

Nucleotide

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Fusarium solani strain OKr3 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: MF510823.1

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

LOCUS MF510823 543 bp DNA linear PLN 25-JUL-2017

DEFINITION *Fusarium solani* strain OKr3 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 MF510823

VERSION MF510823.1

KEYWORDS .

SOURCE *Fusarium solani*

ORGANISM [Fusarium solani](#)
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina; Sordariomycetes; Hypocreomycetidae; Hypocreales; Nectriaceae; *Fusarium*; *Fusarium solani* species complex.

REFERENCE 1 (bases 1 to 543)

AUTHORS Abdalmoohsin,R.G., Lahuf,A.A., Alhamiri,Y.N. and Ablany,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JOURNAL Unpublished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 543)

AUTHORS Abdalmoohsin,R.G., Lahuf,A.A., Alhamiri,Y.N. and Ablany,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##