Plant		Su	su	o/o su	P#	P* (comb. data)	Crossover** value
# 3	self	321	76	19.1	<.01	. 07	38.3%
	BÇ	248	168	40.4	<.01	<.01	40.4%
#11	self	350	97	21.7	>•05	. O1	43.4%
	BC	222	173	43.8	<.05	<.01	43.8%
#12	self	334	76	18.5	<.01	<.01	37.1%
	BC	221	176	44.3	<.05		44.3%

<sup>\*</sup> For deviations from 3:1 and 1:1

Crossover values are rather high but comparable to those for high-sugary plants in the same culture.

Since all, or nearly all, of each tassel presumably was composed of cell descendants of a somatic crossover cell, it is suggested that the putative crossing over occurred early in ontogeny--possibly in zygotic mitosis.

Perhaps the use of a third gene such as Ts, will clarify the situation by making possible a regular four-class backcross test.

H. S. Perry

## HARVARD UNIVERSITY Cambridge, Massachusetts

## 1. Further data on the components of the tunicate locus.

In a previous News Letter (No. 35) we reported that the two components of the Tu locus which had been separated by crossing over appeared to have slightly different effects. After additional backcrosses to the inbred Al58 to render them more nearly isogenic, this is still true. In the summer of 1961 we grew seven strains heterozygous for tuh-d, two of which were 13/16 Al58 and five were 29/32 Al58. These were compared with ten strains heterozygous for tuh-1, three of which were 5/8 Al58 and seven were 29/32 Al58.

The two types of heterozygous half-tunicate genotypes could not be distinguished by their tassels or by the external characteristics of the mature ears, but examinations of longitudinal sections of both young and mature ears revealed differences. Pistillate spikelets from plants heterozygous for tuh-1 consistently had longer rachillae, slightly thinner lower glumes, and longer and more numerous cupule hairs than those from plants heterozygous for tuh-d. In 71 paired comparisons of longitudinal sections of young ears the two types were consistently distinguishable. However, we cannot yet be certain that the differences are inherent in the two components and are not due to other genes closely linked with them.

P. C. Mangelsdorf

<sup>\*\*</sup> Computed for coupling