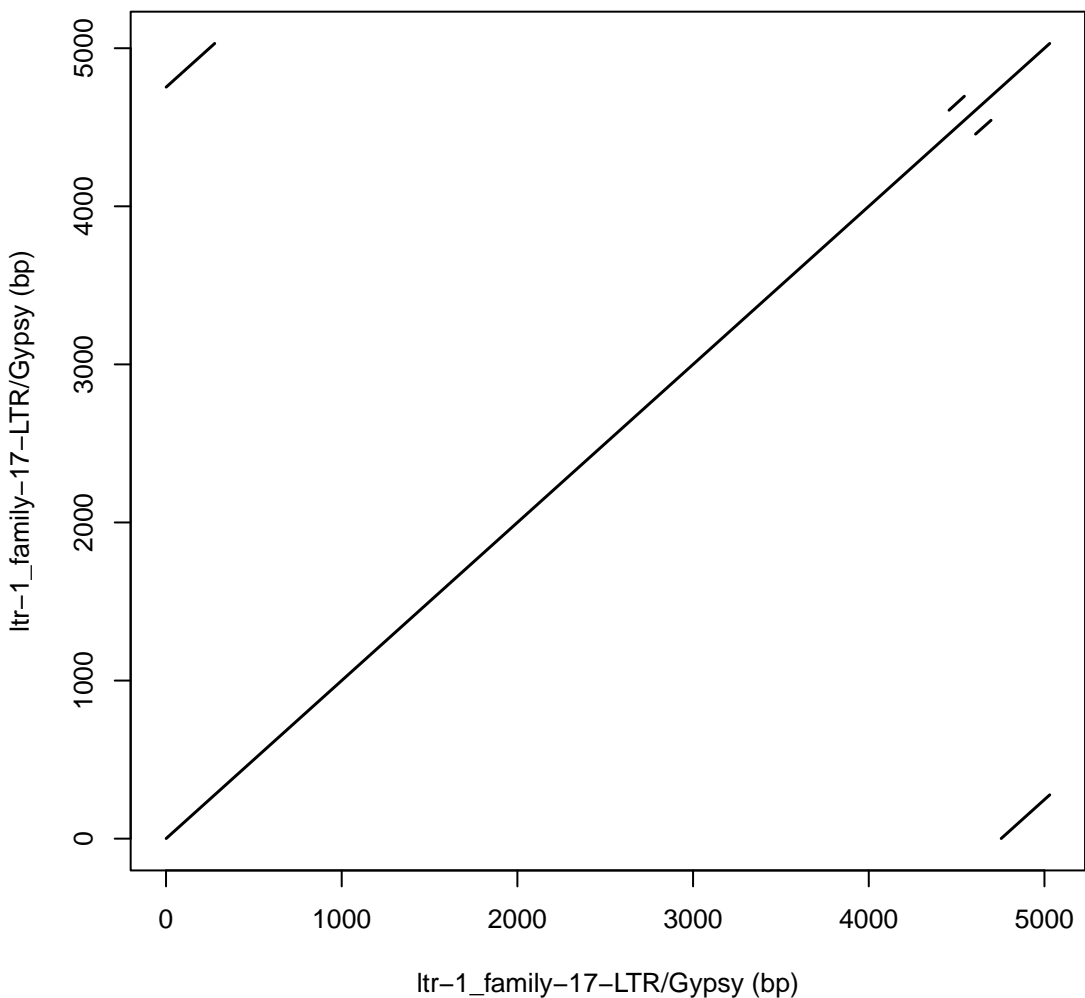
TE: ltr-1_family-17-LTR/Gypsy
consensus size: 5030bp; fragments: 66; full length: 16 (≥ 4527 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

