contaminants (which would be \underline{Bz} \underline{V}) could be detected, the percentage of \underline{Bz} \underline{V} gametes was much lower. This suggests that some of the \underline{Bz} \underline{V} recombinants detected in 1960 were due to contamination.

Table 1
The assortment of outside markers in $\frac{wx}{y}$ recombinants from the cross $\frac{Bz}{1960}$ and $\frac{wx}{1963}$

	1963 No. %		1960 No. <u>%</u>	
Bz v	18	62	63	58
bz v	9	31	27	25
Bz V	1	3.4	15	14
bz V	1	3.4	<u>3</u> 108	2.7
	29		108	

Oliver Nelson

3. The location of the waxy mutant H21.

One of the waxy alleles with which we originally worked was \underline{wx} H2l. On the basis of recombinational frequencies (\underline{wx}) in intercrosses with \underline{c} , $\underline{90}$, \underline{B} , and \underline{a} , it was felt that the most probable order was \underline{c} , $\underline{90}$, $\underline{H2l}$. It has since been shown by conventional genetic analyses that \underline{c} (Coe) is located distally to $\underline{90}$ as (\underline{Bz}) \underline{c} $\underline{90}$ (\underline{V}) .

A similar analysis has now been made for $\frac{H21}{}$. Pollen from plants of the F₁ $\frac{Bz}{bz} \frac{wxH21}{wx^{coe}} \frac{v}{v}$; $\frac{ae}{ae}$ was used to

pollinate the tester stock \underline{bz} \underline{wx}^{Coe} \underline{v} \underline{ae} . Tassel collections were also made for estimates of \underline{wx} frequency by our standard pollen scoring techniques.

In a total population of 1,571,000 pollen grains from 9 plants, 776 $\frac{Wx}{46}$ were detected or 49 x 10-5. This compares with $\frac{46}{46}$ x 10-5 estimated for the cross between \underline{C} and $\underline{H21}$ in our original experiments.

Due to poor germination and dry weather, the conventional analysis yielded a total population of only 21,698. Of these 9 (41 x 10^{-5}) were apparently $\underline{\text{Wx}}$, $\underline{\text{ae}}$ seeds. Of the nine apparent recombinants, 8 were carrying the $\underline{\text{bz}}$ marker and 1 $\underline{\text{Bz}}$ indicating a location for $\underline{\text{H21}}$ distal to $\underline{\text{C}}$ contrary to earlier hypothesis.

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4. Location of miniature seed (mn) on chromosome 2.

Crosses were made between a series of translocation stocks in which waxy (\underline{wx}) was used as a marker for the chromosomal interchanges and a miniature seed (\underline{mn}) \underline{wx} stock. These F_1 plants were then selfed, and the miniature seeds checked with iodine solution for waxy endosperm.

Slightly lower than expected ratios (25%) of waxy were obtained with all translocations except T 2-9 b. Progenies involving T 2-9 b, which has break points on the short arm of chromosome 2 at .18 and on the long arm of chromosome 9 at .22, gave 1.2% waxy seeds. It is therefore apparent that miniature seed is located on Chromosome 2.

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1. Further studies on trivalent frequency in an array of maize chromosome 2-Tripsacum interchange chromosome constitutions.

An attempt was made to synthesize additional 21 chromosome constitutions combining the available primary and secondary maize chromosome 2-Tripsacum interchange chromosomes in various ways. A number of the plants derived repeated constitutions which have been reported earlier (Genetics 51: 23-40. 1965), and showed metaphase I trivalent frequencies very similar to those described before. Four previously unknown 21 chromosome constitutions were also derived. Metaphase I trivalent frequencies from microsporocyte samples of three of these four constitutions were approximately consistent with expectation from previous findings in that: 1. a constitution fitting into the general category