

Aspergillus niger pedigree information for/on

## A. niger FGSC Deposition sheets -->

FGSC# my sg No. Origin: Mutagenesis or segregants (or recombinants) from diploids

A954 = EK053 UVof N724 [EK036] = FGSC# A909 --> *choA101*

A955 = EK054 UV of N724 [EK036] = FGSC# A909 --> *cysA101*

A956 = EK163 UV of EK132 --> *thiB101*

EK132 = cspA1; fwnA1; choA101; pdxA2, from 2n (006)

2n (006) : EK028 / EK053 = FGSC#A954 (above)

EK028 = cspA1; fwnA1; lysA7; leuA1; pdxA2 from 2n (001)

2n (001) : N409 = FGSC#A798 [EK017] / N690 = FGSC#A806 [EK019]

A957 = EK164 UV of EK132 from 2n (006) [see FGSC#A956 above]

A958 = EK211 X-ray of EK146 --> *sftC102*

EK146 = cspA1; fwnA1; leuA1; pdxA2, from 2n (016)

2n (016) : N694 = FGSC#A900 / EK028 from 2n (001) [see FGSC#A956]

A959 = EK217 DEO of EK133

EK133 = cspA1; (acrA1)\* brnA2; choA101; pdxA2 from 2n (006) [see FGSC#A956]

A960 = EK189 from 2n (042) : N814 = FGSC#A920 [EK034] / EK133 (see preceding FGSC#A959)

A961 = EK196 from 2n (037) : EK171 / EK164 = FGSC#A957 (see above)

EK171 cspA1; (acrA1)\* brnA2; fpaD19; lysA14; metB11; oliC2, from 2n (031)

2n (031) : N784 [EK131] / EK138

N784 = new mutant (fpaD19) in cspA1; fwnA1; lysA14 (F.Debets, unpubl.)

EK138 = from 2n (023)

2n (023): N907 = FGSC#A942 [EK106] / EK112, from diploid (002)

EK112 = cspA1; (acrA1)\* brnA2; choA101; nicA1; pdxA2; trpB2

2n (002): N837 = FGSC#A925 [EK040] / EK053 = FGSC#A954 (above)

A962 = EK200 from 2n (036) : EK171 (see preceding FGSC#A961) / EK163 = FGSC#A956 (above)

A963 = EK218 from 2n (052) : EK210 = FGSC#A974 / EK175

EK175 from 2n (031) : N784 [EK131] / EK138 (see FGSC#A961 above)

A964 = EK134 from 2n (008) : N694 = FGSC#A900 [EK032] / EK053 = FGSC#A954 (above)

EK053 UV of N724 = FGSC#A909 --> *choA101*

A965 = EK140 from 2n (016) [see FGSC#A958, above]

A966 = EK141 from 2n (016) [see FGSC#A958]

(acrA1) indicates the likely presence of acrA1, closely linked to brnA2, but not confirmed by growth tests