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Byssochlamys spectabilis strain Pa.Bs1 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: MF511048.1

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

LOCUS MF511048 1107 bp DNA linear PLN 26-JUL-2017

DEFINITION *Byssochlamys spectabilis* strain Pa.Bs1 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 MF511048

VERSION MF511048.1

KEYWORDS .

SOURCE *Byssochlamys spectabilis* (anamorph: *Paecilomyces spectabilis*)

ORGANISM [Byssochlamys spectabilis](#)
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina;
Eurotiomycetes; Eurotiomycetidae; Eurotiales; Thermoascaceae;
Byssochlamys.

REFERENCE 1 (bases 1 to 1107)

AUTHORS Altememe,Z.A. and Lahuf,A.A.

TITLE First report of *Byssochlamys spectabilis* fungus on *Prunus armeniaca* plant in Iraq

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 1107)

AUTHORS Altememe,Z.A. 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

COMMENT ##Assembly-Data-START##
Sequencing Technology :: ABI 3730xI automated Sequencer
##Assembly-Data-END##