

FGSC#	Genotypes	Translocations	Origin	Montreal stock number
57	yA2 pyroA4;malA1	no T	UV of #220	M 895
66	biA1;smA1;lysB5	no T	UV of #26	M 905
68	Mitotic Mapping strain	no T	Cross of Forbes MSD	M 911
69	biA1;wA4;smA1;lysB5	no T	S in #66	M 912
70	Mitotic Mapping strain	no T	C 434 [2]	M 915
71	suA1adE20 riboA1 pabaA1 yA2 adE20;AcrA1 wA2	no T	C KK (3)	M 59
72	adG14 proA1 pabaA1 yA2;wA3	no T	BC I [1]	M 140
75	suA1adE20 adE20 biA1;wA2;choA1;chaA1	no T	C 362 [2]	M 391
78	wA3	no T	C 419 [1]	M 552
79	Mitotic Mapping strain	no T	C 400 (4)	M 626
83	yA2; (veA ⁺)	no T	BC VIII [1]	M 643
84	yA2;chaA1	no T	C 419 [1]	M 701
85	adG14 proA1 pabaA1 yA2;wA3 (veA ⁺)	no T	BC VIII [1]	M 765
87	riboA1 adG14 proA1 pabaA1 yA2	no T	Cross X (2)	M 841
89	biA1;argB2*	T (?)	UV of #26	M 894
90 - 97	not of Glasgow origin (Table 4, Barratt et al 1965)	(insert A)		Grindle
104	Mitotic Mapping strain	no T	C 282 (4)	M 301
105	Mitotic Mapping strain	no T	C 280 (4)	M 276
108	adE20 biA1;wA2 cnxE16;sC12;methG1 nicA2;choA1;chaA1	no T	C 349	M 455
111	proA2 biA1	T1 (II;III)	UV of #26	M 233
112 ^Y	yA2;bwA1	}	set of	M 545
113 ^Y	bwA1		protroph	M 546
114 ^Y	bwA1;chaA1		(not veA ⁺)	M 547
115 ^Y	wA3;bwA1		colour mutants	M 548
116 ^Y	yA2		from C 419	M 549
117 ^Y	prototroph, wild type colour (not veA ⁺)		(includes #78,	M 550
118 ^Y	chaA1	84 of 366	M 551	
120	pabaA1 yA2 adE20;AcrA1;coA1	no T	C 281 (4)	M 286
121	suA1adE20 riboA1 proA1 pabaA1 adE20;chaA1	no T	C 300	M 310
123	biA1 AcrA1 wA3		(=cross d, Kafer 1962)	
122	riboA1 yA2;nicB8	no T	C 4	M 15
125(T)	riboA1 biA1;AcrA1;pyroA4;sB3;nicB8	T1 (I;VIII)	C 62	M 78
126	pabaA1 yA2;wA3;(veA ⁺)	no T	BC VIII [1]	M 740
129(T)	pabaA1 yA2 adE20;choA1	T1 (I;VII)	C 103	M 96
			(=b1, Kafer 1962)	
146	Mitotic Mapping strain	no T	C 434 [2]	M 927
147	AcrA1;lysB5;sB3;chaA1	no T	C 434 [2]	M 940
148	suA1adE20 pabaA1 yA2 adE20;lysB5	no T	C 461 [2]	M 951
150	suA1adE20 adE20 biA1;phenA2;sB3;nicB8	no T	C 434 [2]	M 955
154	Mitotic Mapping strain	no T	C 500	M 1043
155	Mitotic Mapping strain	no T	C 474 [2]	M 1056
158	riboA1 biA1	no T	UV of #26	M 827
159	Mitotic Mapping strain	no T	C 283 (4)	M 1295

^YMay carry suA1adE20

^YMay carry suA1adE20, yA2 and/or chaA1

* = Poor strain, additional morphological mutants

△ meiotic mapping strain (PART)