TE: rnd−1_family−110−LTR/GYPSY
consensus size: 3615bp; fragments: 796; full length: 2 (>=3253.5bp)

divergence to consensus (%)
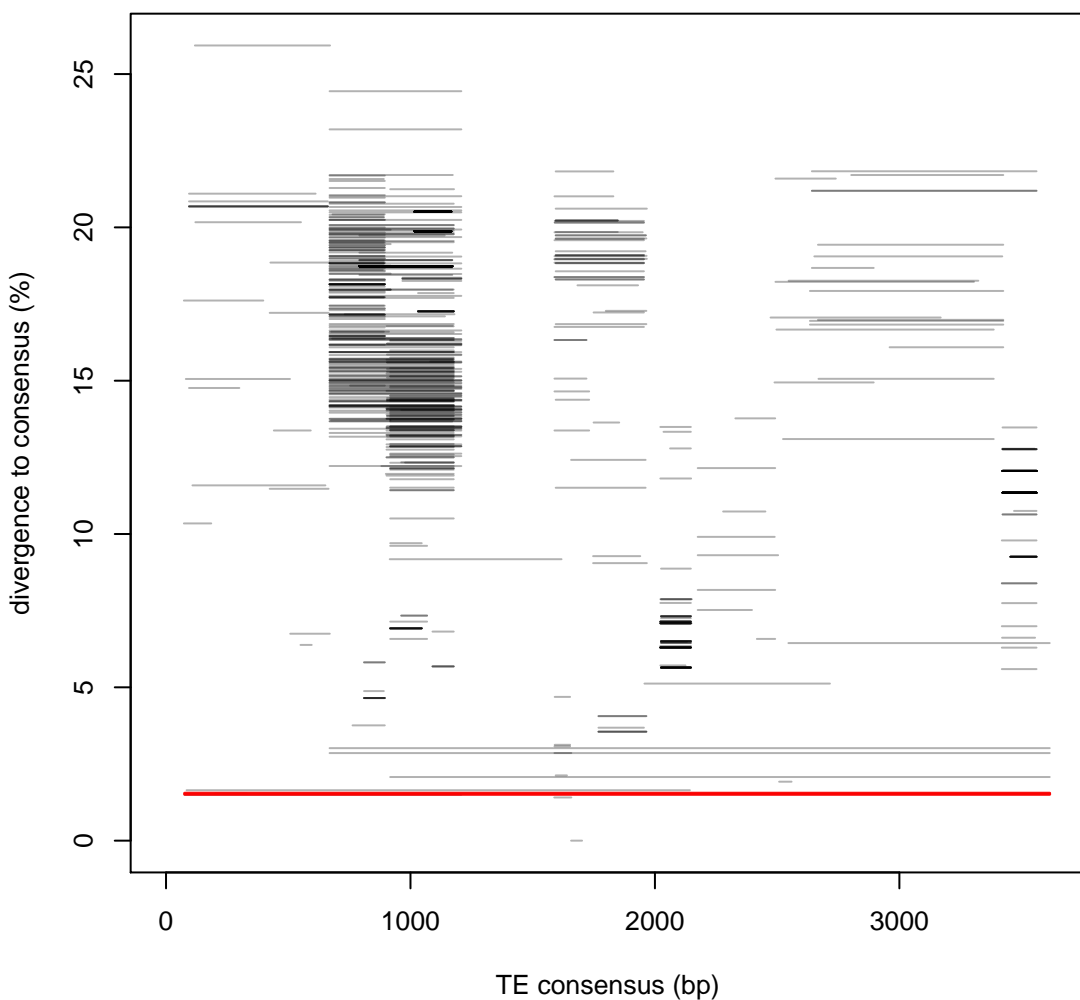
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
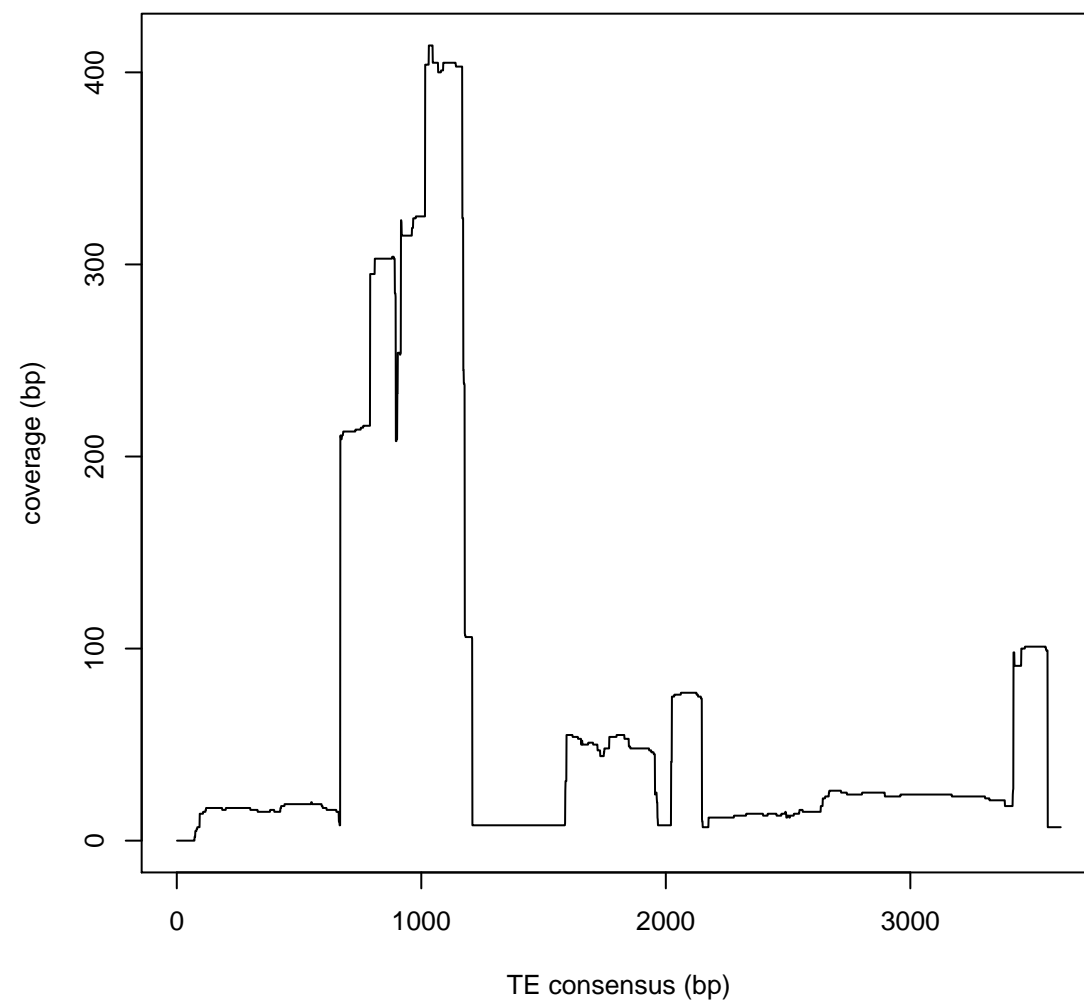
TE consensus (bp)

TE consensus self dotplot (blastn)

rnd−1_family−110−LTR/GYPSY (bp)

rnd−1_family−110−LTR/GYPSY (bp)
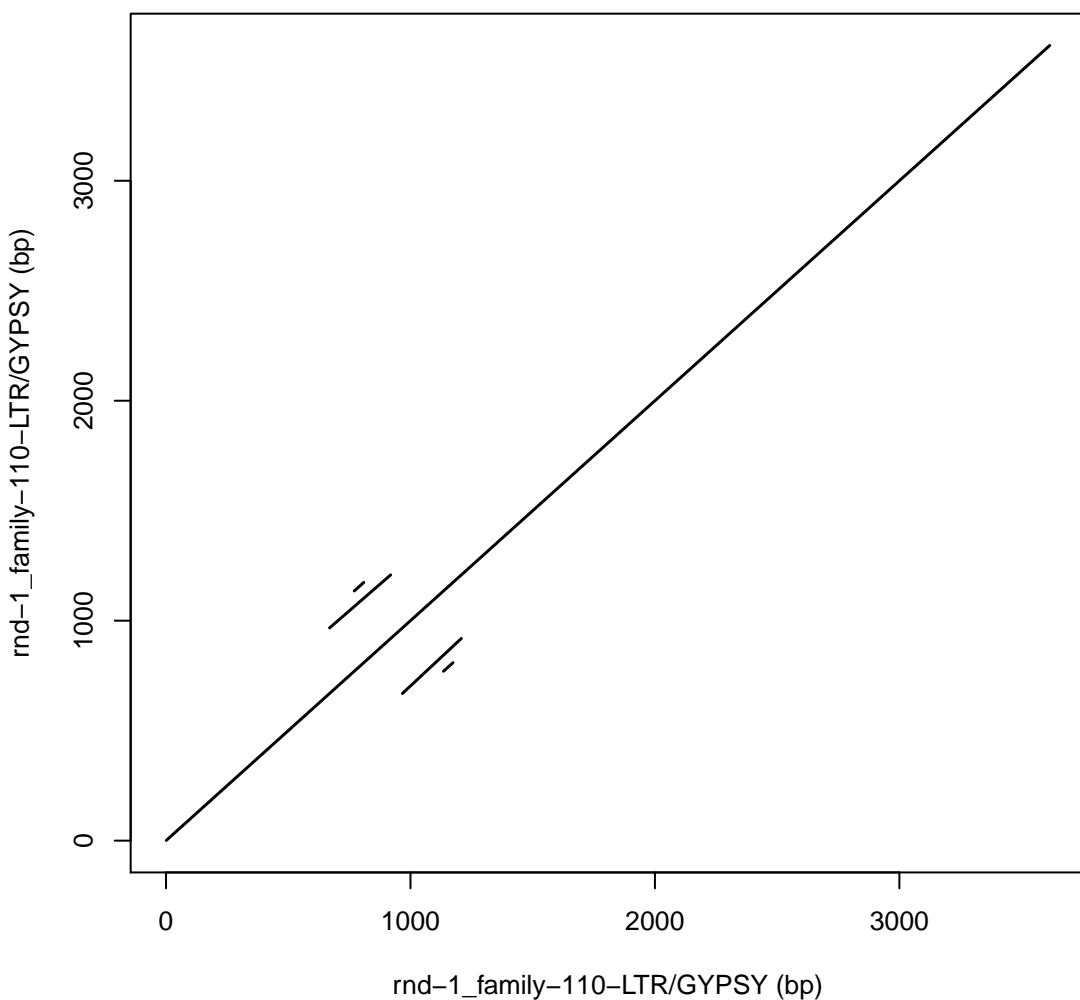
TE consensus structure and protein hits

ERV1−N4_DR_gag LTR

rnd−1_family−110−LTR/GYPSY (bp)