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Abstract

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Murray, N. E. Linkage information

for cysteine and methionine mutants.

cys-9 (T156), cys-ys located between cr (crisp) and thi-1 (thiamine-1) in lininkagegroup IR (see Table 1).

cys-f0 (339816). The tentative location of this locus (Murray 1965 Genetics 52: 801) as the most distal marker in the left arm of linkage group IV is supported by information from other workers.

<u>cys-11</u> (NM86). A cluster of cysteine mutants is located in the <u>cys-5</u> region between <u>leu-3</u> and mating type. The evidence is consistent with the region comprising two loci, <u>cys-5</u> and <u>cys-11</u>. The mutants NM44 and R83R1-1-271 gave very low recombination frequencies when crossed to <u>cys-5</u> (35001). The recombinants from the latter cross were scored for flanking markers and all four flanking marker combinations were represented. When a fourth nutant (NM86) was crossed to <u>cys-5</u> (35001) the recombination frequency was much higher and there war no or little "negative interference". Complementation tests showed that NM86 is physiologically different from the heterocaryon compatible <u>cys-5</u> (NM44) strain, and more specifically Leinweber (personal communication) has shown that while NM86 lacks ATP-sulfurylase, the <u>cys-5</u> alleles tested (35001 and NM44) lack PAPS-reductase. It is proposed that NM86 is an allele at locus <u>cys-11</u>. The combination of flanking markers found for cysteine independent recombinants from a cross of <u>cys-5</u> by <u>cys-11</u> indicare the order mating type, <u>cys-11</u>, <u>cyr-5</u>, leu-3. Adequate genetic information is lacking far a cross of <u>cys</u> (85518) by <u>cys-5</u>, but <u>cys</u> (85518) gave a very low recombination frequency (1 in 200,000) when crossed to <u>cys-11</u> (NM86).

cys-12 (NM268). cys-12: is an additional cysteine locus in linkage group I distal to ad-9 and close to at (0 recombinants among 76 isolates) (see Table).

me-6 (35809) and mac (65108). These mutants or closely linked. Methionine independent recombinants have been isolated from crosses of me-6 by mac and classified with respect to the flanking markers thi-1 and ad-9 (odenine-9). The order indicated is thi-1, mac, me-6, ad-9, but it is probable that mac and me-6 is allelic.

me-7 and me-9. Methionine independent recombinants have been isolated from crosses of me-7 (NM73) by me-9 (NM43t) and classified with respect to flanking markers (thiamine-3 and white collar). The methionine loci are very cl-linked in the order thi-3, me-7, me-9, wc.

Zygote genotype and % recombination	Parental combinations	Recombination			Total and	Marker isolation
		Singles region	Singles region 2	Doubles regions 1 and 2	% germination	numbers
+ thi-1 ad-9	3 4	7	10	0	9 3 (93%)	T156
cys-9 + +	3 3	5	4	0		56501
12.9 15.1						Y 154M37
+ cyr-9 +	2 6	2	20	0	.88	8122
cr + 0 ^{s-1}	2 3	1	16	0	(63%)	T156
3.4 40.9						B 135
<u>+ + cys-12</u>	48	8	8	0	120 (83%)	56501
thi-} ad-9 +	44	5	6	1		Y 154M37
11.7 1 2 . 5						NM268

Table 1. Linkage data on random segregants from crosses involving cys-9 or cys-12.

(The top number in each p-air represents the class that has the + allele of the leftmost marker).

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