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## Fusarium fujikuroi strain OKr1 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: MF510817.1

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LOCUS MF510817 534 bp DNA linear PLN 25-JUL-2017

DEFINITION *Fusarium fujikuroi* strain OKr1 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 MF510817

VERSION MF510817.1

KEYWORDS .

SOURCE *Fusarium fujikuroi* (*Gibberella fujikuroi*)

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

REFERENCE 1 (bases 1 to 534)

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JOURNAL Unpublished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 534)

AUTHORS Lahuf,A.A., Abdalmoohsin,R.G., Alhamiri,Y.N. and Alhblany,A.Z.

TITLE Direct Submission

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