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# Isolation and map location of a new acetate-requiring mutant, ace-9, of Neurospora crassa

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#### **Abstract**

A new acetate-requiring mutant strain of *Neurospora crassa* (ace-9), has been isolated from the double mutant strain *cot-1;inl a*, by inositol-less death (Lester and Gross 1959. Science 129:572) in Vogel's medium supplemented with 0.3% sodium acetate and 2% sucrose. The mutant grows well on complex medium and on Vogel's medium supplemented with casamino acids, acetate, or acetate plus ethanol. Like *ace-2*, *ace-3* and *ace-4* (Okumura and Kuwana 1979. Japan J. Genet. 54:235-244), it shows very weak activity of pyruvate dehydrogenase compelx, but has normal activities of pyruvate carboxylase, pyruvate kinase and glucose-6-phosphate dehydrogenase.

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### Isolation and map location of a new acetate-requiring mutant, ace-9, of Neurospora crassa

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A new acetate-requiring mutant strain of *Neurospora crassa* (ace-9), has been isolated from the double mutant strain cot-1;inl a, by inositol-less death (Lester and Gross 1959. Science 129:572) in Vogel's medium supplemented with 0.3% sodium acetate and 2% sucrose. The mutant grows well on complex medium and on Vogel's medium supplemented with casamino acids, acetate, or acetate plus ethanol. Like ace-2, ace-3 and ace-4 (Okumura and Kuwana 1979. Japan J. Genet. 54:235-244), it shows very weak activity of pyruvate dehydrogenase compelx, but has normal activities of pyruvate carboxylase, pyruvate kinase and glucose-6-phosphate dehydrogenase.

The *ace-9* gene maps between *nuc-2* and *arg-13* on the right arm of the linkage group II (Table 1). Thus the gene sequence near *ace-9* is :-pyr-4 - thr-2 -Centromere- arg-5 - nuc-2 - ace-9 - arg-12 - aro-1.

**Table 1.** Linkage data of *ace-9* in the format of Perkins (1959, Genetics 44:1185-1208)

	Recombinations					
Zygote genotype and recombination percent	Parental combination	Singles region 1	Singles region 2	Doubles regions 1 and 2	Total and percent germination	Marker isolation numbers
+ + ace-9 thr-2 arg-5 + 10.2 12.2	95 212	38 1	33 14	1	394 53%	35423 27947 KG601
+ ace-9 + pyr-4 + arg-12 19.8 3.2	51 2 144	36 14	1 7	0	253 63%	36601 KG601 UM107
+ ace-9 + nuc-2 + aro-1 1.7 9.6	15 87	1 1	1 10	0	115 76%	T28-M2 KG601 Y7655

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