

Table 1 (continued)

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
iodA1	II L	biA1; wA3; nicB8	T1(III→VIII)	S	none?		492
lysA1	VI R	wA1 (veA ⁺)	none	X-rays	"AbV1"	38,350	none
lysF88	I R	biA1; sB3	none?	NA	none	293	376
malA1	VII R	yA2; pyroA4	none	UV	?	57	437
meaB6	III L					451	496
ni3		see cnxE16					
ni21		see cnxB2					
ni51		see nirA14					
nirA14	VIII R	biA1; phenA3	T1(I;IV)	S	none	260	472
nicA2	V	wild type (veA ⁺)		X-rays	?		437
nicB8	VII R	biA1	none	UV	?		122
oliA2	VII R	pabaA1 yA2	none		none	450	453
pA2	V R	adG14 biA1	none	S	none	35	460
pabaA1	I R	biA1	none	X-rays	T1(VI;VII)		187,408
pabaB22	IV R	biA1	none	UV	?		426,493
palB7	VII R	rA1 biA1	none	UV	?	244	381
palC4	IV R	rA1 biA1	none	UV	?		497
palF15	VII R	rA1 biA1	none	UV	?	247	400
pantoA1	III or VIII	yA2; thiA1	T1(III;IV;VIII)	X-rays	T(VII;III or VII;VIII)	340	none
pantoB100	VII R			UV	?	397	453
phenB6	VII R	biA1	none	UV	?	394	499
proA94	I L	wild type (veA ⁺)		UV	?	386	393
riboD5	V R	biA1	none	UV	?	258	504
sA4	III L	biA1	none	UV	?	2	339
sB3	VI R	biA1	none	UV	?	41	369
sC12(=s0)	III L	wild type (veA ⁺)		NM	?	24	488
sD85	VIII R	biA1	none	NG	?	419	421
sE15	VIII R	biA1	none	NG	?	385	399
sF211	VII R	biA1	none	NG	none	388	403
sbA3	VI R	biA1; wA3	{ T1(III→VIII) T1(VI;VII)	UV	T2(I;VII)		500,510
smA1	III L	biA1; lysB5	none	S	none	66	501