

sectored kernel selected from a self-red ear of one of these families. The phenotype of this mutant was non-variegated. Further study will be conducted in an attempt to determine the cause of the pericarp sectoring observed in this group of families.

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III. REPORT ON MAIZE COOPERATIVE

Work of the past season was concentrated principally on improving plant vigor and increasing seed supplies of the traits and tester combinations listed in the accompanying catalogue of stocks. Approximately 35,000 plants, comprising 2,200 families, were grown last summer; about 15,000 pollinations were made. Additional plantings were made in greenhouse and Florida generations.

Further crosses were made to derive new tester combinations or to determine chromosome positions of unmapped traits. It is hoped that this work, along with research on newly-acquired traits, may be intensified next season.

As time permits, all stocks are being gradually converted to the inbred lines M14, W23, and Oh51A. As a consequence of this program of developing stocks adapted to the Corn Belt, a considerable amount of effort is required to re-extract tester combinations and to confirm genetic constitutions.

Seed requests have risen sharply during the past few years. Two to three months each year are now required to supply stocks and provide information on the classification, use, and linkage relations of genetic traits.

The following listing of Maize Cooperative stocks includes the more useful combinations now available. Seed requests should be sent to the Botany Department, University of Illinois, Urbana, Illinois.

Chromosome 1ad₁ an₁ bm₂ad₁ Knan₁ Kn bm₂

as

Hm

Kn

Kn Ts₆lw₁

necrotic 8147-31

pCR

pCW

pMO

PRR ad₁ an₁PRR ad₁ bm₂PRR an₁ gs₁ bm₂PRR br₁ f₁ an₁ gs₁ bm₂PRR br₁ f₁ gs₁ bm₂

pVV

pWR bm₂pWR gs₁ bm₂pWW br₁ f₁ bm₂pWW br₁ f₁ an₁ gs₁ bm₂sr₁ pWR an₁ bm₂sr₁ pWR an₁ gs₁ bm₂sr₁ zb₄ pWWts₂ pWW br₁ bm₂Ts₃Ts₆v₁₉ bm₂V₈V₈ an₁ bm₂vP₅vP₈zb₄ ms₁₇ pWW

Chromosome 1 (Continued)zb₄ pWW bm₂zb₄ pWW br₁zb₄ ts₂ pWWChromosome 2al lg₁al lg₁ gl₂ B skal lg₁ gl₂ b skba₂fl₁lg₁ gl₂ Blg₁ gl₂ blg₁ gl₂ b fl₁ v₄lg₁ gl₂ b fl₁ v₄ Chlg₁ gl₂ B gs₂lg₁ gl₂ b gs₂ v₄lg₁ gl₂ b gs₂ v₄ Chlg₁ gl₂ B sk v₄lg₁ gl₂ b sk v₄lg₁ gl₂ b sk fl₁ v₄lg₁ gl₂ B v₄lg₁ gl₂ b v₄lg₁ gl₂ b v₄ Chlg₁ gs₂ b v₄ws₃ lg₁ gl₂ Bws₃ lg₁ gl₂ bws₃ lg₁ gl₂ b fl₁ v₄ws₃ lg₁ gl₂ B skws₃ lg₁ gl₂ b sk

Chromosome 3

A₁ ga₇; A₂ C R
 A₁ sh₂; A₂ C R
 Ad-31; A₂ C R
 Ad-31 sh₂; A₂ C R
 a^P et; A₂ C R Dt₁
 a₁; A₂ C R B Pl dt₁
 a₁ et; A₂ C R Dt₁
 a₁ sh₂; A₂ C R Dt₁
 a₁ sh₂; A₂ C R dt₁
 a₁ sh₂ et; A₂ C R Dt₁
 a₁st sh₂; A₂ C R Dt₁
 a₁st et; A₂ C R Dt₁
 a_x-1; A₂ C R
 a_x-3; A₂ C R
 a_x-3 et; A₂ C R
 an₂ = allele of d₁
 ba₁
 Cg
 cr₁
 d₁
 d₁ Cg
 d₁ gl₆
 d₁ gl₆ Lg₃
 d₁ lg₂
 d₁ Lg₃
 d₁ Lg₃ Rg
 d₁ Pg₂
 d₁ Rg
 d₁ rt
 d₁ ts₄ lg₂
 d₂
 gl₆
 gl₆ lg₂ a₁ et; A₂ C R Dt₁
 gl₆ Lg₃

Chromosome 3 (Continued)g¹₆ Rgg¹₆ v¹⁷g¹₆ v¹⁷ lg₂g¹₇lg₂ A₁^b et; A₂ C R Dt₁lg₂ a₁ et; A₂ C R Dt₁lg₂ a₁ sh₂ et; A₂ C R Dt₁lg₂ a₁st et; A₂ C R Dt₁lg₂ pmLg₃pg₂

pm

ra₂ra₂ lg₂ pmra₂ Rg

Rg

rt; A₁ A₂ C Rts₄ na₁v¹⁷vP₁

Primary trisome 3

Chromosome 4bm₃bt₂

de (1 or 16?)

Ga₁ Su₁ga₁ su₁g¹₃j₂j₂ g¹₃la su₁ g¹₃la su₁ Tu g¹₃

Chromosome 4 (Continued)

lo Su₁
 lo su₁
 lw₄; lw₃
 o₁
 sp₁ su₁
 st
 su₁ bm₃
 su₁ gl₃
 su₁ gl₄
 su₁ j₂ gl₃
 su₁ o₁
 su₁ ra₃
 su₁ Tu
 su₁ Tu gl₃
 su₁ zb₆
 su₁ zb₆ gl₃
 su₁ zb₆ Tu
 su₁ am
 Ts₅
 Ts₅ st
 Ts₅ su₁
 Tu gl₃
 v₈

Chromosome 5

a₂; A₁ C R
 a₂ bm₁ bt₁ bv₁ pr; A₁ C R
 a₂ bm₁ pr v₂; A₁ C R
 a₂ bm₁ pr ys₁; A₁ C R
 a₂ bt₁ pr; A₁ C R
 a₂ bt₁ pr ys₁; A₁ C R
 a₂ pr; A₁ C R
 ae

Chromosome 5 (Continued)bm₁ pr; A₁ A₂ C Rbm₁ pr v₂; A₁ A₂ C Rbm₁ pr ys₁; A₁ A₂ C Rbm₁ pr ys₁ v₂; A₁ A₂ C Rbm₁ yg₁bt₁ pr; A₁ A₂ C RGa Bt₁ga bt₁g¹⁵g¹⁸g¹⁷ a₂ bt₁ v₂; A₁ C Rg¹⁷ v₂intensifier of pr closely linked to bt₁lw₂lw₃; lw₄na₂na₂ prpr; A₁ A₂ C Rpr ys₁; A₁ A₂ C Rsh^{fl} = "sh₄""sh₃" = allele of bt₁

tn

v₃ pr; A₁ A₂ C Rv₁₂vp₂ g¹⁸vp₂ pr; A₁ A₂ C Rvp₇vp₇ pr; A₁ A₂ C RChromosome 6at = allele of si₁po Y₁ plpo y₁ pl

Chromosome 6 (Continued)

Pt
 si₁ Y₁ Pl
 si₁ Y₁ pl
 si₁ y₁ pl
 y₁ l₁₀
 Y₁ ms (1?)
 y₁ ms (1?)
 y₁ pb₄ Pl
 y₁ pb₄ pl
 Y₁ PG11: wx PG12
 y₁ PG11: wx PG12
 y₁ Pl Bh
 y₁ pl Bh
 Y₁ Pl sm py; A₁ A₂ b PRR
 Y₁ pl su₂
 y₁ pl su₂
 Y₁ Pl; seg w₁
 Y₁ pl; seg w₁
 y₁ Pl; seg w₁
 y₁ pl; seg w₁
 "male sterile-silky" = allele of si₁
 "orobanche" (seedling)
 "ragged" (seedling)
 "white 8522" (seedling)
 "white 8896" (seedling)

Chromosome 7

bd
 Bn₁
 g₂
 gl₁ ij bd
 gl₁ sl Bn₁
 Hs

Chromosome 7 (Continued)

ij

in; pr A₁ A₂ C Ro₂o₂ gl₁ slo₂ gl₁ sl Bn₁o₂ ra₁ gl₁o₂ v₅ gl₁; seg ra₁o₂ v₅ ra₁ gl₁o₂ v₅ ra₁ gl₁ Hsra₁ gl₁Tp₁v₅ gl₁ Tp₁va₁vp₉ gl₁; wxChromosome 8

mn

v₁₆ ms₈ j₁v₁₆ ms₈ j₁; l₁

"necrotic 6697" (seedling)

"sienna 7748" (seedling)

Chromosome 9au₁ au₂Bf₁bk₂ ms₂₀bk₂ Wcbm₄C sh₁ wx; A₁ A₂ Rc sh₁ wx; A₁ A₂ Rc sh₁ wx gl₁₅; A₁ A₂ Rc wx; A₁ A₂ R

Chromosome 9 (Continued)

c wx bk₂; A₁ A₂ R
 Dt₁ (See Chromosome 3 stocks)
 I wx; A₁ A₂ R Pr B pl
 I wx; A₁ A₂ R pr B pl
 K₉^L C sh₁ wx; A₁ A₂ R
 l₇
 ms₂
 ms₂ sh₁; A₁ A₂ C R
 ms₂₀
 sh₁ wx d₃
 sh₁ wx l₇
 sh₁ wx pg₁₂; Y PG₁₁ pl
 sh₁ wx v₁
 wx ar
 wx Bf₁
 wx bk₂
 wx d₃
 wx da₁; A₁ A₂ C R
 wx g₄
 wx l₆
 wx pg₁₂; Y PG₁₁
 wx PG₁₂; Y PG₁₁
 wx^a
 yg₂ c sh₁ wx; A₁ A₂ R
 yg₂ C sh₁ bz wx; A₁ A₂ R
 Primary trisome 9

Chromosome 10

a₃
 bf₂
 du₁
 g₁
 g₁ l₂

Chromosome 10 (Continued)g₁ r_B; A₁ A₂ Cg₁ r sr₂gl₉l₁; v₁₆ ms₈ j₁li g₁ R; A₁ A₂ Cli g₁ r; A₁ A₂ Cli g₁ r; A₁ A₂ C; carries abnormal 10nl₁ g₁ R; A₁ A₂ COg R; A₁ A₂ C B PlR_B sr₂r^r sr₂Rmb; A₁ A₂ CRnj; A₁ A₂ CRst; A₁ A₂ Cv₁₈w₂

zn

"oil yellow" (seedling and plant)

Primary trisome 10

Unplaced genes

cl

ct

de₁₇

dv

dy

fl₂gl₁₁gl₁₂gl₁₄gl₁₆gl_g

h

Unplaced genes (Continued)

l₃
 ms₅
 ms₆
 ms₇
 ms₉
 ms₁₀
 ms₁₁
 ms₁₂
 ms₁₃
 ms₁₄
 Mt
 New starchy
 rd
 Rs₁
 rs₂
 "sh₅"
 lw₁
 lw₂
 v₁₃
 va₂
 vp₆
 wi
 ws₁ ws₂
 zb₁
 zb₂
 zb₃

Multiple gene stocks

A₁ A₂ C R^F Pr B Pl
 A₁ A₂ C R^E Pr B Pl
 A₁ A₂ C R^E Pr B pl lg₁ y
 A₁ A₂ C R Pr
 A₁ A₂ C R Pr wx

Multiple gene stocks (Continued)

A₁ A₂ C R Pr wx gl₁
 A₁ A₂ C R Pr wx y
 A₁ A₂ C R pr
 A₁ A₂ C R pr su₁
 A₁ A₂ C R pr su₁ y wx
 A₁ A₂ C R pr y gl₁
 A₁ A₂ C R pr y wx
 A₁ A₂ C R pr y wx gl₁
 A₁ A₂ c R Pr su₁
 A₁ A₂ c R Pr y wx
 A₁ A₂ c R Pr y sh₁ wx
 A₁ A₂ C r Pr su₁
 A₁ A₂ C r Pr su₁ y gl₁
 A₁ A₂ C r Pr y wx
 A₁ A₂ C r Pr y sh₁ wx
 bm₂ lg₁ a₁ su₁ pr y₁ gl₁ j₁ wx gl₁
 colored scutellum
 lg₁ su₁ bm₁ y₁ gl₁ j₁
 su₁ y₁ wx a₁ A₂ C R^B pr
 y₁ su₁ ra₁ gl₁
 y₁ wx gl₁

Popcorns

Amber Pearl
 Black Beauty
 Hulless
 Ladyfinger
 Ohio Yellow
 Red
 South American
 Supergold
 White Rice

Exotics and Varieties

Argentine Popcorn

Black Mexican Sweet Corn (with B chromosomes)

Black Mexican Sweet Corn (without B chromosomes)

Gourdseed

Maiz chapolote

Papago Flour Corn

Parker's Flint

Strawberry Popcorn

Tama Flint

Tom Thumb Popcorn

Zapaluta chica

Chromosome rearrangements

The following rearrangements are being maintained primarily for use in determining the chromosome locations of new traits. All are marked with closely-linked endosperm or seedling traits.

The cytological positions of Inv 2a were determined by Dr. Morgan. those of Inv 9a were determined by Dr. Li. The indicated interchange points of the reciprocal translocations are taken from published work of Dr. Longley.

Inversions

lg ₁ or gl ₂ Inv 2a (also available with Ch)	2S.7; 2L.8
wx Inv 9a	9S.7; 9L.9

Reciprocal translocations

wx 1-9c	1S.48; 9L.22
wx 1-9 4995	1L.19; 9S.20
wx 2-9b	2S.18; 9L.22
wx 3-9c	3L.09; 9L.12
wx 3-9 5775	3L.09; 9S.24
wx 4-9b	4L.90; 9L.29
wx 4-9 5657	4L.33; 9S.25
wx 4-9g	4S.27; 9L.27
wx 5-9a	5L.69; 9S.17
wx 5-9c	5S.07; 9L.10
wx 5-9 4817	5L.06; 9S.07
wx 5-9 5614	5L.09; 9L.06
wx 6-9a	6S.79; 9L.40
wx, y 6-9b	6L.10; 9S.37
wx 6-9 4505	6L.13; 9 cent
wx 6-9 4778	6S.80; 9L.30
wx 7-9a	7L.63; 9S.07
wx or gl ₁ 7-9 4363	7 cent; 9 cent
wx 8-9d	8L.09; 9S.16
wx 8-9 6673	8L.35; 9S.31
wx 9-10b	9S.13; 10S.40
su 1-4a (also available with pRR)	1L.51; 4S.69
su 1-4d (also available with pRR)	1L.27; 4L.30
su 4-5j	4L.21; 5L.36
su, y 4-6a	4L.37; 6L.43
su 4-8a	4S.59; 8L.19
su, R 4-10b	4L.15; 10L.60
y 1-6c (also available with pRR)	1S.25; 6L.27
gl ₂ 2-3c	2S.46; 3S.52
gl ₂ 2-3 5304	2S.62; 3L.29
gl ₂ 2-6b	2S.69; 6L.49
gl ₂ , R 2-10b	2S.50; 10L.75
gl ₁ 6-7 4545	6L.25; 7S.73

Stocks of A-B chromosome translocations

B-1a	1L. 2	Proximal to <u>Hm</u>
B-1b	1S. 05	
B-3a	3L. 1	
B-4a	4S. 25	Proximal to <u>su</u> ₁
B-7b	7L. 3	Proximal to <u>ta</u> ₁
B-9a	9L. 5	
B-9b	9S. 4	Between <u>C</u> and <u>wx</u> ; close to <u>wx</u>
B-10a	10L. 35	Proximal to <u>g</u> ₁

-- Earl B. Patterson