Cephalosporium gramineum culture CBS:391.70 strain CBS 391.70 small subunit ribosomal RNA gene, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence

GenBank: MH859752.1

FASTA Graphics PopSet

Go to:

LOCUS MH859752 585 bp DNA linear PLN 29-JUN-2022
DEFINITION Cephalosporium gramineum culture CBS:391.70 strain CBS 391.70 small
subunit ribosomal RNA gene, partial sequence; internal transcribed
spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer
2, complete sequence; and large subunit ribosomal RNA gene, partial
sequence.
ACCESSION MH859752
VERSION MH859752.1
DBLINK BioProject: PRJNA422523
KEYWORDS .
SOURCE [Hymenula] cerealis
ORGANISM [Hymenula] cerealis
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina;
Pezizomycotina incertae sedis; Hymenella.
REFERENCE 1 (bases 1 to 585)
AUTHORS Vu,D., Groenewald,M., de Vries,M., Gehrmann,T., Stielow,B.,
Eberhardt,U., Al-Hatmi,A., Groenewald,J.Z., Cardinali,G.,
Houbraken,J., Boekhout,T., Crous,P.W., Robert,V. and Verkley,G.J.M.
TITLE Large-scale generation and analysis of filamentous fungal DNA
barcodes boosts coverage for kingdom fungi and reveals thresholds
for fungal species and higher taxon delimitation
JOURNAL Stud. Mycol. 92, 135-154 (2019)
PUBMED 29955203
REFERENCE 2 (bases 1 to 585)
AUTHORS Vu,D., Groenewald,M., De Vries,M., Gehrmann,T., Stielow,B.,
Eberhard,U., Al-Hatmi,A.M., Groenewald,J.Z., Cardinali,G.,
Boekhout,T., Crous,P., Robert,V. and Verkleij,G.J.
TITLE Direct Submission
JOURNAL Submitted (15-DEC-2017) Bioinformatics group, Westerdijk Fungal
Biodiversity Institute, Upsalalaan, Utrecht 3584CT, Netherlands
COMMENT This submission replaces the Targeted Locus Study (TLS) project
KBUW00000000, which included the individual sequences
KBUW01000001-KBUW01024118.

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                     transcribed spacer 1, 5.8S ribosomal RNA, internal
                     transcribed spacer 2, and large subunit ribosomal RNA"
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