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1. Interchromosomal effects of deficiencies in chromosome 1 on association.

Homozygotes for zb_1 P^{WW} As br_1 in chromosome 1 were crossed with X-rayed pollen carrying Zb_1 P^{WF} As Br_1 . Forty-four plants hemizygous for one or more of the three recessive genes were amenable to analysis at diakinesis or metaphase I. Fifteen of the deficient plants, including three monosomics showing loss of all dominant morphological markers, were variably asynaptic. Syncytes, curved spindles, and fragmentation--characteristic of asynaptic homozygotes--occurred in the deficient plants exhibiting failure of association. It seems likely that As was deleted along with linked dominant markers in the X_1 asynaptic plants. The single dose of As , contributed by the female parent, was insufficient to control normal first division association, thereby simulating the homozygous recessive.

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1. Genetic correspondence of *Tripsacum* chromosomes to their homeologs from corn.

Further progress has been made during the year in identifying *Tripsacum* chromosomes both genetically and cytologically. When a *Tripsacum* chromosome in a $2n+1$ stock substitutes in physiological function for a dominant gene in corn by covering its recessive marker allele present in the maize chromosome complement, we can locate a *Tripsacum* gene on a particular *Tripsacum* chromosome. Thus we can map the *Tripsacum* chromosomes, not by their own recessive genes, but by the ability of their dominant genes to prevent the expression

of recessive marker genes, either singly or in linked series, from corn. The data so far obtained, presented below, show that one Tripsacum chromosome carries dominant genes preventing the expression of three recessives on the short arm of chromosome 2 while another Tripsacum chromosome corresponds to the other arm. A similar situation exists with respect to chromosome 4. A single Tripsacum chromosome carries dominant genes which mask four recessive genes on chromosome 7 and another Tripsacum chromosome masks five recessive genes on the short arm of chromosome 9.

<u>Corn chromosome</u>	<u>Dominant from Tripsacum</u>
1	<u>Bm</u> ₂ *
2S	<u>Ws</u> <u>lg</u> ₁ <u>Gl</u> ₂ (does not cover <u>v</u> ₄)
2L	<u>V</u> ₄ * (does not cover <u>lg</u> ₁ <u>gl</u> ₂)
3	<u>A</u> ₁ *
4S	<u>Su</u> ₁ (does not cover <u>gl</u> ₃)
4L	<u>Gl</u> ₃ (does not cover <u>su</u> ₁)
7	<u>V</u> ₅ <u>Ra</u> ₁ <u>Gl</u> ₁ <u>I</u> _j
8	<u>J</u> ₁ *
9	<u>Yg</u> <u>C</u> <u>Sh</u> ₁ <u>Bz</u> <u>Wx</u>

*Allo-trisomic stocks which were lost but are being re-developed. The data for a more complete genetic map of Tripsacum will be forthcoming as crosses and backcrosses to multiple marker gene stocks of corn are made.

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2. A planting in Florida of perennial relatives of maize.

Arrangements have been made during the year to establish and maintain at the Montgomery Foundation of the Fairchild Tropical Garden, Miami, Florida, a collection of the perennial relatives of maize. Representative specimens of the collections of Tripsacum from Mexico and Guatemala made by Dr. Raju Chaganti and Mr. Garrison Wilkes (mentioned in last year's News Letter) were delivered to Florida and are now well established in a planting protected from frost by a sprinkler system. The planting includes all of the known species of Tripsacum, some interspecific hybrids in Tripsacum, perennial teosinte, three species of Manisuris, and Elyonurus tripacoides. In February all species except T. australe were in flower. The National Science Foundation has made a grant to the Fairchild Garden to