

FGSC#	Genotypes	Translocations	Origin	Montreal stock number
335	proA1 pabaA125 biA1;pyroA4;uvsD153	T (?) } T (?) }	UV of proA1	Jansen
336	proA1 pabaA125 biA1;pyroA4;uvsE182		pabaA125 biA1;pyroA4	Jansen
337	riboA1 biA1;wA3 thiA4 cnxE16 (=ni3)	no T	C 50	M 23
338	yA2; wA3 puA1 cnxE16 adC1 AcrB2	T (?)	C 85	M 94
339	suA1adE20 pabaA1 yA2 adE20;sA4	no T	C 136	M 146
340	proA1 biA1;phenA2;pantoA1	T1(III;VIII;VII)	C 243	M 347
341	suA1adE20 proA1 pabaA1 yA2 adE20;AcrA1 puA1; choA1 nicB8;chaA1	no T	C 347	M 426
342	suA1adE20 adE20 biA1;AcrA1;choA1;chaA1	no T	C365	M 771
343	adG14 proA1 pabaA1 yA2;wA3 (ve <sup>+</sup> ) (from BC IX)	no T	C 1620	M 799
344	yA2;wA2 adC1; sC12; (ve <sup>+</sup> )	no T	C J (2)	M 814
345	biA1;sC12;pyroA4;T1(II;IV)	T1(II;IV)	C s (2)	M 854
346	biA1;AcrA3 adC1;sC12;pyroA4	no T (?)	S in #259	M 875
347	anA1 adE20 biA1;wA2;lysB5 T1(I;V;VII;VIII)	T spont. in #348		M 900
348	anA1 adE20 biA1;wA2;lysB5 (from cross of dupl. T1(III→VIII))	no T	C 512	M 921
349	adE20 biA1;lacA1;choA1;chaA1	no T	C 496	M 917
350	biA1 phenA2;lysA1 sB3 ("Ab VI" = Inversion?)	no T	C 346	M 918
352	riboA1 adG14 yA2;phenA2 T1(I;VIII)	T1(I;VIII)	C 2 (6)	M 962
353	biA1;ornB9 riboB2	T (?)	C 494	M 966
354	AcrA1;lysB5 T1(I;VII)	T1(I;VII)	C 515	M 1014
355	anA1 adE20 biA1;wA2;lysB5 T2(I;VII) (from cross of #347 or 348)		C 542b	M 1054
356	riboA1 biA1 (stand.)	no T	C 630 [1]	M 1076
357	biA1;wA3 (replaces #51)	no T	C 640 [1]	M 1089
358	riboA1 adG14 proA1 yA2;ornA4	T (?)	C 3	M 1101
359	pabaA1;wA3	no T	C 639 [1]	M 1231
360	AcrA1;lysB5;chaA1 (5 <sup>th</sup> generation back cross)	no T	C 752	M 1239
361	riboA1 yA2	no T	C 630 [1]	M 1265
362	riboA1 anA1 adG14 proA1 pabaA1 yA2	no T	C 773	M 1267
363	suA1adE20 riboA1 anA1 luA1 pabaA1 yA2 adE20 biA1;AcrA1	no T	C 792	M 1274
364	suA1adE20 proA2 yA2 adE20;AcrA1 wA2	no T	Dipl. 629	M 1277
365	yA2;AcrA1;lysB5;chaA1 (5 <sup>th</sup> back cross)	no T	C 752	M 1279
366	yA2;chaA1;bwA1	no T (?)	C 789	M 1282
367	suA1adE20 proA1 yA2 adE20;AcrA1 wA2	no T	Dipl. 628	M 1285
368	riboA1 yA2;sA2	no T	C 785	M 1301
369	pabaA1 biA1;sB3	no T	C 788	M 1303
370	adG14;sA1;pyroA4;chaA1 T1(V;VI)	T1(V;VI)	C 838	M 1309
371	AcrA1;lysB5;fwA2	no T	S in M 1241	M 1318
372	biA1;sA2;chaA2 T1(V;VIII)	T1(V;VIII)	S in #316	M 1319
373	anA1;sA1;lysB5;chaA1 T1(V;VI)	T1(V;VI)	C 838	M 1324
374	Mitotic Mapping strain	no T	C 569	M 1335
375	Mitotic Mapping strain	no T	C 569	M 1343
376	suA1adE20 lysF88 pabaA1 yA2 adE20	no T (?)	C 841	M 1346
377	riboA1 yA2 adE20	no T	C 847	M 1368
378	biA1;fwA1	no T	S in #26	M 1370
379	(veA <sup>+</sup> ) fwA1 ornB7	no T (?)	cross of J.C.	M 1371
380	lysB5;fwA2 cnxB2 facB101 riboB2 palB7 chaA1	no T	C 804	M 1373

▲ mitotic mapping strain (table)

Not ribo B2