TE: ltr−1_family−53_s_1−LTR/Gypsy
consensus size: 3084bp; fragments: 590; full length: 4 (>=2775.6bp)

divergence to consensus (%)
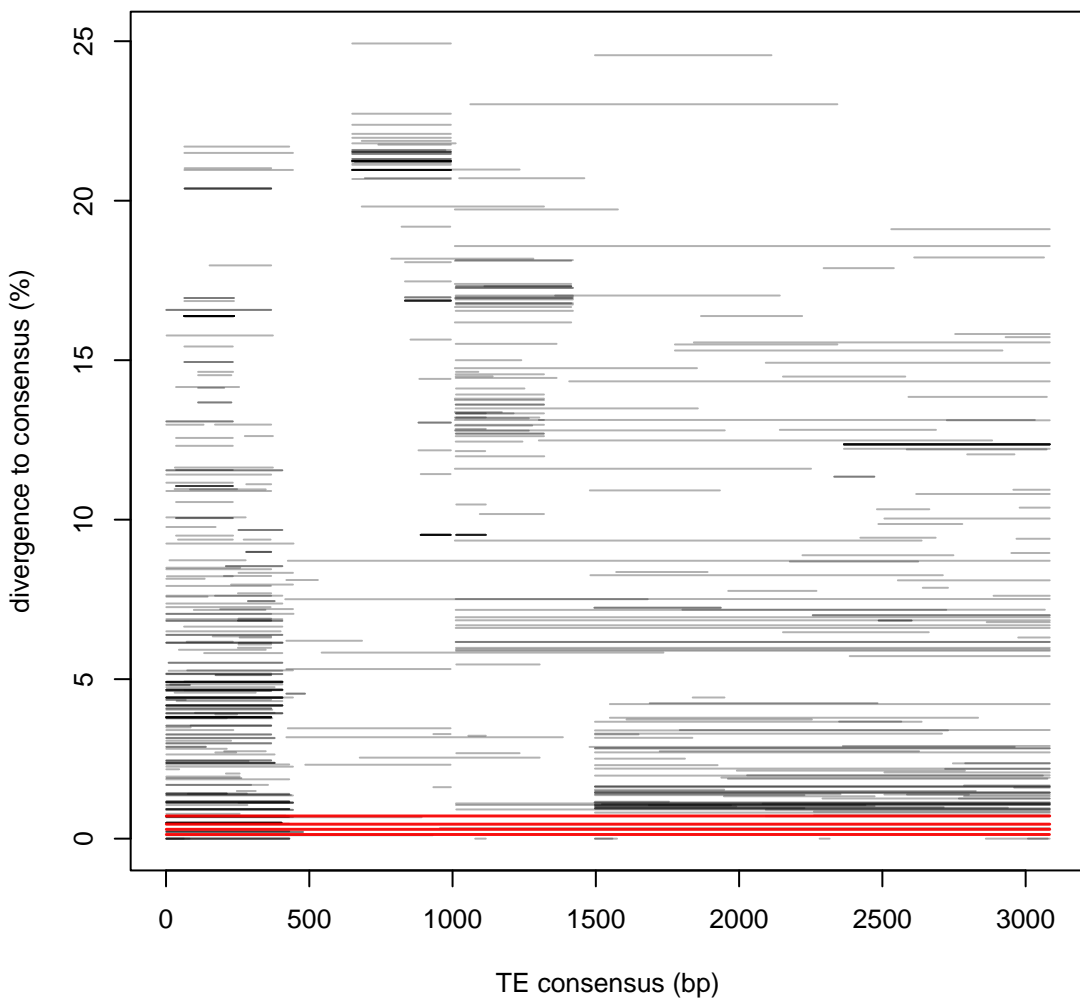
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
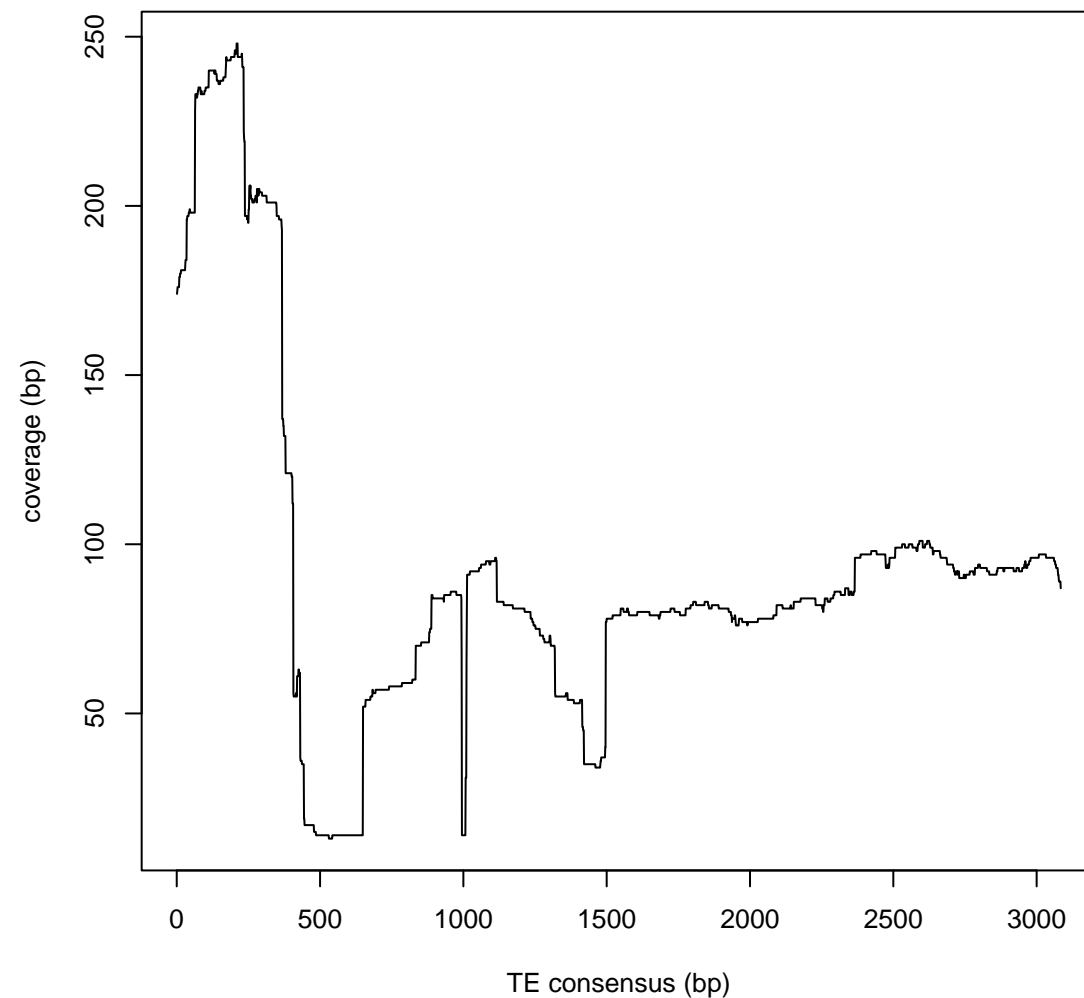
TE consensus (bp)

TE consensus self dotplot (blastn)

ltr−1_family−53_s_1−LTR/Gypsy (bp)

ltr−1_family−53_s_1−LTR/Gypsy (bp)
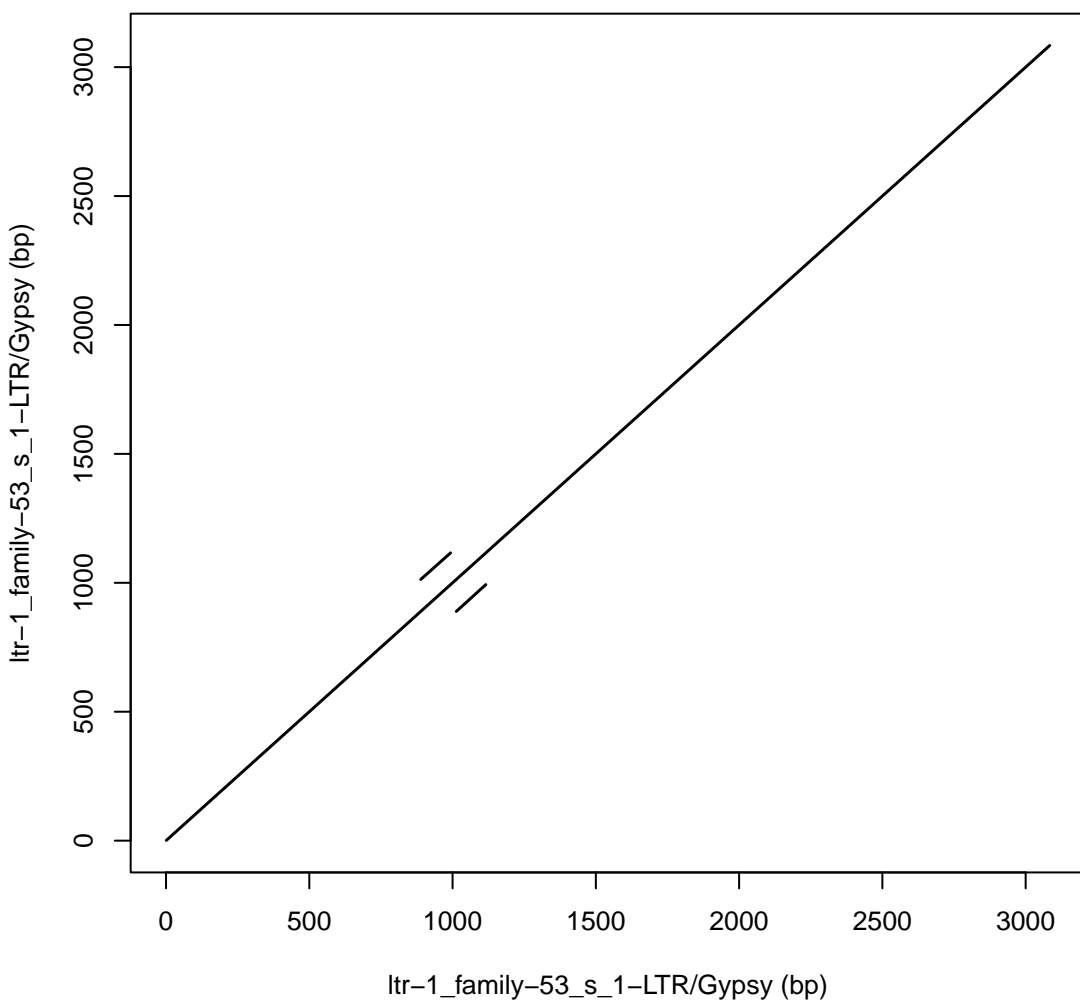
TE consensus structure and protein hits

no orf >400 bp detected

ltr−1_family−53_s_1−LTR/Gypsy (bp)