

Appendix I
NUCLEIC ACID SEQUENCES

Nucleotide

GenBank

Colletotrichum sp. isolate HCS3 internal transcribed spacer 1, partial sequence; 5.8S ribosomal RNA gene and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence

GenBank: MN128230.1

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

LOCUS MN128230 559 bp DNA linear PLN 09-JUL-2019

DEFINITION Colletotrichum sp. isolate HCS3 internal transcribed spacer 1, partial sequence; 5.8S ribosomal RNA gene and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence.

ACCESSION MN128230

VERSION MN128230.1

KEYWORDS .

SOURCE Colletotrichum sp.

ORGANISM [Colletotrichum sp.](#)
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina;
Sordariomycetes; Hypocreomycetidae; Glomerellales; Glomerellaceae;
Colletotrichum.

REFERENCE 1 (bases 1 to 559)

AUTHORS Talukdar,R., Padhi,S. and Tayung,K.

TITLE Antimicrobial secondary metabolites isolated from endophytic fungi inhabiting healthy tissues of Houlttuynia cordata Thunb., an ethnomedicinal plant of North East India

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 559)

AUTHORS Talukdar,R., Padhi,S. and Tayung,K.

TITLE Direct Submission

JOURNAL Submitted (03-JUL-2019) DEPARTMENT OF BOTANY, GAUHATI UNIVERSITY, GOPINATH BORDOLOI NAGAR, GAUHATI, ASSAM 781014, India

COMMENT ##Assembly-Data-START##
Sequencing Technology :: Sanger dideoxy sequencing
##Assembly-Data-END##

FEATURES Location/Qualifiers

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[misc_RNA](#)
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/note="contains internal transcribed spacer 1, 5.8S ribosomal RNA, internal transcribed spacer 2, and large subunit ribosomal RNA"

ORIGIN

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121 tggcacaagc aaataatcaa aacttttaac aacggatctc ttggttctgg catcgatgaa
181 gaacgcagcg aaatgcgata agtaatgtga attgcagaat tcagtgaatc atcgaatctt
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421 gactcttgcc gtaaaacccc ccaattttta caaggttgac ctcgcatgag gtaggaatac
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541 actgttgctt ctctagtga

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Nucleotide

GenBank

Phyllosticta sp. isolate HCS6 internal transcribed spacer 1, partial sequence; 5.8S ribosomal RNA gene and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence

GenBank: MN170572.1

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

LOCUS MN170572 607 bp DNA linear PLN 17-JUL-2019

DEFINITION Phyllosticta sp. isolate HCS6 internal transcribed spacer 1, partial sequence; 5.8S ribosomal RNA gene and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence.

ACCESSION MN170572

VERSION MN170572.1

KEYWORDS .

SOURCE Phyllosticta sp.

ORGANISM [Phyllosticta sp.](#)
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina; Dothideomycetes; Dothideomycetes incertae sedis; Botryosphaerales; Phyllostictaceae; Phyllosticta; unclassified Phyllosticta.

REFERENCE 1 (bases 1 to 607)

AUTHORS Talukdar,R., Padhi,S. and Tayung,K.

TITLE Antimicrobial secondary metabolites isolated from endophytic fungi inhabiting healthy tissues of Houlttuynia cordata Thunb., an ethnomedicinal plant of North East India

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 607)

AUTHORS Talukdar,R., Padhi,S. and Tayung,K.

TITLE Direct Submission

JOURNAL Submitted (12-JUL-2019) DEPARTMENT OF BOTANY, GAUHATI UNIVERSITY, GOPINATH BORDOLOI NAGAR, GAUHATI, ASSAM 781014, India

COMMENT ##Assembly-Data-START##
Sequencing Technology :: Sanger dideoxy sequencing
##Assembly-Data-END##

FEATURES

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/country="India"
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[misc_RNA](#)
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/note="contains internal transcribed spacer 1, 5.8S ribosomal RNA, internal transcribed spacer 2, and large subunit ribosomal RNA"

ORIGIN

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121 tggctaagt cccgccagta tacaactc aagcgattat tttgtgaagt cctgatatat
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241 cagcgaatg cgataagtaa tgtgaattgc agaattcagt gaatcatcga atctttgaac
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