

PLEASE PROVIDE COMPLETE INFORMATION

Reprints or other data relating to this deposit will aid the Stock Center and recipients of the strain.

Accession number

SPECIES Aspergillus nidulans A 866

GENOTYPE SulA1 anA1 biA1; wA2 cnxE16; galA1; methG1; facA303; sbA3; choA1; fwA1

&
LINKAGE GROUP(S) I I I II II III IV V VI VII VIII

YOUR STOCK NUMBER FOR THIS CULTURE EK 3615
include stock no. from other collections

ORIGIN OF STOCK from diploid ^{EK} 2214

2n 2214 = ^M 3544 (from 2n 2721) + + YA2 + ^I AcuA1 + + ^{II} + + ^{III} + + ^{IV} + + ^V nicA2 + ^{VI} sB₃ + ^{VII} malA1 ^{VIII} fwA2 +
M 2488, (from 2n 2478) SulA1 anA1 + biA1 + wA2 cnxE16 galA1 methG1 + facA₃₀₃ + sbA₃ choA1 + ⁿ p₂ w

← see back ←
for example - obtained from, genetic background, from cross with; or if collected from nature, collection point, substrate and collector.

PUBLISHED REFERENCES _____

(for any information regarding this stock)

IF UNPUBLISHED, please indicate strain of origin, mutagen, worker, genetic background, important characteristics _____

cnxE16 can be identified by chlorate resistance,
use 150 mM KClO₃ in CM

COMMENTS (special growth requirements, aberrations, heterokaryon compatibility, special uses of strain, etc.)

white* Master (=mitotic mapping) strain ^(markers on all 8 chromosomes)

* When conidiation is poor on haploidizing media, white
segregants are the most visible types
(use back of page if necessary)

YOUR NAME Etha Kafer DATE March 29/98