TE: rnd−1_family−346_s_2−LTR/Gypsy
consensus size: 1355bp; fragments: 175; full length: 1 (>=1219.5bp)

divergence to consensus (%)
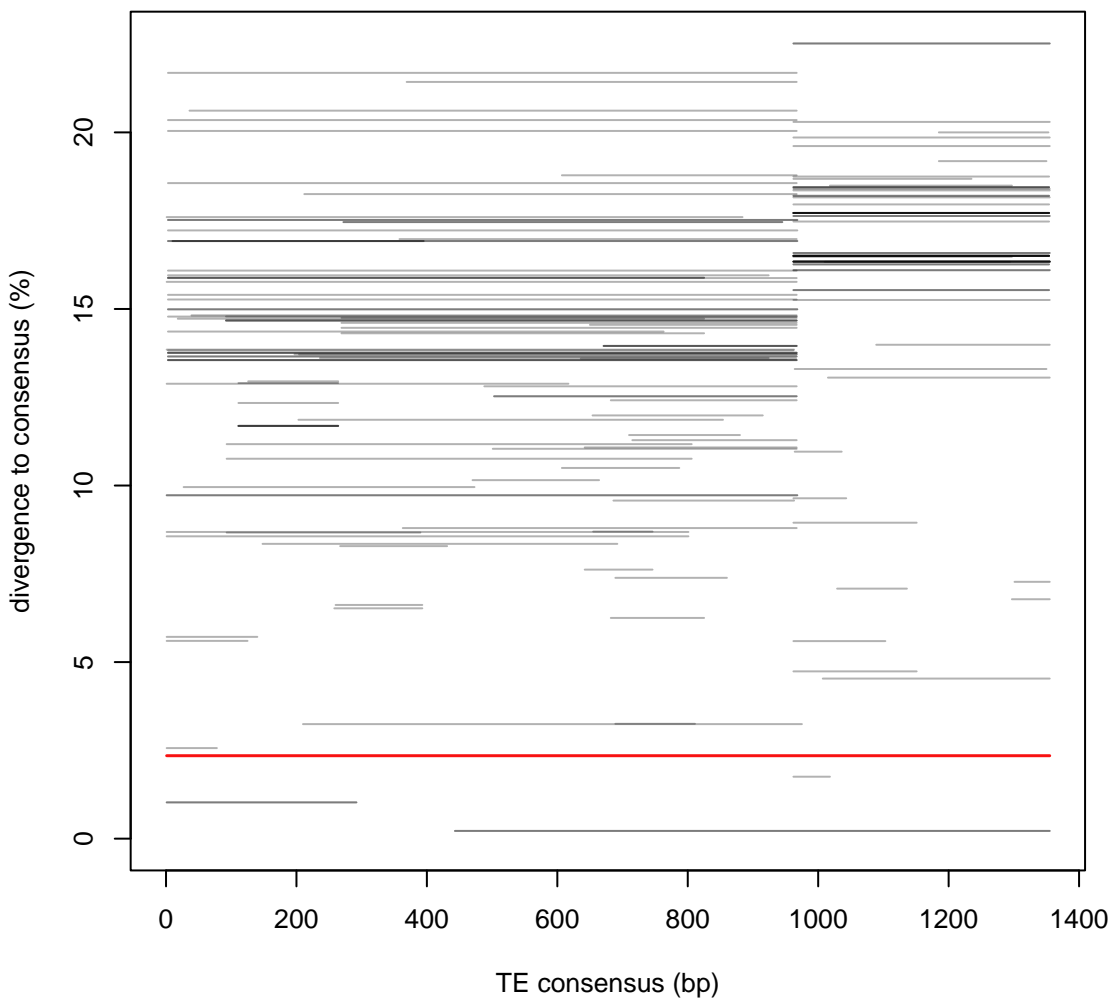
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
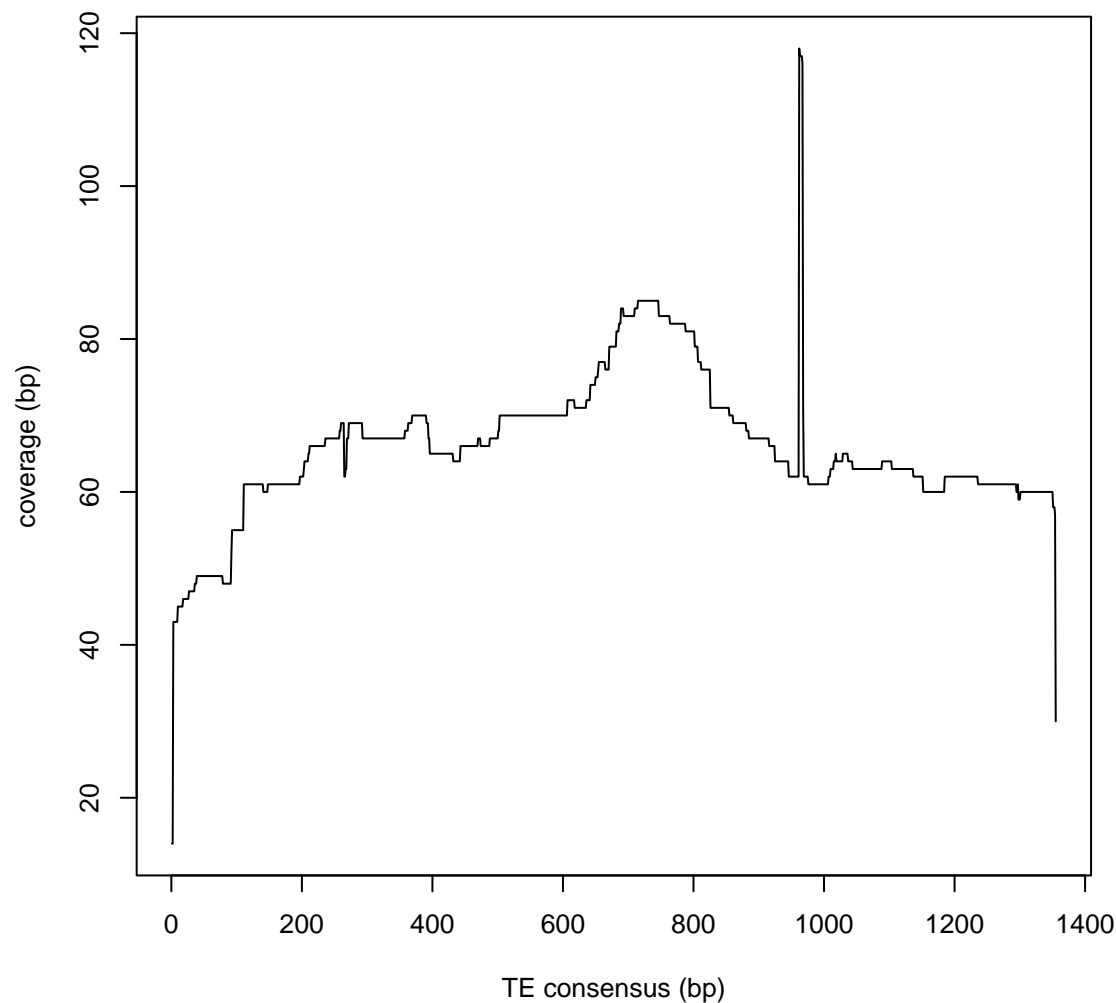
TE consensus (bp)

TE consensus self dotplot (blastn)

rnd−1_family−346_s_2−LTR/Gypsy (bp)

rnd−1_family−346_s_2−LTR/Gypsy (bp)
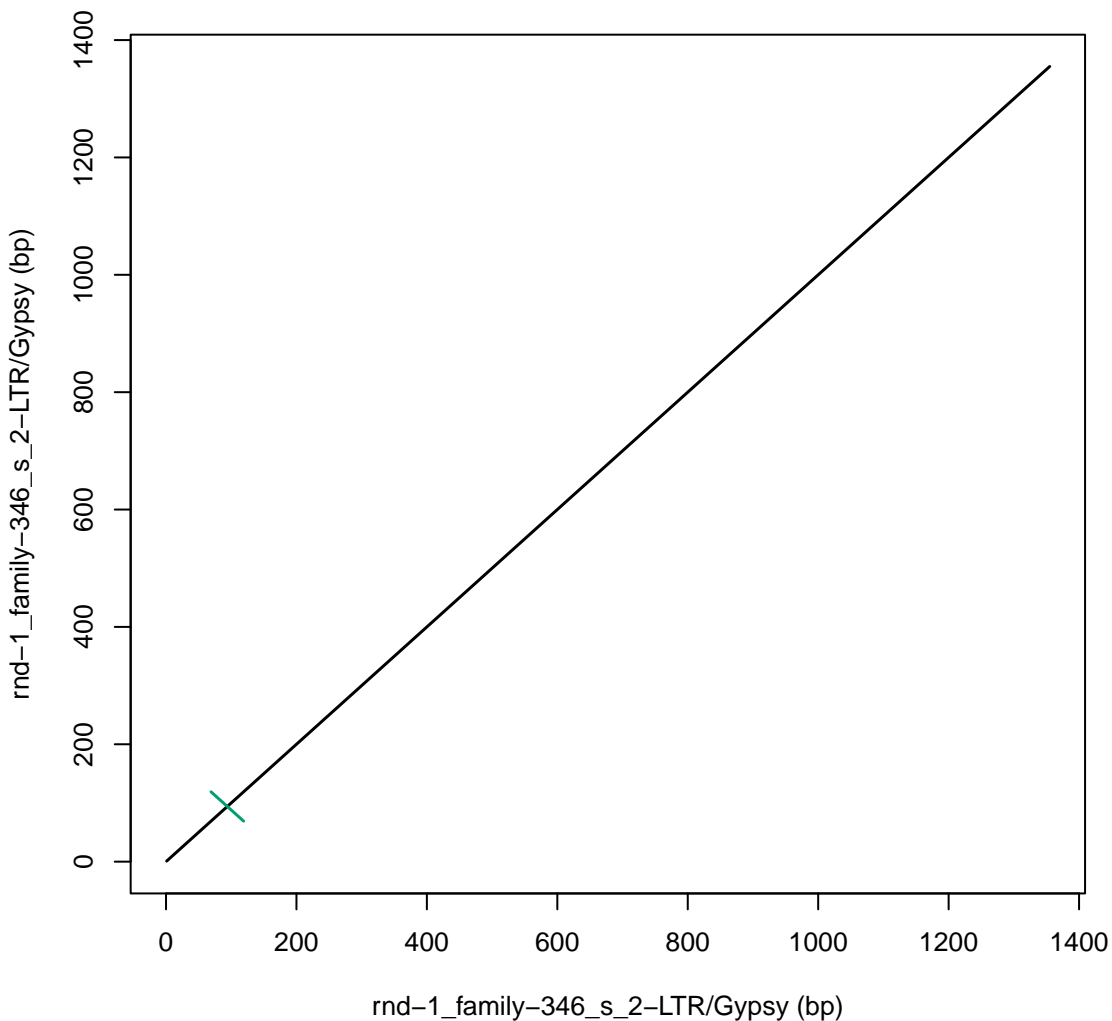
TE consensus structure and protein hits

no orf >400 bp detected

rnd−1_family−346_s_2−LTR/Gypsy (bp)