

Nucleotide

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

[GenBank](#)

## Nicotiana glutinosa tRNA-His (trnH-GUG) gene and psbA-trnH intergenic spacer, partial sequence; chloroplast

GenBank: MK072596.1

[FASTA](#) [Graphics](#)

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LOCUS MK072596 474 bp DNA linear PLN 06-FEB-2019

DEFINITION Nicotiana glutinosa tRNA-His (trnH-GUG) gene and psbA-trnH intergenic spacer, partial sequence; chloroplast.

ACCESSION MK072596

VERSION MK072596.1

KEYWORDS .

SOURCE chloroplast Nicotiana glutinosa

ORGANISM [Nicotiana glutinosa](#)

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; Gunneridae; Pentapetalae; asterids; lamiids; Solanales; Solanaceae; Nicotianoideae; Nicotianeae; Nicotiana.

REFERENCE 1 (bases 1 to 474)

AUTHORS Prabhakar Rao,K., Sarala,K., Saroja,T. and Damodar Reddy,D.

TITLE Direct Submission

JOURNAL Submitted (18-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..474

/organism="Nicotiana glutinosa"

/organelle="plastid:chloroplast"

/mol\_type="genomic DNA"

/db\_xref="taxon:35889"

/tissue\_type="leaf"

[gene](#)

complement(<1..30)

/gene="trnH-GUG"

[tRNA](#)

complement(<1..30)

/gene="trnH-GUG"

/product="tRNA-His"

[misc feature](#)

complement(31..>474)

/note="psbA-trnH intergenic spacer"

ORIGIN

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121 ctatttcaat gatattatta ttgcaaagat aagaatatga agtcaaaatt cgattttttt
181 ttgaaataaa aatcaaaaag atatagtaac attagcaaga agagaacaa gtcatttttc
241 tccaatttta aataaatata aatcaaaat agaatactca atcatgaata aatgcaagca
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361 gtaataaaat accagtaaat aaataaataa aaagaaaaaa agaaaggagc aatagcacc
421 tcttgataga acaagaaaat gattattgct ctttcttttt caaaacctcc tata

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