

Nucleotide

The Nucleotide database will include EST and GSS sequences in early 2019. [Read more.](#)

GenBank

Nicotiana trigonophylla psbA-trnH intergenic spacer region, partial sequence; chloroplast

GenBank: MK075954.1

[FASTA](#) [Graphics](#)
[Go to:](#)

LOCUS MK075954 489 bp DNA linear PLN 28-APR-2019
 DEFINITION Nicotiana trigonophylla psbA-trnH intergenic spacer region, partial sequence; chloroplast.
 ACCESSION MK075954
 VERSION MK075954.1
 KEYWORDS .
 SOURCE chloroplast Nicotiana trigonophylla
 ORGANISM [Nicotiana trigonophylla](#)
 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; Gunneridae; Pentapetalae; asterids; lamiids; Solanales; Solanaceae; Nicotianoideae; Nicotianeae; Nicotiana.
 REFERENCE 1 (bases 1 to 489)
 AUTHORS Prabhakar Rao,K., Sarala,K., Saroja,T. and Damodar Reddy,D.
 TITLE Nicotiana trigonophylla - trnH-psbA -Intergenic Spacer Region
 JOURNAL Unpublished
 REFERENCE 2 (bases 1 to 489)
 AUTHORS Prabhakar Rao,K., Sarala,K., Saroja,T. and Damodar Reddy,D.
 TITLE Direct Submission
 JOURNAL Submitted (20-OCT-2018) Biotechnology, division of crop improvement, ICAR-Central Tobacco Research Institute, Bhaskara nagar, Rajahmundry, Andhra Pradesh 533105, India
 COMMENT ##Assembly-Data-START##
 Sequencing Technology :: Sanger dideoxy sequencing
 ##Assembly-Data-END##
 FEATURES Location/Qualifiers
 source 1..489
 /organism="Nicotiana trigonophylla"
 /organelle="plastid:chloroplast"
 /mol_type="genomic DNA"
 /db_xref="taxon:[118711](#)"
 /tissue_type="leaf"
[misc feature](#) <1..>489
 /note="psbA-trnH intergenic spacer region"
 ORIGIN
 1 tactgccttg atccacttgg ctacatccgc cccctcgctt acttaaattc cgtttttaca
 61 ttattttaat tagaaaacaa aagattcaag ttcgaatatt tctcttcttt cttatttcaa
 121 tgatattatt atttcaaaga taagaatag aagtaaaatc tcgatttttt ttttgaata
 181 aaaatcaaaa agatatagta acattagcaa gaagagaaac aagtcatttt tctacaattt
 241 taaataaata caaaatcaaa atagaatact caatcatgaa taaatgcaag caaataacct
 301 ctcttctttt ttctataatg taaacaaaa agtctatgta agtaaaatc tagtaataa
 361 ataaaaagaa aaaaagaag gagcaatagc accctcttga tagaacaaga aaatgattat
 421 tgctcctttc ttttcaaac ctctataga ctgactggg atcttatcca tttgatagatg
 481 gagcttcga
 //