

TABLE 1

Mutant	Linkage group and arm	Strain of Origin		Mode of origin	New or Induced Translocations	Fungal Genetics Stock Center Nos.	
		Genotype	Translocation			Original strain or untested descendant	Translocation free descendants
abA1		see ileA3					
AcrA1	II L	pabaA1 yA2; coA1	T1(VI;VII)	S	none	36	204,360
adC1 (wA2)	II R	yA2 (veA ⁺)	T1(III;IV;VIII)	X-rays	none		259,344
adI50	III L	biA1	none	UV	?	285	
biA1 (veA1)	I R	wild type (veA ⁺)		X-rays	none	26	194,357,391 392
bwA1	VI R	proA1 yA2; wA3 adG14 pabaA1 biA1	T1(III→VIII) T1(VI;VII)	S	none	lost	112-115,366
chaA1	VIII R	biA1; choA1	T1(I;VII)	S	none	lost	204,360
chaA2	VIII R	biA1; sA2	T1(V;VIII)	S	none	372	none
cnxB2	VIII R	AcrA1 wA3; nicB8	none	?	?		449
cnxE16	II R	biA1; wA3	T1(III→VIII) T1(VI;VII)	UV	none		337,462
cnxH3	III L						490
coA1	VIII R	pabaA1	T1(VI;VII)	S	none	29	120,449
dilA1	III L	yA2; pyroA4	none	UV	?		489,503
facA303	VR	wA3; pyroA4	T1(VI;VII)	S	?		502
facB101	VIII R	wA3; pyroA4	T1(VI;VII)	S	?		409
facC102	VIII R	wA3; pyroA4	T1(VI;VII)	S	none?	255	427
fpA91	I L	proA11 yA2; pyroA4	(no T?)	NG	?	387	390
fpaB37	IL	riboA1 biA1	none	S	none	422	507
frA1	IV R	yA2; pyroA4	none	UV	T1(IV;VIII)	470	none
fwA1	VIII R	biA1	none	S	none	378	379
fwA2	VIII R	AcrA1; lysB5	none	S	none	371	469,485
galA1	III L	biA1; wA3	T1(III→VIII) T1(VI;VII)	UV	?		211
galD5	I L	biA1; wA3	" "	UV	?		494
galE9	III L	biA1; wA3	" "	UV	?		214
hxA2	IV R	biA1	none	DES	?		258,491
ileA1	II R	biA1; sB3	none	UV	none?	295	
ileA3	II R	biA1	none	UV	?		505