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## Curvularia tsudae strain OKr8 5.8S ribosomal RNA gene, partial sequence; internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence

GenBank: MF510824.1

[FASTA](#) [Graphics](#)

LOCUS MF510824 312 bp DNA linear PLN 25-JUL-2017

DEFINITION Curvularia tsudae strain OKr8 5.8S ribosomal RNA gene, partial sequence; internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence.

ACCESSION MF510824

VERSION MF510824.1

KEYWORDS .

SOURCE Curvularia tsudae (Cochliobolus australiensis)

ORGANISM [Curvularia tsudae](#)

Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina; Dothideomycetes; Pleosporomycetidae; Pleosporales; Pleosporineae; Pleosporaceae; Curvularia.

REFERENCE 1 (bases 1 to 312)

AUTHORS Abdalmoohsin,R.G., Lahuf,A.A., 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JOURNAL Unpublished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 312)

AUTHORS Abdalmoohsin,R.G., Lahuf,A.A., 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

COMMENT ##Assembly-Data-START##

Sequencing Technology :: ABI 3730xI automated Sequencer

##Assembly-Data-END##

FEATURES Location/Qualifiers