Phytophthora infestans strain IMI66006 internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, complete sequence

GenBank: AF266779.1

FASTA Graphics

Go to:

LOCUS AF266779 795 bp DNA linear PLN 29-JAN-2024

DEFINITION Phytophthora infestans strain IMI66006 internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, complete sequence.

ACCESSION AF266779

VERSION AF266779.1

KEYWORDS .

SOURCE Phytophthora infestans (potato late blight agent)

ORGANISM Phytophthora infestans

Eukaryota; Sar; Stramenopiles; Oomycota; Peronosporales; Peronosporaceae; Phytophthora.

REFERENCE 1 (bases 1 to 795)

AUTHORS Cooke,D.E., Drenth,A., Duncan,J.M., Wagels,G. and Brasier,C.M.

TITLE A molecular phylogeny of Phytophthora and related oomycetes

JOURNAL Fungal Genet. Biol. 30 (1), 17-32 (2000)

PUBMED 10955905

REFERENCE 2 (bases 1 to 795)

AUTHORS Drenth,A., Cooke,D.E.L., Duncan,J.M., Wagels,G. and Brasier,C.M.

TITLE Direct Submission

JOURNAL Submitted (24-MAY-2000) CRC for Tropical Plant Protection, University of Queensland, John Hines Building, Brisbane, QLD 4072, Australia

FEATURES Location/Qualifiers

source 1..795

/organism="Phytophthora infestans"

/mol_type="genomic DNA"

/strain="IMI66006"

/db_xref="taxon:4787"

/note="cause of late blight"

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/product="internal transcribed spacer 1"

rRNA 219..380

/product="5.8S ribosomal RNA"

misc_RNA 381..795

/product="internal transcribed spacer 2"

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