Fungal Genetics Stock Center Dept. of Microbiology Univ. of Kansas Medical Center Kansas City, KS 66103-7240

PLEASE PROVIDE COMPLETE INFORMATION

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

Accession number

GENOTYPE..... cspA1; (acrA1)* brnA2; fpaD19; choA101; thiB101; metB11; pdxA2; oliC2; cmB12..... DESIGNATION OF MUTANT ALLELES...1.....1.....2.....19.....101.....101.....11.....2......2.....12...... STRAIN DESIGNATION IF WILD-TYPE YOUR STOCK NUMBER FOR THIS CULTURE...... EK226 include stock no. from other collections * acrA1 is closely linked to brnA2, and crossovers were encountered very rarely; (acrA1) means not retested ORIGIN OF STOCK..... from 2n (051): EK200 / EK193 from 2n (032).... EK200 = FGSC# <u>A962</u>.... for. 2n (032)...see <u>A979</u>..... for example - obtained from, genetic background, from diploid with; or if collected from nature, collection point, substrate and collector. PUBLISHED REFERENCES..None; this is the second Super Master Strains (see FGSC# A982) Review ref. for strains, mutants and mapping strains: <u>Bos</u> et al.(<u>1993</u>) Appl. Microbiol. <u>Biotechnol.</u> <u>38</u>:742-745 Basic reference for mapping by mitotic crossing over: <u>Debets</u> et al. (<u>1993</u>) Curr. Genet. <u>23</u>: 47-53)..... (for any information regarding this stock) IF UNPUBLISHED, please indicate strain of origin, mutagen, worker, genetic background, important characteristics. Strain of origin for all strains: FGSC# A733 New mutants are mapped by haploidization of well-marked heterozygous 2n and crossovers for mutations located on the same chromosome are rare; however, in A. niger, they are more frequent than in A.nidulans. COMMENTS (special growth requirements, aberrations, heterokaryon compatibility, special uses of strain, etc.) As described by <u>Debets</u> et al.<u>1993</u> (see above, & <u>1990</u>, Mol. Gen. Genet. <u>221</u>: 453-458) haploid crossover segregants are often found in A. niger, especially for mutations that map on opposite chromosome arm. This strain is a useful mapping strain which grows well on standard CM (addition of MET may improve conidiation). (use back of page if necessary)