TE: rnd−5_family−472−Unclassified
consensus size: 890bp; fragments: 16; full length: 1 (>=801bp)

divergence to consensus (%)
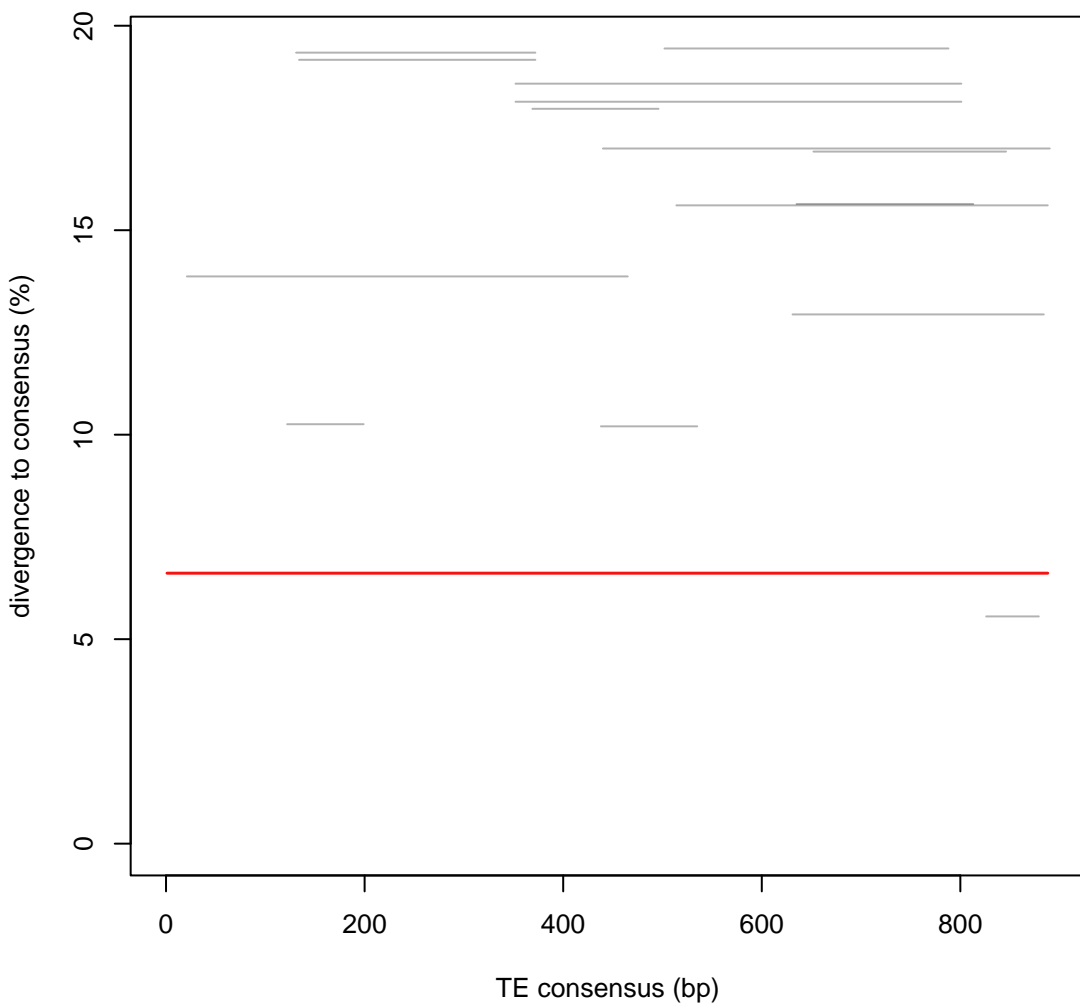
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
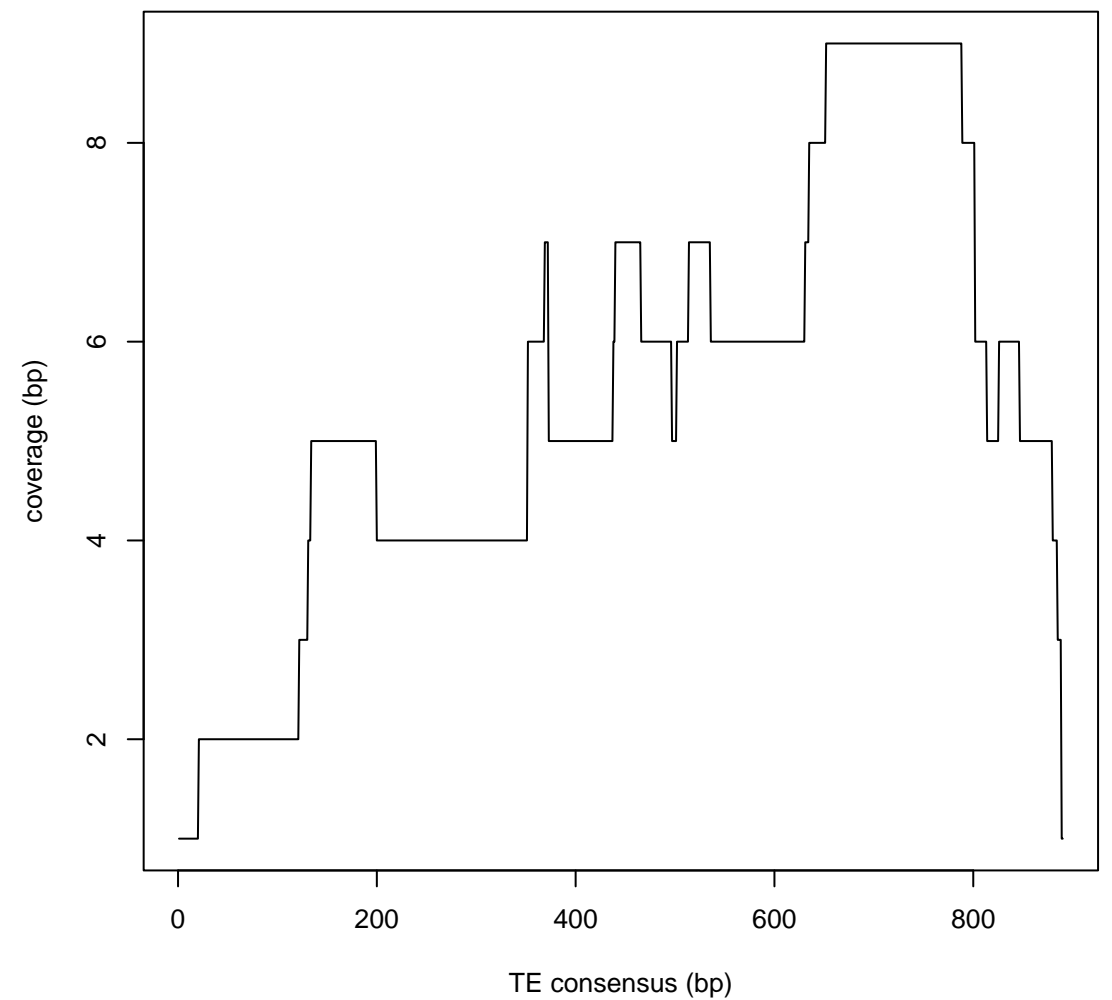
TE consensus (bp)

TE consensus self dotplot (blastn)

rnd−5_family−472−Unclassified (bp)

rnd−5_family−472−Unclassified (bp)
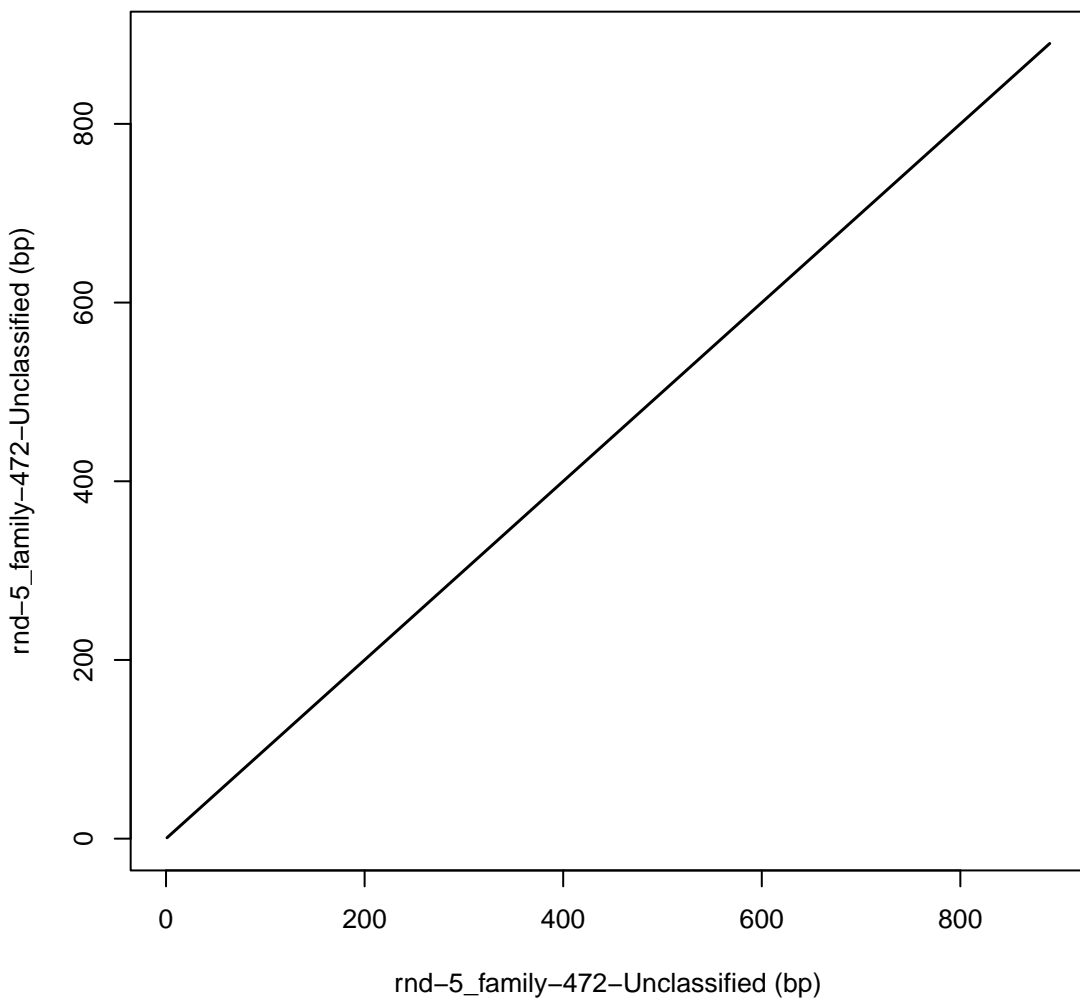
TE consensus structure and protein hits

ATCopia18A LTR/Copia

rnd−5_family−472−Unclassified (bp)