

Nucleotide

Nucleotide

Advanced

GenBank

Send to:

Ascomycota sp. strain OKr4 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: MF510819.1

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

LOCUS MF510819 548 bp DNA linear PLN 25-JUL-2017

DEFINITION Ascomycota sp. strain OKr4 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 MF510819

VERSION MF510819.1

KEYWORDS .

SOURCE Ascomycota sp.
ORGANISM [Ascomycota sp.](#)
Eukaryota; Fungi; Dikarya; Ascomycota.

REFERENCE 1 (bases 1 to 548)
AUTHORS Abdalmoohsin,R.G., Lahuf,A.A., Alhamiri,Y.N. and Alhblany,A.Z.
TITLE Molecular characterization of the pathogenic fungi causing the root rot disease on okra plants in Kerbala governorate/Iraq
JOURNAL Unpublished

REFERENCE 2 (bases 1 to 548)
AUTHORS Abdalmoohsin,R.G., Lahuf,A.A., Alhamiri,Y.N. and Alhblany,A.Z.
TITLE Direct Submission
JOURNAL Submitted (20-JUL-2017) Plant protection Department, Agriculture College-University of Kerbala, City center, Kerbala, Kerbala KK1 3DR, Iraq

COMMENT ##Assembly-Data-START##
Sequencing Technology :: ABI 3730xI automated Sequencer
##Assembly-Data-END##

FEATURES Location/Qualifiers