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Fusarium fujikuroi strain Lahuf10 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: MF510829.1

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LOCUS MF510829 534 bp DNA linear PLN 26-JUL-2017
 DEFINITION *Fusarium fujikuroi* strain Lahuf10 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 MF510829
 VERSION MF510829.1
 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., Jaafar,O.H. and Li,J.
 TITLE First report of *Gibberella fujikuroi* on *Mangifera indica* in Iraq
 JOURNAL Unpublished
 REFERENCE 2 (bases 1 to 534)
 AUTHORS Lahuf,A.A., Jaafar,O.H. and Li,J.
 TITLE Direct Submission
 JOURNAL Submitted (21-JUL-2017) Plant protection Department, Agriculture College-University of Kerbala, City center, Kerbala, Kerbala KK1 3DR, Iraq
 COMMENT ##Assembly-Data-START##