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Aspergillus nomius strain OKr7 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: MF510822.1

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LOCUS MF510822 924 bp DNA linear PLN 25-JUL-2017
DEFINITION *Aspergillus nomius* strain OKr7 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 MF510822
VERSION MF510822.1
KEYWORDS .
SOURCE *Aspergillus nomius*
ORGANISM [Aspergillus nomius](#)
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina; Eurotiomycetes; Eurotiomycetidae; Eurotiales; Aspergillaceae; *Aspergillus*.
REFERENCE 1 (bases 1 to 924)
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JOURNAL Unpublished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 924)
AUTHORS Lahuf,A.A., Abdalmoohsin,R.G., Alhamiri,Y.N. and Alhblany,A.Z.
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