

FG-SC#	Genotypes	Translocations	Origin	Montreal stock number
283	Mitotic Mapping strain	no T	MSF of McCully	GD (M 1655)
284	biA1;suB2palB7;palB7	no T	S in #244	G. Dorn
285	biA1;adI50 (not growing on acetate)	no T	UV of #26	G. Dorn
286	riboA1 biA1;fpaC43	no T	S in #158	G. Dorn
287	biA1;suD2palA1;palA1	no T	S in #243	G. Dorn
288	Mitotic Mapping strain	no T	MSE of McCully	G. Dorn
291 (T)	biA1;wA3;galC4	} both & with }	UV in #51	C.F. Roberts
292 (T)	biA1;wA3 lacB3			
293	lysF88 biA1;sB3	no T (?)	NA of #41, Pees	GD (M 1305)
294	biA1;sB3;lysD18 (BL)	T (?)	NA of #41, Pees	G. Dorn
295	biA1;ileA1 (in group II);sB3	no T (?)	NA of #41	E. Pees (M 1497)
296	biA1;lysE13 (EL on V);sB3	T (?)	NA of #41	E. Pees
297	lysF51 biA1;sB3	T (?)	NA of #41	E. Pees
298	biA1;lysE14 (EL on V);sB3	T (?)	NA of #41	E. Pees
299	biA1;sB3;his*L122 (EL) (*L = Leiden)	T (?)	NA of #41	E. Pees
300	biA1;sB3;lysD7 (BL on VII)	T (?)	NA of #41	E. Pees
301	biA1;lysE23T (EL on V);sB3	T (?)	NA of #41	E. Pees
303	proA1 yA2;AcrA1	T(III→VIII)?	Cross 38 (3)	R.H. Pritchard
304 (T)	proA1 pabaA1 yA2;phenA2	T(I;VIII)	Cross 2 (6)	RHP (M 1156)
305	proA2 pabaA18 biA1;phenA2;lysB5	Teaching strains	Cross of RHP	R.H. Pritchard
306	adF17 pabaA1 yA2		Cross of RHP?	R.H. Pritchard
3	proA1 pabaA1;nicB8	(Clowes and Hayes, 1968)	Cross of RHP?	R.H. Pritchard
308	pabaA18 biA1	may have T	UV of #26?	R.H. Pritchard
309	proA2 pabaA18 yA2		Cross of ?	R.H. Pritchard
310	proA1 biA1;AcrA1;pyroA4	T(III→VIII)?	Cross 38 (3)	RHP (M 1148)
311	adE20 biA1;wA2;riboD3;nicA2	} T (?)	} Cross of RHP	R.H. Pritchard
312	suA1adE20 yA2 adE20;AcrA1;phenA2;lysB5			
314	proA1 pabaA1 yA2;wA3 adC1 (veA <sup>+</sup> )	no T	Cross of Forbes, #13 x adC1 (veA <sup>+</sup> ) strain	M 821
315	adG14 pabaA1 yA2	no T	Cross V (2) of G. Ponte #37 x pabaA1 yA2;coA1	M 825
316 (T)	biA1;sA2;T1(V;VIII)	T1(V;VIII)	UV of #26	M 845
319	anA1 yA2;wA3 adC1;sC12	no T	Cross n, EKB	M 1128
320	suA1adE20 riboA1 proA1 adE20 biA1;pyroA4	T (?)	Cross g, EKB	GD (M 1100)
326	biA1;methG1;flA1	T (?)	mutant in #219?	Azevedo
327	adE20 biA1;wA3;methG1 pyroA4;uvs-5	T (?)	UV of #139	Tuveson
328	biA1 ad-20;met-3;uvsA1 (probably allele of metG)	T (?)	strain of Lanier	Tuveson
329	adE20 biA1;wA3;methG1 pyroA4;uvs-4	T (?)	UV of #139	Tuveson
330	adE20 biA1;wA3;methG1 pyroA4;uvs-77	T (?)		
332	mel (different wild type, from Birmingham)] 8-azaguanine		AZA	?
333	pabaA108 biA1;uvsB110	T (?)	UV of pabaA108 biA1	Jansen
334	pabaA108 biA1;uvsC114	T (?)		

\*See uvs- terminology.

A mitotic Mapping strain (Table)  
(T) Translocation