3. Two new alleles of the E gene found in teosinte.

Three alleles of the E gene, E^F, E^N, and E^S, have been found in maize. In a study of 550 strains of maize from South and Central America, we found that the E^N allele was the most common and the E^F allele the least common. Two other alleles, E^L and E^R, have been found in some teosinte strains. E^L and E^R specify enzymes with electrophoretic migration rates intermediate between the enzymes specified by E^F and E^N, and E^N and E^S, respectively. The teosinte heterozygotes E^L/E^R form a hybrid enzyme with an intermediate migration rate just as do the maize heterozygotes. A hybrid enzyme is also found in the E^F maize E^R teosinte hybrid plants. Other hybrid crosses are presently being made for further tests.

We have tested eight teosinte lines supplied by Dr. P. C. Mangels-dorf. Chapingo, Chilpancingo, Chalco, Arcelia, and Huixta carry only the alleles found in maize, EF, EN, or ES, while the Lake Retana, Florida, and El Valle teosintes carry only the new alleles EL or ER. Dr. Mangelsdorf has pointed out the interesting comparison that the latter three teosinte lines are quite Tripsacoid. Surprisingly, this esterase is not found in young Tripsacum seedlings. We have tested two species, T. floridamum and T. dactyloides. At the moment we cannot tell whether the absence of the enzyme is due to absence of the E gene or simply a different distribution of gene activity in the life cycle of Tripsacum.

Drew Schwartz

OHIO AGRICULTURAL EXPERIMENT STATION Wooster, Ohio Department of Agronomy

1. Evidence for linkage of genes for stalk-rot susceptibility with genes located in chromosomes 7 of inbreds Mo2lA and NC34.

Translocation studies (Agron. Jour. 49:197-201) have shown that major genes for resistance to Helminthosporium turcicum leaf blight are located in the short arms of chromosomes 7 of inbreds Mo21A and NC34. Tests designed to locate these genes more precisely were grown at Columbus, Ohio, using a chromosome 7 marker gene stock. Plants from the backcrosses (02 V5 ral gl x Mo21A) x 02 V5 ral gl and (02 V5 ral gl x NC34) x 02 V5 ral gl were inoculated with ground leaf inoculum of H. turcicum. Difficulties of classification made the gene V5 unusable.

Leaf blight failed to reach a high level of disease incidence by rating time. Although it appeared doubtful that good differences in disease reaction could be obtained, individual plants from each of the backcrosses were scored for H. turcicum. Statistical analyses of the data failed to show any significant differences between the various gene classes for H. turcicum reaction.

A natural infestation of stalk rot, most likely <u>Gibberella zeae</u>, caused many of the plants to die prematurely. These were noted as dead plants since it was difficult, as well as impracticable, to score them for leaf blight.