

Nucleotide

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Fusarium keratoplasticum strain OKr2 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: MF510816.1

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LOCUS MF510816 530 bp DNA linear PLN 25-JUL-2017
DEFINITION *Fusarium keratoplasticum* strain OKr2 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 MF510816
VERSION MF510816.1
KEYWORDS .
SOURCE *Fusarium keratoplasticum*
ORGANISM [Fusarium keratoplasticum](#)
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina; Sordariomycetes; Hypocreomycetidae; Hypocreales; Nectriaceae; *Fusarium*; *Fusarium solani* species complex.
REFERENCE 1 (bases 1 to 530)
AUTHORS Lahuf,A.A., Abdalmoohsin,R.G., 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JOURNAL Unpublished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 530)
AUTHORS Lahuf,A.A., Abdalmoohsin,R.G., 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