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Cladosporium oxysporum strain SL.Co1 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: MF511908.1

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LOCUS MF511908 576 bp DNA linear PLN 26-JUL-2017
DEFINITION Cladosporium oxysporum strain SL.Co1 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 MF511908
VERSION MF511908.1
KEYWORDS .
SOURCE Cladosporium oxysporum
ORGANISM [Cladosporium oxysporum](#)
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina; Dothideomycetes; Dothideomycetidae; Capnodiales; Cladosporiaceae; Cladosporium.
REFERENCE 1 (bases 1 to 576)
AUTHORS Jaafar,O.H. and Lahuf,A.A.
TITLE Molecular characterization of the pathogenic fungus causing the root rot disease on Tomato plants in Kerbala governorate/Iraq
JOURNAL Unpublished
REFERENCE 2 (bases 1 to 576)
AUTHORS Jaafar,O.H. and Lahuf,A.A.
TITLE Direct Submission
JOURNAL Submitted (21-JUL-2017) Plant protection Department, Agriculture College-University of Kerbala, City center, Kerbala, Kerbala KK1 3DR, Iraq