List of Aspergillus niger strains for deposit at FGSC (Kafer Lab, SFU; Febr. 1996)

Four types of strains are being deposited at FGSC:

i) mutations in new genes,

- ii) strains with more than one marker on a chromosome (i.e. rare crossovers)
- iii) strains suitable for transformation
- iv) strains for mitotic mapping

EK numbers = chronological numbers for our stocks in silicagel

Roman numerals in genotypes indicate Linkage groups of each marker. Note:

- a) All A.niger strains and progeny currently used for genetic analysis carry cspA1.
- b) When the cloning of genes identifies A. niger genes as homologous to those of A. nidulans a change in terminology is made to indicate such findings; therefore pyrA1(5,6) => pyrG1(5,6); argB2 => argL2, a new mutation is argB13; niaA2 => niaD2. Only the new Terminology is shown.
- c) Mutants growing on proline are all called **pro**, even if they also grow on arginine &/or ornithine; all other mutants growing on arginine are called **arg**, whether they grow on ornithine or citrulline or not.

FGSC	Our	Genotype	Origin: 2n or
# sg No. # of strain of origin i) Mutations in new genes (or new alleles)			
		cspA1; (I) acrA1* brnA2; (III) choA101 ; (V) nicA1	`(UVof N724)
		cspA1; (I) acrA1 brnA2; (III) cysA101; (V) nicA1	(UVof N724)
A956	EK163	cspA1; (I) fwnA1; (III) choA101; (IV) thiB101; (VI) pdxA2	(UVof EK132 from 2n 006)
A957	EK164	cspA1; (I) fwnA1; (III) choA101; (VI) pdxA2; (VII) nicB101	(UVofEK132)
A958	EK 211	cspA1; (I) fwnA1; (IV) leuA1; (VI) pdxA2; (VII) sftC102 (se ^r =se	elenate resist.)(X-ray of EK146)
A959	EK 217	cspA1; (I) (acrA1)* brnA2;(III) choA101;(VI) pdxA2; (?) sftB103	(DEO of EK133)
* acrA1 is closely linked to brnA2, and no crossovers were encountered; (acrA1) means not retested.			

ii) Double mutants in same chromosome (rare crossovers)

A960 EK189 cspA1; (I) fwnA1; (III) pyrG5 choA101; (VII) nicB5 (042.5.8)
A961 EK196 cspA1; (I-V) acrA1 brnA2; fpaD19; choA101 lysA14; (±ntrB3); metB11; (VII) nicB101 (037.1.6)
A962 EK200 cspA1; (I-IV) fwnA1; fpaD19; lysA14; ntrB3_thiB101; (VII) oliC2 (036.2.19)
A963 EK218 cspA1; (I) fwnA1; (III) choA101; (VI-VIII) pdxA2; oliC2 sftC101; crnB12 (052.2.2)