TE: rnd−5_family−54−LINE/R1
consensus size: 6696bp; fragments: 393; full length: 0 (>=6026.4bp)

divergence to consensus (%)
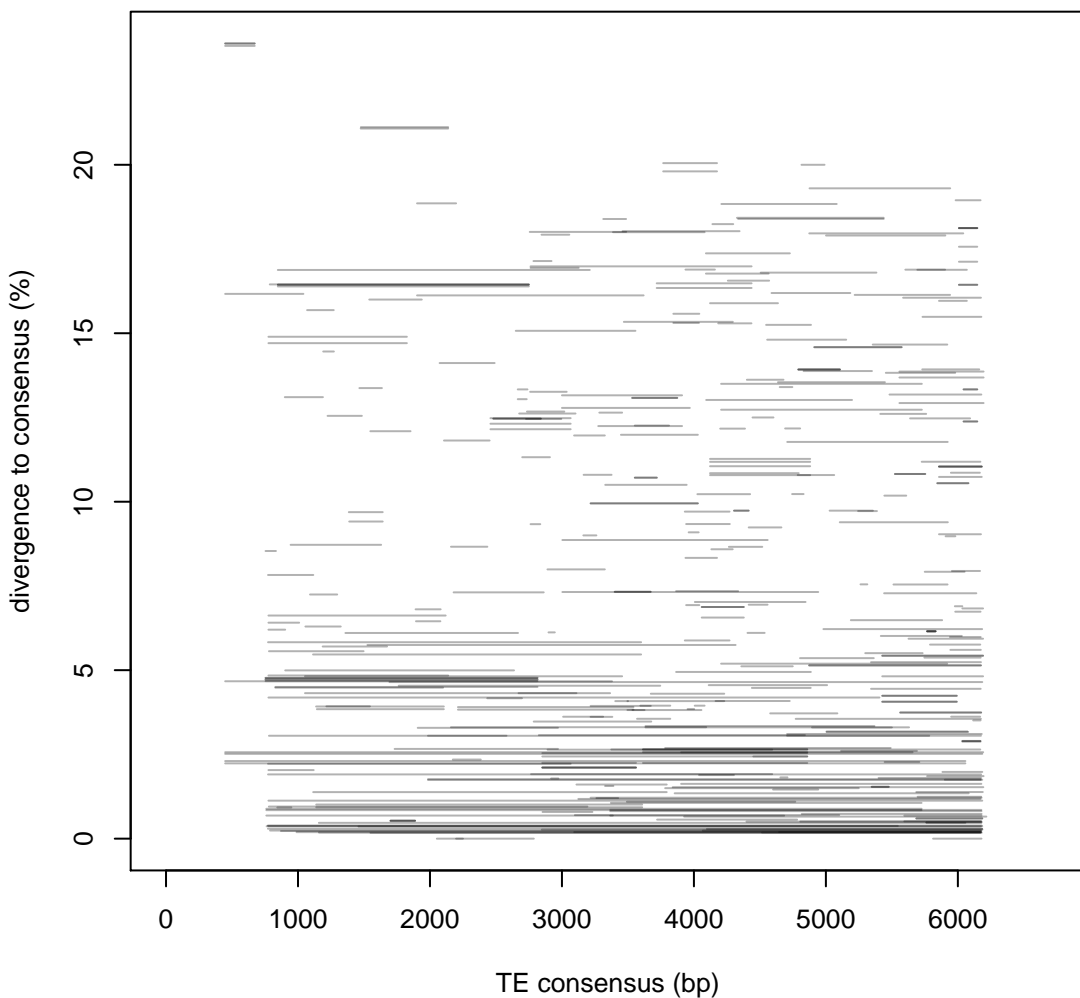
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
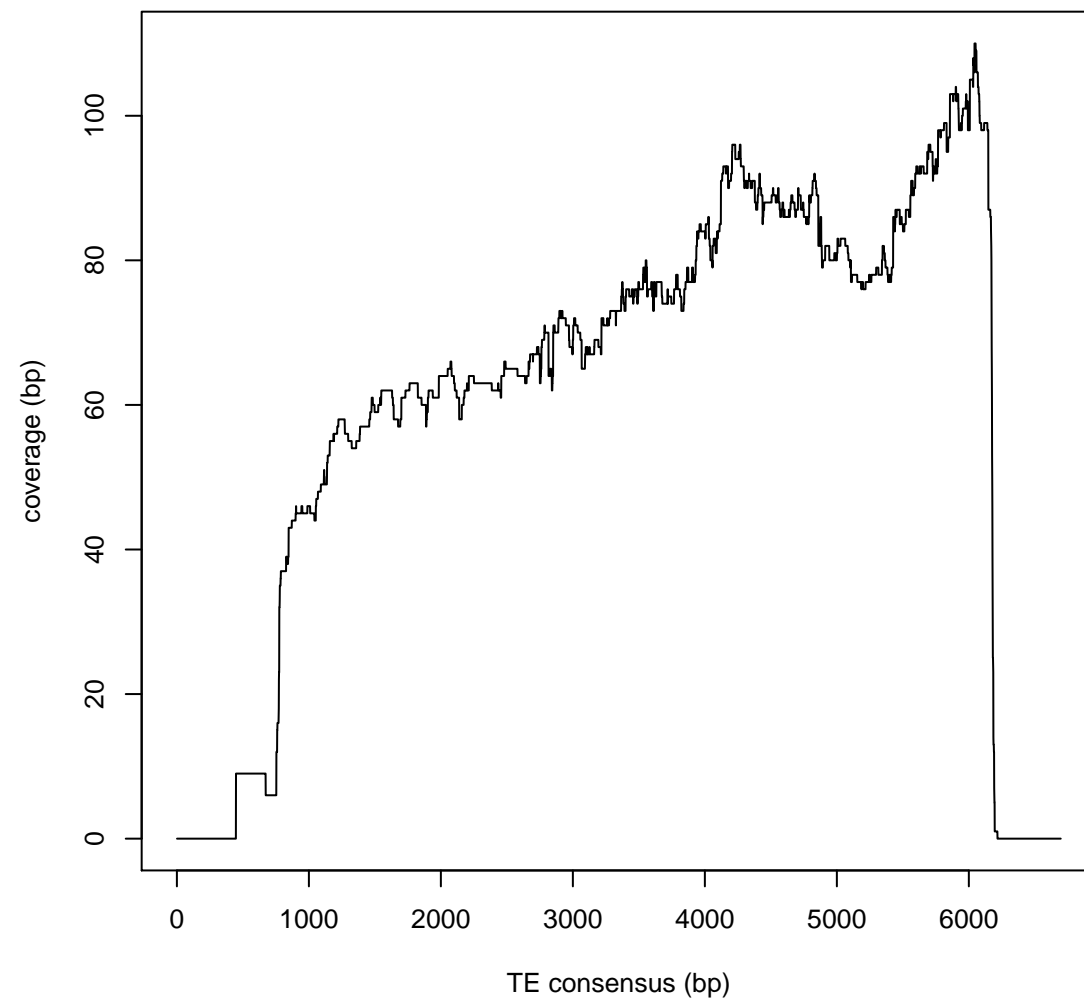
TE consensus (bp)

TE consensus self dotplot (blastn)

rnd−5_family−54−LINE/R1 (bp)

rnd−5_family−54−LINE/R1 (bp)
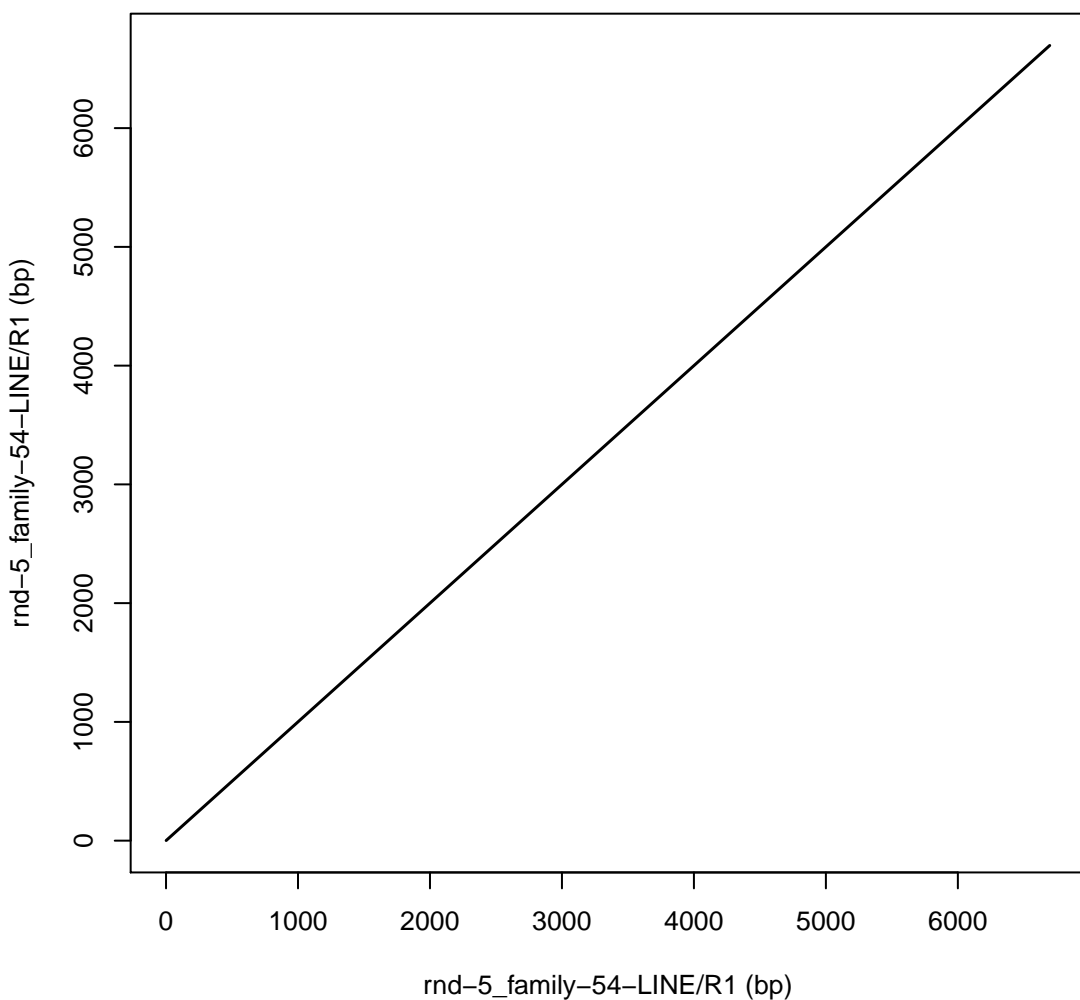
TE consensus structure and protein hits

no orf >400 bp detected

rnd−5_family−54−LINE/R1 (bp)