TE: rnd-5_family-295-LINE/I
consensus size: 5270bp; fragments: 200; full length: 10 (>=4743bp)

divergence to consensus (%)
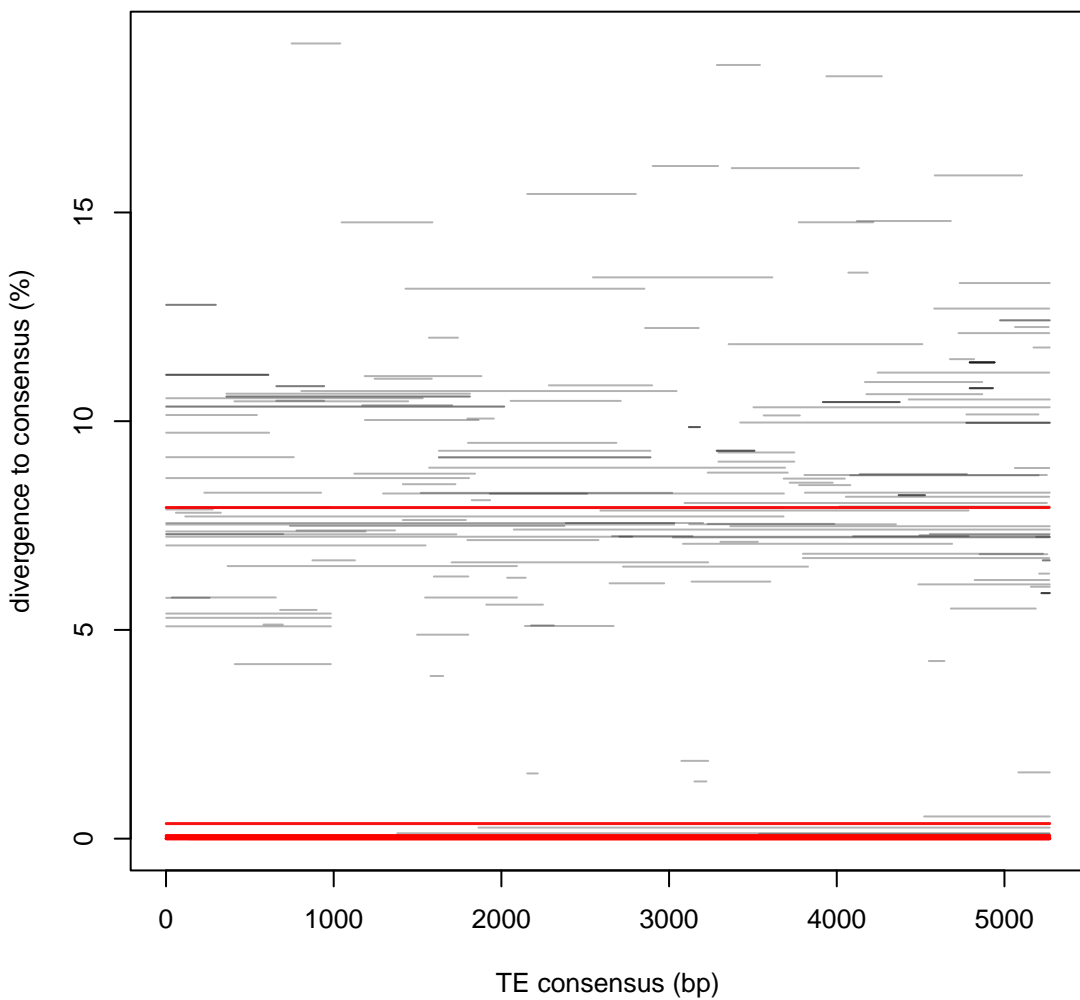
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
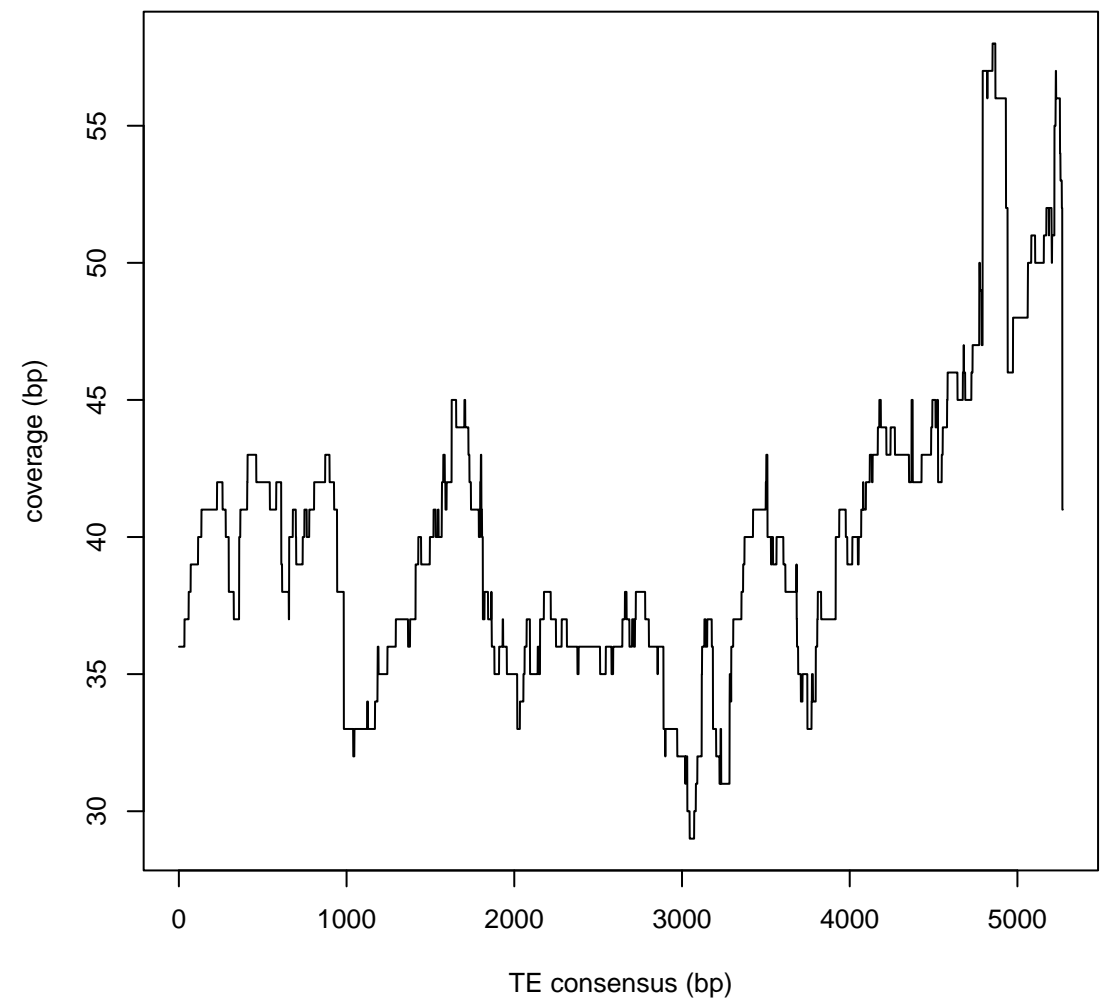
TE consensus (bp)

TE consensus self dotplot (blastn)

rnd-5_family-295-LINE/I (bp)

rnd-5_family-295-LINE/I (bp)
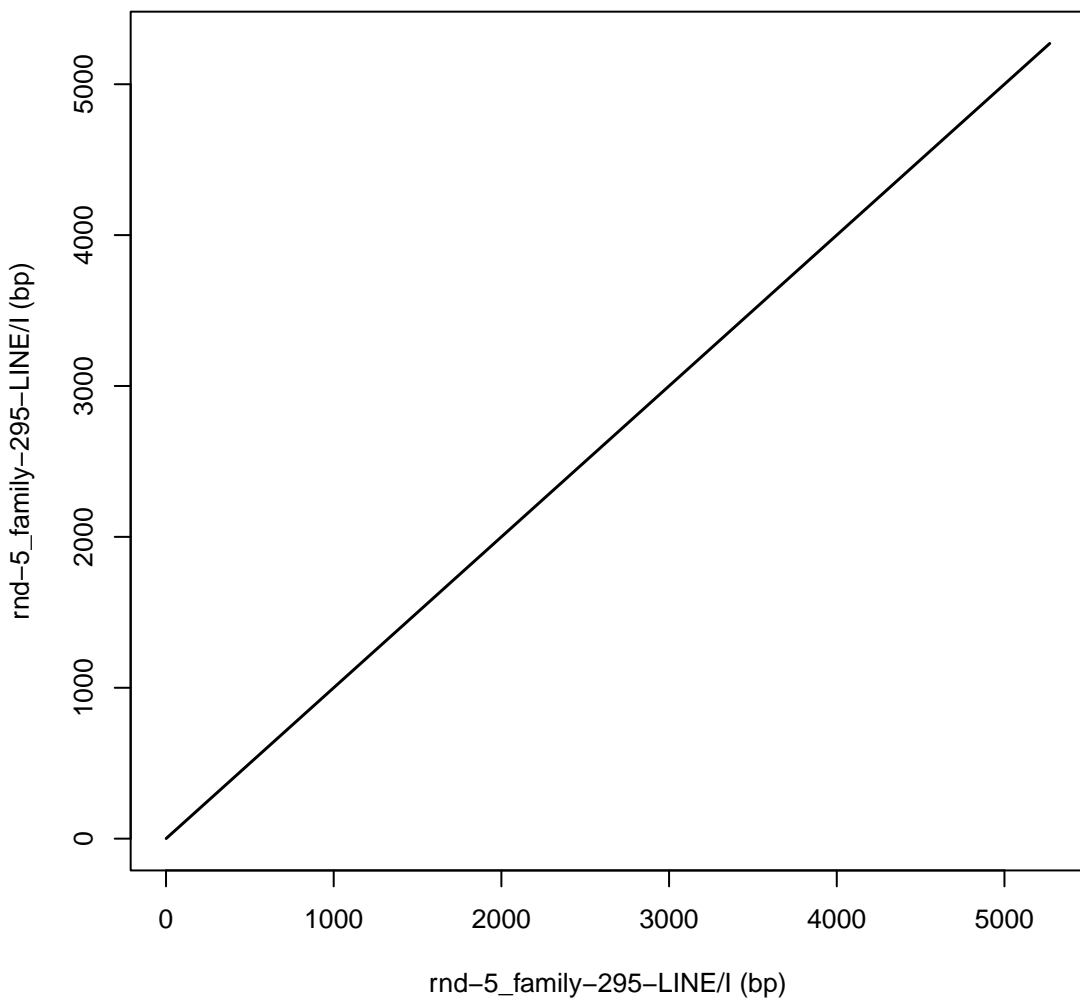
TE consensus structure and protein hits

no orf >400 bp detected

rnd-5_family-295-LINE/I (bp)