

Aspergillus flavus NRRL3357 uncharacterized protein (G4B84_005804), partial mRNA

NCBI Reference Sequence: XM_041285628.1

[FASTA Graphics](#)

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LOCUS XM_041285628 1335 bp mRNA linear PLN 15-SEP-2021

DEFINITION Aspergillus flavus NRRL3357 uncharacterized protein
(G4B84_005804),

partial mRNA.

ACCESSION XM_041285628

VERSION XM_041285628.1

DBLINK BioProject: [PRJNA727281](#)

BioSample: [SAMN14089210](#)

KEYWORDS RefSeq.

SOURCE Aspergillus flavus NRRL3357

ORGANISM [Aspergillus flavus NRRL3357](#)

Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina;
Eurotiomycetes; Eurotiomycetidae; Eurotiales; Aspergillaceae;
Aspergillus; Aspergillus subgen. Circumdati.

REFERENCE 1 (bases 1 to 1335)

AUTHORS Fountain, J.C., Clevenger, J.P., Nadon, B., Youngblood, R.C.,
Korani, W., Chang, P.-K., Starr, D., Wang, H., Isett, B.,
Johnston, H.R.,

Wiggins, R., Chu, Y., Agarwal, G., Kemerait, R.C., Pandey, M.K.,
Bhatnagar, D., Ozias-Akins, P., Varshney, R.K., Scheffler, B.E.,
Vaughn, J.N. and Guo, B.

TITLE Two New Chromosome-Level Aspergillus flavus Reference Genomes
Reveal a Large Insertion Potentially Contributing to Isolate
Stress

Tolerance and Aflatoxin Production

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 1335)

CONSRM NCBI Genome Project

TITLE Direct Submission

JOURNAL Submitted (15-SEP-2021) National Center for Biotechnology
Information, NIH, Bethesda, MD 20894, USA

REFERENCE 3 (bases 1 to 1335)

AUTHORS Fountain, J.C., Clevenger, J.P., Nadon, B., Youngblood, R.C.,
Korani, W., Chang, P.-K., Starr, D., Wang, H., Isett, B.,
Johnston, H.R.,

Wiggins, R., Chu, Y., Agarwal, G., Kemerait, R.C., Pandey, M.K.,
Bhatnagar, D., Ozias-Akins, P., Varshney, R.K., Scheffler, B.E.,
Vaughn, J.N. and Guo, B.

TITLE Direct Submission

JOURNAL Submitted (30-JUL-2020) Plant Pathology, University of Georgia,

2747 Davis Rd., Tifton, GA 31793, USA

COMMENT PROVISIONAL [REFSEQ](#): This record has not yet been subject to final NCBI review. This record is derived from an annotated genomic sequence (NC_054693).

COMPLETENESS: incomplete on both ends.

FEATURES Location/Qualifiers

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[CDS](#) 1..1335

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ORIGIN

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