

Diaporthe phaseolorum isolate TW42 small subunit ribosomal RNA gene, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence

GenBank: MH930433.1

[FASTA Graphics](#)

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LOCUS MH930433 601 bp DNA linear PLN 21-AUG-2019

DEFINITION Diaporthe phaseolorum isolate TW42 small subunit ribosomal RNA gene, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence.

ACCESSION MH930433

VERSION MH930433.1

KEYWORDS .

SOURCE Diaporthe phaseolorum

ORGANISM [Diaporthe phaseolorum](#)
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina;
Sordariomycetes; Sordariomycetidae; Diaporthales; Diaporthaceae;
Diaporthe.

REFERENCE 1 (bases 1 to 601)

AUTHORS Leung,F.-M., Feng,J., Wong,H.-N. and Cheung,H.-Y.

TITLE Application of Various DNA Sequences for Establishing Phylogenetic Relationship of Endophytic Fungi Isolated from *Tripterygium wilfordii* Hook f.

JOURNAL Hong Kong Pharm J 25 (4), 127-136 (2018)

REFERENCE 2 (bases 1 to 601)

AUTHORS Cheung,H.Y.

TITLE Direct Submission

JOURNAL Submitted (20-SEP-2018) Department of Biomedical Sciences, City University of Hong Kong, 83 Tat Chee Avenue, Kowloon Tong, Hong Kong SAR 999077, China

COMMENT ##Assembly-Data-START##
Assembly Method :: BioEdit v. 7.0.5
Sequencing Technology :: Illumina
##Assembly-Data-END##

FEATURES Location/Qualifiers

source 1..601
/organism="Diaporthe phaseolorum"
/mol_type="genomic DNA"
/isolate="TW42"
/db_xref="taxon:[36923](#)"

[misc_RNA](#) <1..>601
/note="contains small subunit ribosomal RNA, internal transcribed spacer 1, 5.8S ribosomal RNA, internal

transcribed spacer 2, and large subunit ribosomal RNA"

ORIGIN

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541 aatttgacct cggatcaggt aggaataccc gctgaactta agcatatcaa taagcggagg
601 a
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