

Hsu, K. S. Drug resistant mutants.

Drug resistant mutants have been isolated and tested for their mode of inheritance, whether

Mendelian or non-Mendelian. The aim was to add a new class of Mendelian markers besides the visible and nutritional ones, and to search for cytoplasmic determinants. No cytoplasmic mutants have been found, but gene controlled resistance to the drugs acriflavine, actidione, and caffeine has been demonstrated. The results obtained are summarized in the following table:

Isolation No.	Drug	Resistance level ($\mu\text{g/ml}$)	Origin	Locus and linkage group
KH1	acriflavine	2	spontaneous	acr-1 (II.)
KH2	"	10	spontaneous	acr-2 (IIIL)
KH3	"	10	U.V.	- -
KH4	"	50	U.V.	acr-2 (IIIL)
KH5	"	50	spontaneous	- -
KH6	"	50	spontaneous	acr-2 (IIIL)
KH7	"	50	U.V.	- -
KH51	actidione	10	U.V.	- -
KH52	"	10	U.V.	act-1 (IR)
KH53	"	10	U.V.	- -
KH101	caffeine	2500	spontaneous	- -

All in St. Lawrence background: A dash indicates that analysis is in progress.

With the exception of KH1 and KH2, which were picked up during serial subculture on acriflavine-containing minimal medium, all of the mutants were recovered by plating conidia on Vogel's minimal medium plus the drug. The concentration of the drug was such that no growth was observable when 10^6 to 10^7 conidia from the sensitive strain were plated. U.V.-irradiation giving 50-75% killing was applied in some cases. Strains displaying colonial growth and conidiating well were selected for use in the experiments. These included cr;cot; cr;cot;ylo; cr;bal; and rq cr, all in a background of St. Lawrence stocks 74A and 73a.

Scoring for resistant vs. sensitive is clear-cut for all of the three categories of mutants. The gene acr-2 is the first gene to be located in the left arm of linkage group III. In a three-point cross acr-2 (allele KH2) x sc tryp-1, where sc is the centromere marker, the following phenotypic classes were observed:

(acr + + 74	(+ + + 1
(+ sc tryp 76	(acr sc tryp 6
(acr + tryp 28	(+ + tryp 0
(+ sc + 31	(acr sc + 0