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TO STUDENTS OF MAIZE GENETICS:-

You who attended the "cornfab" in my hotel room at the time of the winter science meetings in New York will recall that I promised to prepare a summary of the published data involving linkage groups in maize, to add my own unpublished data, and to send these records to each of you for criticism and the addition of such unpublished records as you may care to furnish me. I am now enclosing the records promised, but can claim no credit for having assembled them. Professor Fraser had, before leaving for a year in Europe, abstracted the available published papers. Mr. Beadle has completed that work, has assembled my own unpublished records, and has arranged all the tables and charts.

I hope that each of you, whether or not you attended the New York meeting, will send me such relevant data as you have not yet published, showing either linkage or independent inheritance. In so far as you have data ready for publication, I prefer to receive a copy of your manuscript, but shall be glad to have also records which you are not ready to publish, if you care to send them. I agree not to publish any such data without your consent and in any case to give proper credit. Any records sent, however, should be with the understanding that I am at liberty to use them in an early revision of the mimeographed sheets for distribution to other workers, pending the publication of the general linkage paper which I have been threatening to bring out for some years now.

I indicated at New York that the records were too incomplete to warrant publication now, a fact made strikingly obvious by the "rainbows" on the maps. The distribution of the data in mimeographed form should serve temporarily the needs of those actively studying maize genetics; and others can wait. The coordination of effort agreed to in New York should go far toward straightening out many of the question marks in the next year or two.

In this connection, I add here, as a reminder, a list of those to whom linkage groups were parcelled out at New York.

- C-Wx group - Eyster, Bucknell; Beadle, Cornell.
 R-G group - Lindstrom, Jenkins, Wentz, Ames.
 Su-Tu group - Emerson, Cornell.
 B-Lg group - Stadler, Missouri; McClintock, Cornell.
 Y-Pl group - Hill, Cornell.
 P-Br group - Emerson, Cornell.
 Ra-Gl₁ group - Brewbaker, Minnesota; Brunson, Manhattan; Fraser, Cornell.
 Pr-V₂ group - Eyster, Bucknell; Jorgenson, Ohio; Li, Cornell.
 D₁-Pg₂ group - Not assigned.
 A-Ts₄ group - Brink, Wisconsin; Li, Cornell.

To those not at the New York meeting, it should be explained that this assignment of groups was, so far as possible, made in accordance with the expressed interests of those assuming the responsibilities entailed. It was far from our purpose to preempt groups for ourselves and thereby warn off other workers. Our purpose rather was to make sure that each known group would be given immediate and adequate attention to the end that the not very exciting job of chromosome mapping may go forward with some dispatch, thereby making possible an attack on certain important genetic problems now awaiting just such tools as accurate linkage maps afford. It should go without saying therefore that the help of those of you who were not at the New York conference will be welcomed.

I suggest that those who have made themselves responsible for any group, request needed material directly from the workers most likely to have it, as indicated by the names in the last column of the table for that group. We at Cornell shall be glad to furnish on request tester stocks in so far as our somewhat limited supply will permit. It would doubtless be helpful if those who have particularly desirable testers for any group would proffer them to the ones who are primarily responsible for that group.

Sincerely,

R. A. Emerson

GENERAL NOTES

Linkage data.-

In the last column of the tables giving the linkage data for the several linkage groups, papers from which the records have been summarized are indicated by author and year. Not all published data are included. For instance, F_2 data are omitted when abundant back-cross data are available. Records credited to an author without indication of the year are unpublished. In general, unpublished data received in personal correspondence are not included, except when no published records are available. Such data are doubtless incomplete. It is thought, therefore, that workers will prefer to add their complete data as of the spring of 1929.

X and Y in the column headings of the several tables indicate the dominant genes of the first column and x and y their respective recessive allelomorphs.

In the second column under the heading "Link. phase", C = coupling and R = repulsion, Bc = back-crossed and S = selfed.

Data presented in the table of three-point tests are included in, not additional to, data in the several group tables. The first column of this table shows the genotype of one parent only, the other parent having obviously the respective allelomorphs of the genes of parent no. 1. The genotypes involved in columns 2 - 5 will be clear from the following illustration:

Parent No. 1	Parental combinations		Region 1	Region 2	Regions 1 and 2
	No. 1	No. 2			
C sh Wx	C sh Wx-c	Sh wx	C Sh wx-c	sh Wx	C Sh Wx-c
I Sh wx	I Sh wx-i	sh Wx	I sh Wx-i	Sh wx	I sh wx-i

Maps.-

No attempt has been made to indicate map distance other than by observed cross-over percentages; 3 mm. = 1 per cent crossing over.

Starred genes (*) are those located with reasonable certainty; others probably belong in the general region indicated.

A gene tested with only one of the located genes is placed opposite that gene at a distance determined by the cross-over percentage, its locus being approximately at one end or other of the "rainbow".

Independence of linkage groups.-

This chart shows what tests have been made between genes of any one linkage group and those of other, presumably independent, groups. Thus, there are records involving approximately 9900 individuals from selfed parents indicating independence between C or I and A and approximately 2000 individuals in back-cross progenies indicating independence of sh and A. It is obvious that the data are not adequate to establish the independence of all the groups, and it is hoped that other workers will have unpublished data to fill in some of the "holes". As an example of the necessity of obtaining more nearly adequate data, a manuscript by Hayes and Brubaker (received after the stencils for the linkage tables had been cut) indicates that gl_2-fl belong to the B-Ig group, while Beadle's unpublished records suggest that fl is in the C-wx group. The independence of these two groups is, therefore, questionable.

C-SH-WX GROUP

List of Genes

ar	Argentia - finely striped leaf	Eyster 1929
au ₁	Aurea chlorophyll-yellow plant	Eyster 1929
au ₂	Aurea chlorophyll-yellow seedling	Eyster 1929
kp	Brown pericarp with a	Meyers 1927
C	Colored aleurone with A and R	East and Hayes 1911
d ₃	Dwarf plant	Suttle (Unpub.)
de ₁₅	Defective endosperm	Brink 1927
fl	Floury endosperm	Hayes and East 1915
gl₂	Glossy seedling	Hayes and Brewbaker 1925
gm ₁	Germless	Eyster 1929
I	Inhibitor for aleurone color	East and Hayes 1911
pk	Polkadot leaf	Eyster 1929
v ₁	Virescent seedling	Demerec 1924
v ₁₄	Virescent seedling	Phipps (Unpub.)
v ₁₅	Virescent seedling	Phipps (Unpub.)
w ₁₁	White seedling	Demerec 1926
wx	Waxy endosperm	Collins 1909
yg	Yellow-green plant	Jenkins 1927

Notes

- pk The 1929 data of Eyster on pk are not consistent with his earlier data. He makes the statement in his 1929 paper that pk and ar show relatively close linkage--hence pk probably lies on the wx side of C.
- d₃
w₁₁ In the material on which the d₃ and w₁₁ counts were made, the C and R factors were segregating. Demerec states that a calculation of the recombination percentage with C would suggest that both d₃ and w₁₁ were on the wx side of sh but that a calculation on such material could not be depended on.
- au₁ The location of au₁ to the right of sh is somewhat doubtful. Recombination values with C and sh are based on separate progenies. Neither au₁ or au₂ have been tested with yg for allelomorphism.
- v₁₄ V₁₄ is known to be located in the C-sh-wx linkage group but the data (Phipps unpub.) are of such a nature that a recombination value cannot be calculated.

Linkage Data

Genes X Y	Link. phase	Number of individuals					Recombina- tions		Authority
		XY	Xy	xY	xy	Total	No.	%	
C Wx	R Bc	115	340	298	92	845	207	24.5	Breggar '13
	C Bc	858	310	311	781	2260	621	27.5	Breggar '13
	C Bc	371	115	125	397	1008	240	23.8	Kempton '19
	C Bc	2542	717	739	2710	6708	1456	21.7	Hutchison '22
						42511	9388	22.1	Stadler '25
					249663	67402	27.0	Collins and (Kempton '27	
					302995	79314	26.2		
C Sh	C Bc	4032	149	152	4035	8368	301	3.6	Hutchison '22
	C Bc	10077	366	397	9866	20706	763	3.7	Eyster '29
	R Bc	638	21379	21096	672	43735	1310	3.0	Hutchison '22
					72849	2374	3.3		
Sh Wx	R Bc	1531	5991	5885	1488	14895	3019	20.3	Hutchison '22
X Sh	C Bc	9452	384	402	9377	19615	786	4.0	Hutchison '22
I Wx	C Bc	1487	584	547	1520	4138	1131	27.3	Hutchison '22
	R Bc	790	2217	2283	792	6082	1582	26.0	Hutchison '22
						10220	2713	26.6	
C V ₁	R Bc	300	676	711	294	1981	594	30.0	Demerec '24
Wx V ₁	R	70	84	40	3	197		7.2	Demerec '24
J Pk	C S ¹	128	6	54	56	244		2	Eyster '24
	C S ²	148	5	128	92	373		2	Eyster '24
Sh Pk	R S ³	140	61	60	2	263		10	Eyster '24
	R S ³	382	173	173	11	739		24.5	Eyster '29
	R Bc ³	73	363	366	70	872	143	16.4	Eyster '29
Sh D ₃	R S	329	162	138	8	637		22.8	Demerec '26
Sh W ₁₁	R S	487	193	161	16	857		31.2	Demerec '26
	C S	320	26	25	67	438		13.4	Demerec '26
J Yg	C S					30817		20.5	Jenkins '27
	R S					3885		23.0	Jenkins '27
	R Bc	10	57	52	7	126	17	13.5	Jenkins '27
Sh Yg	R Bc	193	546	429	99	1267	292	23.0	Jenkins '27
	R S	2583	1212	1057	89	4941		28.6	Jenkins '27
Wx Yg	C Bc	397	289	297	412	1395	586	42.0	Jenkins '27
	R Bc	78	120	136	80	414	158	38.2	Jenkins '27
						1809	744	41.1	Jenkins '27
De ₁₅ Wx	C S	4075	461		1609	6145		19.4	Brink '27
De ₁₅ Sh	R S	2449	1146		1237	4832		16.5	Brink '27
Wx Bp	R Bc	9	56	49	9	123	18	14.6	Meyers '27
C Ar	R Bc ⁴	2178	4692	4166	1507	12543	3685	29.4	Eyster '29
Sh Ar	R Bc ⁴	1925	4763	4177	1221	12086	3146	26.0	Eyster '29
Sh Au ₁	C S	2108	311	310	492	3221		21.6	Eyster '29
C Au ₁	C S ¹	546	79	638	305	1568		26.5	Eyster '29
Sh Au ₂	R S ⁵	340	133	146	10	629		28.0 ⁵	Eyster '29
Sh Gm ₁	C S	2693	301	258	702	3954		15.3	Eyster '29
Wx V ₁₅	R S ³	297	128	139	2	566		19	Phipps
Sh V ₁₅	R S ³	366	171	172	5	714		20	Phipps

¹ C and R segregating - 9:7 ratio

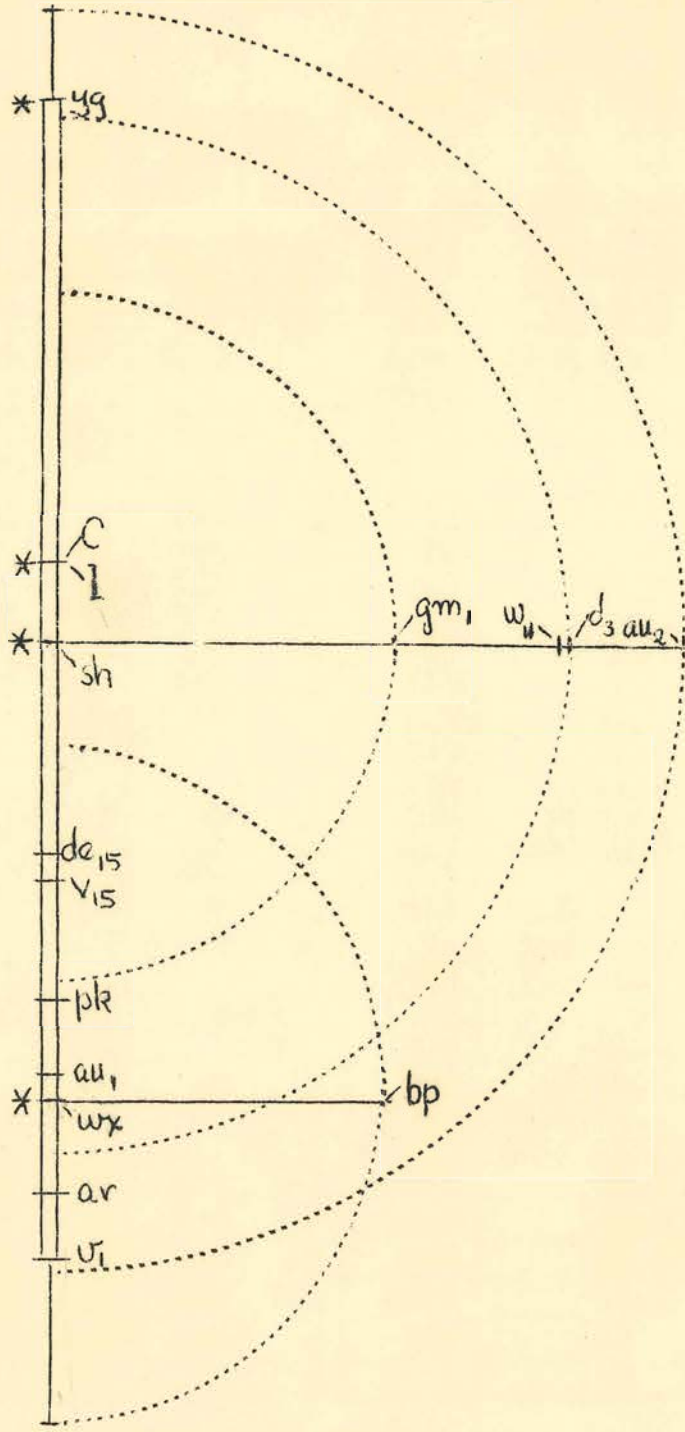
² A, C and R segregating - 27:37 ratio

³ Ratio corrected for germination by author

⁴ See Three-point test data

⁵ Recombination value recalculated - author's calculation given as 39.7

C sh wx Group



R-G GROUP

List of Genes

df	Flint defective	Lindstrom 1925
g	Golden plant	Lindstrom 1918
gm ₂	Germless	Demerec 1926
li ₁	Lineate - striped leaves	Kempton 1920
l ₁	Luteus seedlings	Lindstrom 1917
l ₂	Luteus seedlings	Lindstrom 1925
nl	Narrow-leaf	Emerson (Unpub.)
pg ₁	Pale-green seedling	Brunson 1924
R	Aleurone color	East and Hayes 1921
S	Spotted aleurone with Rrr	Kempton 1919
v ₁₈	Virescent seedling	Phipps (Unpub.)
v ₂₀	Virescent seedling	Phipps (Unpub.)
w ₂	White seedling	Carver 1924

Linkage Data

Genes X Y	Link. phase	Number of individuals				Recombina- tions		Total	Authority	
		X Y	X y	x Y	x y	No.	%			
R G	C Bc	200	55	58	174	487	113	23.2	Lindstrom '17 & '18	
	C Bc	227	36	33	195	491	69	14.1	Emerson	
	R Bc	29	81	86	18	214	47	22.0	Lindstrom '18	
	R Bc	18	117	156	28	319	46	14.4	Emerson	
						1511	275	18.2		
R L ₁	C S	303	2	5	121	431		1.6	Lindstrom '21	
G L ₁	R Bc	8	35	21	5	69	13	18.8	Lindstrom '18	
R Pg ₁	C S ²	1907	300	1053	686	3946		23.3	Brunson '24	
	R S	1199	506	445	32	2182		27.2	Brunson '24	
G Pg ₁	C S	628	59	57	146	890		14.6	Wentz	
Li ₁ Pg ₁	R S	194	71			265		45	Brunson '24	
R W ₂	C S	1329	171	202	402	2104		18.5	Carver '24	
	C S	648	74	81	157	960		17.8	Lindstrom '24	
	R S	43	16	22	2	83		30.8	Carver '24	
W ₂ L ₁	R S	815		210	10	1035			Lindstrom '25	
	R S ³	585		348	84	1017			Lindstrom '25	
	R S ⁴	560		318	70	948			Lindstrom '25	
	R S ⁵	380		402	115	897		22.0	Lindstrom '25	
R L ₂	R S	986	405	473	69	1893		33.9	Lindstrom '25	
	C S ²	837	197	582	277	1893		35.4	Lindstrom '25	
R Gm ₂	R S	2239	784	976	84	4083		31	Demerec '26	
	R S	6876	2947	1182	90	11095		27	Wentz	
Gm ₂ G	R S	2810	873			3683		50 ±	Wentz	
Gm ₂ Pg ₁	R S	835	255			1090		50 ±	Wentz	
R V ₁₈	C ⁶	51	15	43	93	202		20	Phipps	
R V ₂₀	C Bc ⁶	77	10	80	152	319		12.5	Phipps	
G Li ₁	R Bc	148	817	924	111	2000	259	13.0	Hutchison	
R Li ₁	C Bc	208	74	86	138	506	160	31.6	Hutchison	
	C Bc ⁶	460	191	282	374	651 ⁷	191	29.3	Hutchison	
						1157	351	30.3		
G NI	R Bc	69	389	382	49	889	118	13.3	Emerson	
R NI	C Bc	219	93	116	191	619	209	33.8	Emerson	

¹1918 data indicate complete linkage

²C and R segregating - 9:7 alleurone ratio

³W₁ and w₂ segregating

⁴W₂ and w₃ segregating

⁵W₁, w₂ and w₃ segregating

⁶C and R segregating

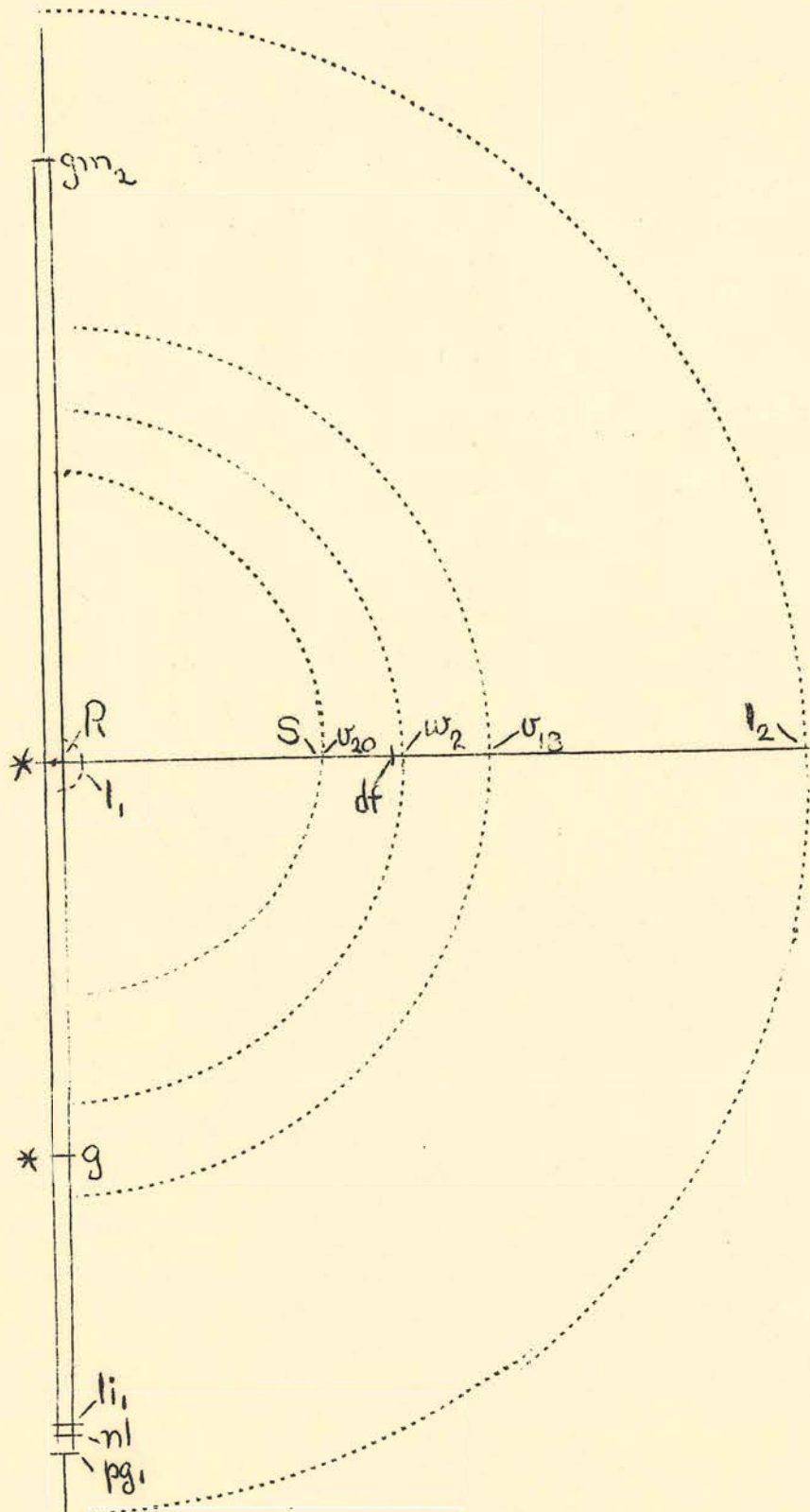
⁷First two classes only

Notes

df Lindstrom states that df and w₃ are very closely linked but presents no data.

S Kempton (1919) postulated this spotting factor, located so as to give about 12.5% recombinations with R. Emerson (Unpub.) has additional evidence in support of this assumption.

R g Group



SU-TU GROUP

List of Genes

de ₁	Defective endosperm	Mangelsdorf/1925 ⁵
de ₆	Defective endosperm	Mangelsdorf 1926
de ₁₆	Defective endosperm	Wentz 1925 + Jones
Ga	Gamete - pollen tube growth	Mangelsdorf/1925
ge ₁	Premature germination	Mangelsdorf 1926
su	Sugary endosperm	East and Hayes 1911
Ts ₅	Tassel-seed	Emerson (Unpub.)
Tu	Tunicate ear	Collins 1917
wl	White-base leaf	Stroman 1925

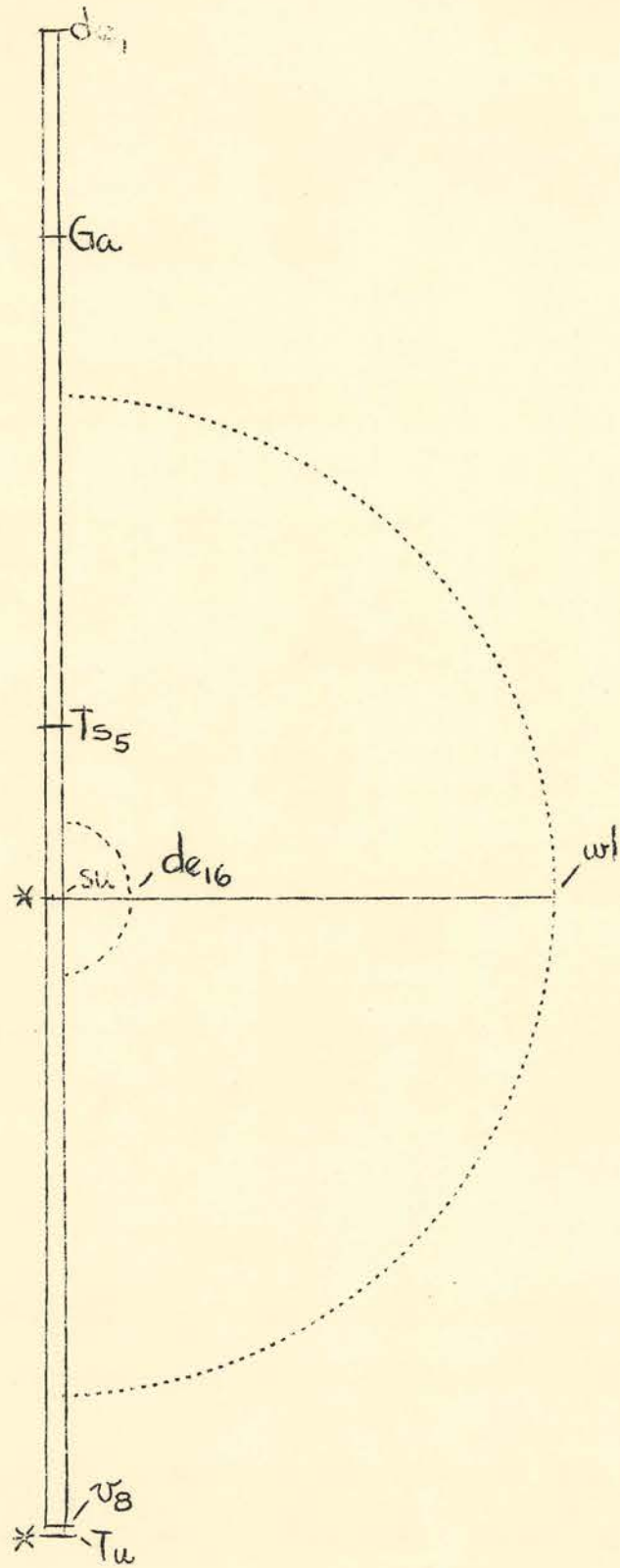
Linkage Data

Genes X Y	Link. phase	Number of individuals						Recombina- tions		Authority
		X Y	X y	x Y	x y	Total	No.	%		
Su Tu	C S	113	4	7	25	149	-	8.3	Jones & Gallastegui '19	
	C Bc	430	175	169	406	1180	344	29.1	Eyster '21	
	C Bc	612	290	208	562	1672	498	29.8	Emerson	
	R Bc	1031	2498	2093	807	6429	1838	28.6	Eyster '22	
	R Bc	63	215	164	57	499	120	24.0	Emerson	
						9780	2800	28.6		
Su Wl	R S	44	19	11	1	75		25.0	Stroman '24	
	R S	4492	2018	1961	93	8564		22.0	Carver '27	
De ₁₆ Su	C S	20622	453		7201	28276	-	3.2	Wentz '25	
Su V ₈	C S	940	214	179	148	1481	-	32.4	Demerec '26	
V ₈ Tu	C S	450	1	Lethal		451		<1	Phipps	
De ₁ Su	R S	601	238	247	64	1150		39	Mangelsdorf & Jones '25	
De ₆ Su	R S	204	92			296		26	Mangelsdorf '25	
Ge ₁ Su	R S	1218	474			1692		40	Mangelsdorf '26	
Su Ts ₅	C Bc	578	41	42	457	1118	83	7.4	Emerson	
Ts ₅ Tu	R Bc	49	166	115	48	378	97	25.7	Emerson	

Notes

- de₁₆ is used instead of de_{su} for sugary defective of Wentz.
- V₈ is very near Tu but whether to the left or right is unknown.
- Ga is to the left of su because it disturbs the Tu-tu ratio very little if at all in pedigrees in which it disturbs the Su-su ratio materially (Emerson, Unpub.).
- de₁ is presumably to the left of Ga, because Ga is between de and su (Mangelsdorf and Jones 1925).

su Tu Group



B-LG GROUP

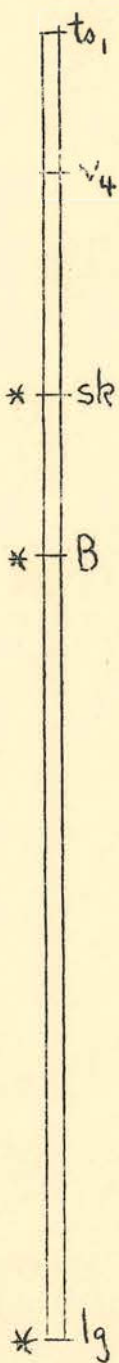
List of Genes

B	Intensifier of plant color	Emerson 1918
lg	Liguleless	Emerson 1918
sk	Silkless	Jones 1925
ts ₁	Tassel-seed	Emerson 1920
v ₄	Virescent seedling	Demerec 1924

Linkage Data

Genes X Y	Link. phase	Number of individuals					Recombina- tions		Authority
		X Y	X y	x Y	x y	Total	No.	%	
B lg	C Bc	240	134	102	243	719	236	32.8	Emerson '18
	C Bc	642	291	282	620	1835	573	31.2	Emerson
	C Bc	2487	1469	1557	2609	8122	3026	37.2	Emerson & Hutchison '21
	R Bc	498	1085	1037	504	3124	1002	32.1	Emerson
						13800	4837	35.0	
Lg Ts ₁	C Bc	117	52	72	74	315	124	39.4	Emerson
	R Bc	51	65	64	42	222	93	41.9	Emerson
						537	217	40.4	
B V ₄	C Bc	113	24	21	110	268	45	16.8	Demerec '24
V ₄ Lg	R Bc	412	501	521	366	1800	778	43.2	Demerec '24
B Sk	C Bc	1332	97	106	1226	2761	203	7.4	Anderson
	R Bc	2	82	66	6	156	8	5.1	Anderson
						2917	211	7.2	
Lg Sk	R Bc	187	288	315	167	957	354	37.0	Anderson
		148	60	67	133	408	127	31.1	Anderson
						1365	481	35.2	

B lg Group



Y-PL GROUP

List of Genes

Bh	Blotched aleurone with A c R	Emerson (Unpub.)
fi	Fine streaked leaves	Anderson 1922
Pl	Purple plant color	Emerson 1918
sm	Salmon silks	Anderson 1921
v ₆	Virescent seedling	Carver 1927
v ₇	Virescent seedling	Carver 1927
w ₁	White seedling	Stroman 1924
w ₅	White seedling with w ₆	Demerec 1924
w ₆	White seedling with w ₅	Demerec 1924

Linkage Data

Genes		Link. phase	Number of individuals				Total	Recombinations		Authority
X	Y		X Y	X y	x Y	x y		No.	%	
Y	Pl	C Bc	79	22	28	71	200	50	25.0	Emerson '18
			545	221	234	506	1506	455	30.2	Anderson '21
		R Bc	20	51	30	55	216	81	37.5	Anderson
			173	46	59	176	454	105	23.1	Hutchison
			367	880	897	372	2516	739	29.4	Anderson '21
		135	398	374	118	1025	253	24.7	Anderson	
						5917	1683	28.5		
Pl	Sm	C Bc	1076	145	146	994	2361	291	12.3	Anderson '21
			84	1014	971	76	2145	160	7.5	Anderson '21
						4506	451	10.0		
Y	Fi	C Bc	353	0		many				Anderson '23
Y	w ₅	C S	250	37	35	54	376		24.3	Demerec '23
Y	{w ₅ w ₆ }	S ¹	349	12	60	33	454		{24.3	Demerec '23
									{24.5	
Y	w ₁	C S	1020	237	259	191	1707		35	Lindstrom '24
		C S	1132	321	347	175	1975		42	Stroman '24
		R S	456	181	186	41	864		42	Stroman '24
Y	v ₆	R S	467	225	209	12	913		23	Carver '27
Y	v ₇	C S	592	149	178	79	998		42	Carver '27
		C S ²	445	277	106	116	944		36	Carver '27
v ₆	v ₇	R S	497	175	237		913		42	Carver '27
Bh	Y	C Bc	144	51	118	210 ³	523	169	32.3	Anderson
Bh	Pl	C Bc	58	1	26	47 ³	132		1.7 ⁴	Anderson

¹w₅ and w₆ duplicate genes

²Segregating for another v - not linked

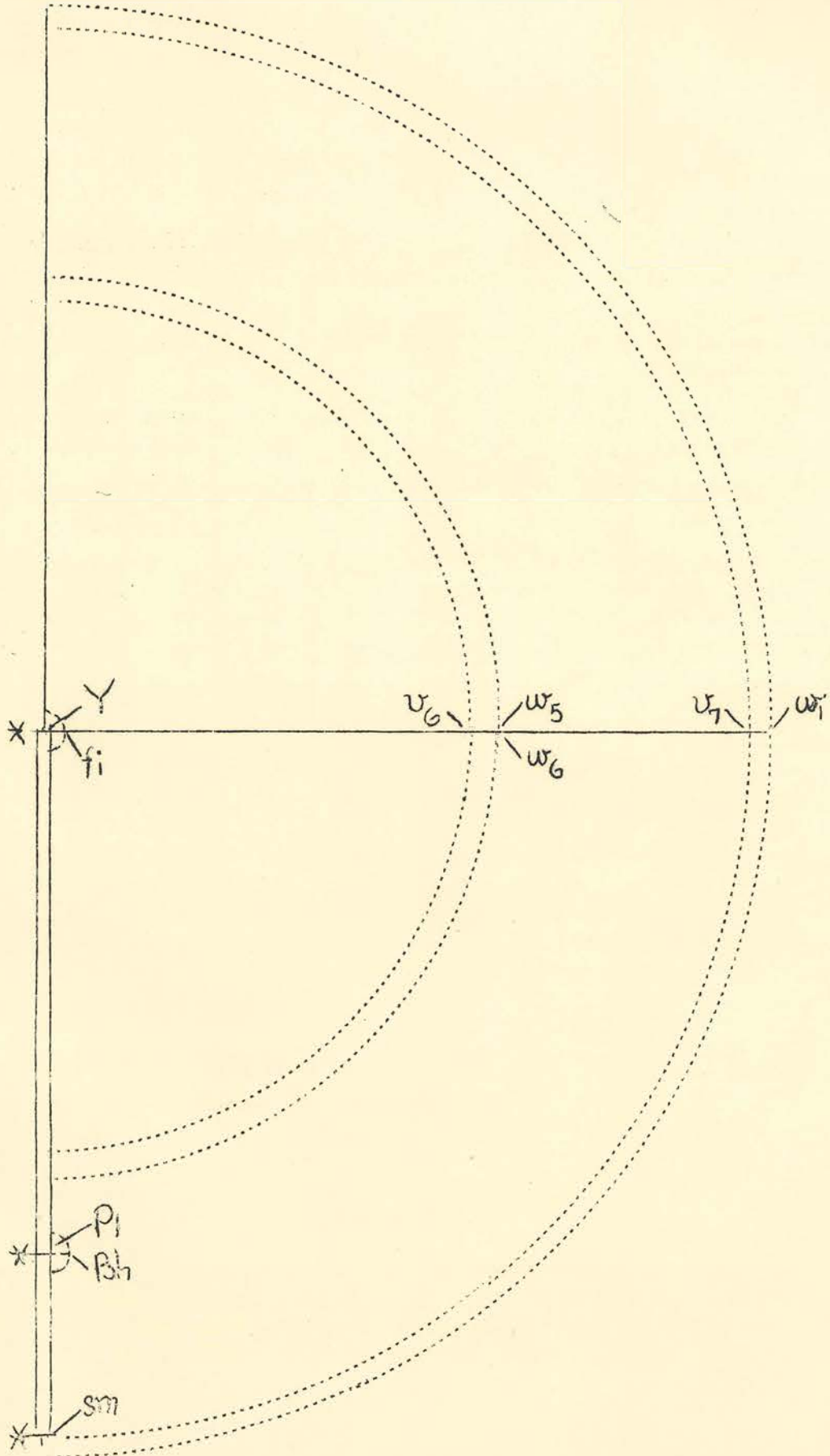
³Probably part of this class actually Bh

⁴From Bh class

Notes

m₁ }
m₂ } Stroman presents data which he interprets as showing linkage between m₁ and m₂ and also between m₁ and Y. His data are sufficiently extensive only to suggest that these factors may belong to this linkage group.

Y Pl Group



RA-GL₁ GROUP

List of Genes

Bn	Brown aleurone	Kvakan 1924
gl ₁	Glossy seedling	Kvakan 1924
in	Intensifier of aleurone	Fraser 1924
pg ₁	Pale-green seedling	Demerec 1925
ra	Ramosa	Gernert 1912
sl	Slashed seedling	Hayes and Brewbaker 1928
sr ₂	Striate - striped leaf	Brunson (Unpub.)

Linkage Data

Genes		Link. phase	Number of individuals				Total	Recombina- tions		Authority
X	Y		X Y	X y	x Y	x y		No.	%	
Bn	Gl ₁	C Bc	177	63	54	192	486	117	24.1	Kvakan '24
Gl ₁	V ₅	C Bc	106	9	6	120	241	15	6.2	Kvakan '24
Bn	V ₅	C Bc	83	31	29	98	241	60	24.9	Kvakan '24
Bn	Ra	C Bc	169	104	100	161	534	204	38.2	Kvakan '24
Bn	Pg ₁	C S	203	8	5	65	281		4.5	Demerec '25
Gl ₁	Sr ₂	R Bc	97	289	342	63	791	160	20.2	Brunson

Notes

sl Hayes and Brewbaker state that sl belongs to this linkage group.

Y₂ } Hayes and Brewbaker present data showing a linkage
Y_p } between two factors for yellow endosperm (Y₂ and Y_p)
and a glossy seedling factor. Since the relation of
the glossy character to gl₁ is not evident, the placing
of these two genes in this linkage group would appear
uncertain.

RA-GL₁ GROUP

List of Genes

Bn	Brown aleurone	Kvakan 1924
gl ₁	Glossy seedling	Kvakan 1924
in	Intensifier of aleurone	Fraser 1924
pg ₁	Pale-green seedling	Demerec 1925
ra	Ramosa	Gernert 1912
sl	Slashed seedling	Hayes and Brewbaker 1928
sr ₂	Striate - striped leaf	Brunson (Unpub.)
v ₅	Virescent	Demerec 1924

dup. - skip 2

Linkage Data

Genes		Link. phase	Number of individuals				Total	Recombina-tions		Authority
X	Y		X Y	X y	x Y	x y		No.	%	
Bn	gl ₁	C Bc	177	63	54	192	486	117	24.1	Kvakan '24
GL ₁	V ₅	C Bc	106	9	6	120	241	15	6.2	Kvakan '24
Bn	V ₅	C Bc	83	31	29	98	241	60	24.9	Kvakan '24
Bn	Ra	C Bc	169	104	100	161	534	204	38.2	Kvakan '24
Bn	pg ₁	C S	203	8	5	65	281		4.5	Demerec '25
GL ₁	Sr ₂	R Bc	97	289	342	63	791	160	20.2	Brunson

Notes

sl Hayes and Brewbaker state that sl belongs to this linkage group.

Y₂ } Hayes and Brewbaker present data showing a linkage
Y_p } between two factors for yellow endosperm (Y₂ and Y_p)
and a glossy seedling factor. Since the relation of
the glossy character to gl₁ is not evident, the placing
of these two genes in this linkage group would appear
uncertain.

RA-GL₁ GROUP

List of Genes

Bn	Brown aleurone	Kvakan 1924
gl ₁	Glossy seedling	Kvakan 1924
in	Intensifier of aleurone	Fraser 1924
pg ₁	Pale-green seedling	Demerec 1925
ra	Ramosa	Gernert 1912
sl	Slashed seedling	Hayes and Brewbaker 1928
sr ₂	Striate - striped leaf	Brunson (Unpub.)
v ₅	Virescent	Demerec 1924

Linkage Data

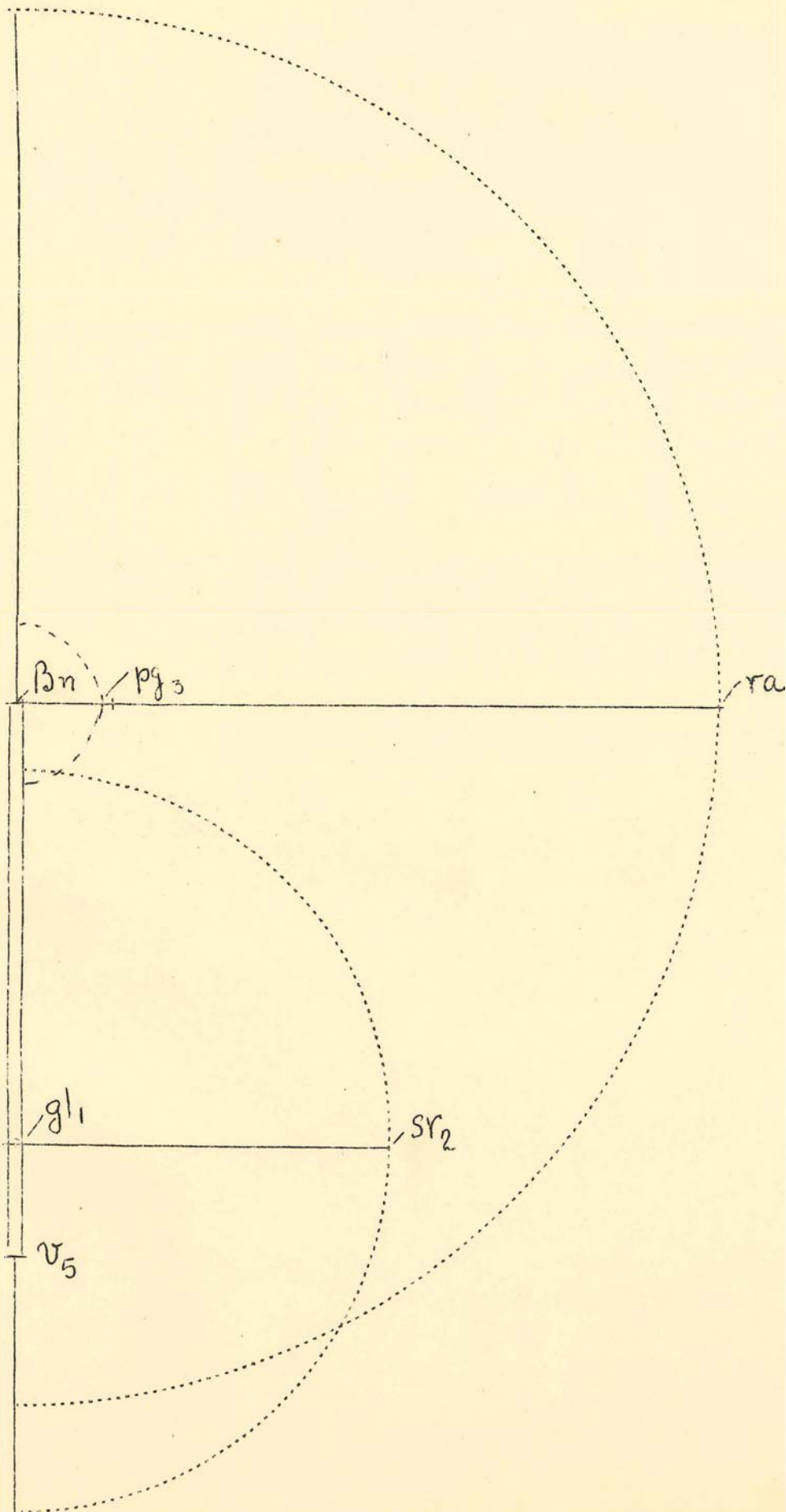
Genes X Y	Link. phase	Number of individuals				Total	Recombina- tions		Authority
		X Y	X y	x Y	x y		No.	%	
Bn GL ₁	C Bc	177	63	54	192	486	117	24.1	Kvakan '24
GL ₁ V ₅	C Bc	106	9	6	120	241	15	6.2	Kvakan '24
Bn V ₅	C Bc	83	31	29	98	241	60	24.9	Kvakan '24
Bn Ra	C Bc	169	104	100	161	534	204	38.2	Kvakan '24
Bn Pg ₁	C S	203	8	5	65	281		4.5	Demerec '25
GL ₁ Sr ₂	R Bc	97	289	342	63	791	160	20.2	Brunson

Notes

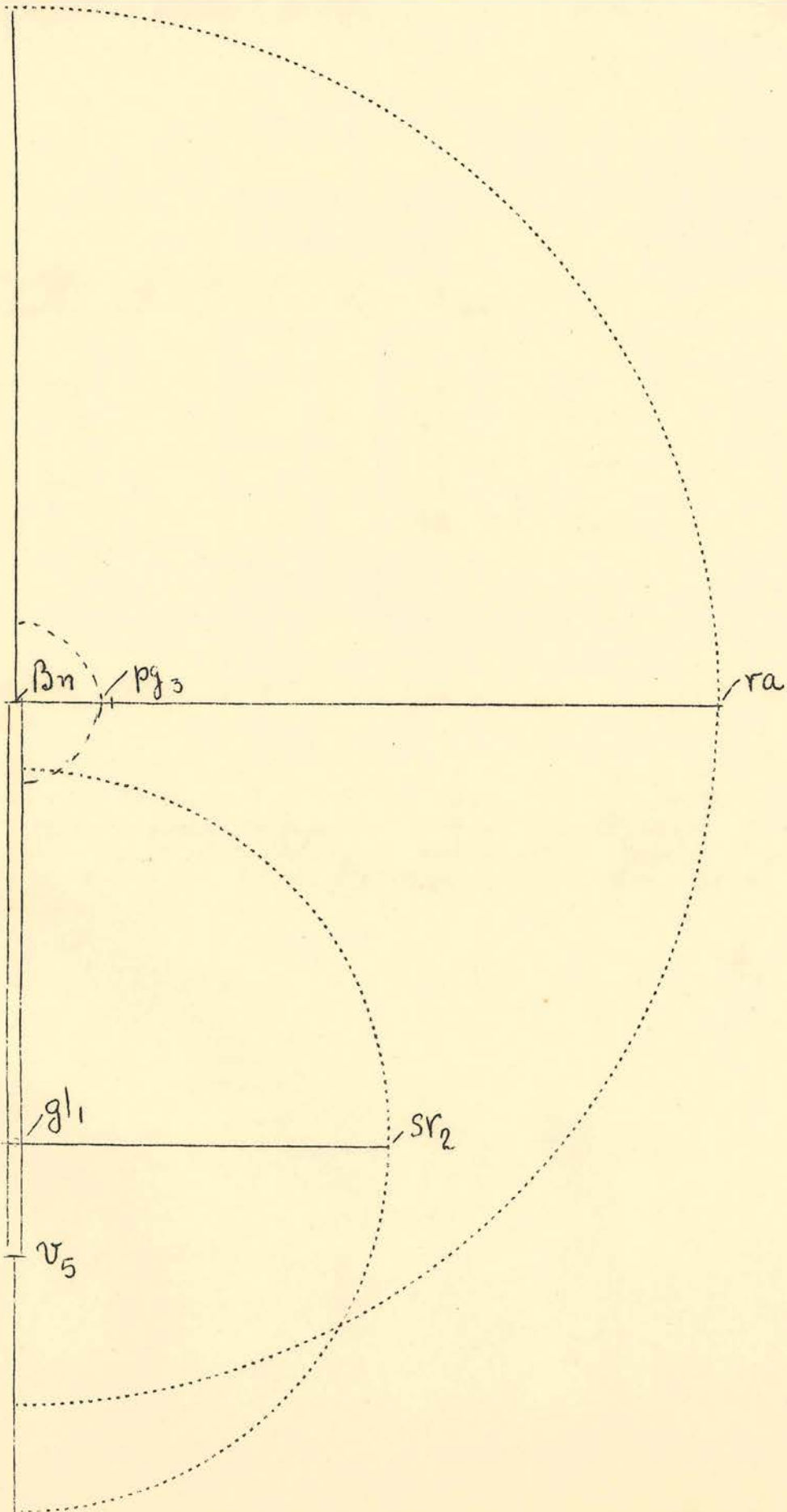
sl Hayes and Brewbaker state that sl belongs to this linkage group.

Y₂ } Hayes and Brewbaker present data showing a linkage
Yp } between two factors for yellow endosperm (Y₂ and Yp)
and a glossy seedling factor. Since the relation of
the glossy character to gl₁ is not evident, the placing
of these two genes in this linkage group would appear
uncertain.

ra-gl₁ Group



ra-gl₁ Group



PR-V₂ GROUP

List of Genes

bm	Brown midrib	Eyster 1926
bv	Brevis - semi-dwarf plant	Suttle (Unpub.)
f ₂	Fine striped leaves	Eyster 1926
Pr	Purple aleurone	East and Hayes 1917
sc ₁	Scarred endosperm	Eyster 1926
tn	Tiny plant	Eyster 1926
v ₂	Virescent seedling	Demerec 1924
v ₃	Virescent seedling	Demerec 1924
v ₁₂	Virescent seedling	Phipps (Unpub.)
yg	Yellow green	Eyster 1926
ys	Yellow-stripe	Beadle 1929

Linkage Data

Genes X Y	Link. phase	Number of individuals				Total	Recombina- tions		Authority
		X Y	X y	x Y	x y		No.	%	
Pr V ₂	R Bc	377	532	499	366	1774	743	41.9	Phipps
	C Bc	67	46	41	51	205	87	42.4	
						1979	830	42.0	
Pr V ₃	R Bc	123	296	320	102	841	225	26.8	Phipps
Pr V ₁₂	C ₁ Bc	61	15	4	75	155	19	12.4	Phipps
	C S	492	37	39	137	705		11.4	
Pr Ys	R	219	323	209	19	770		8.3	Beadle '29

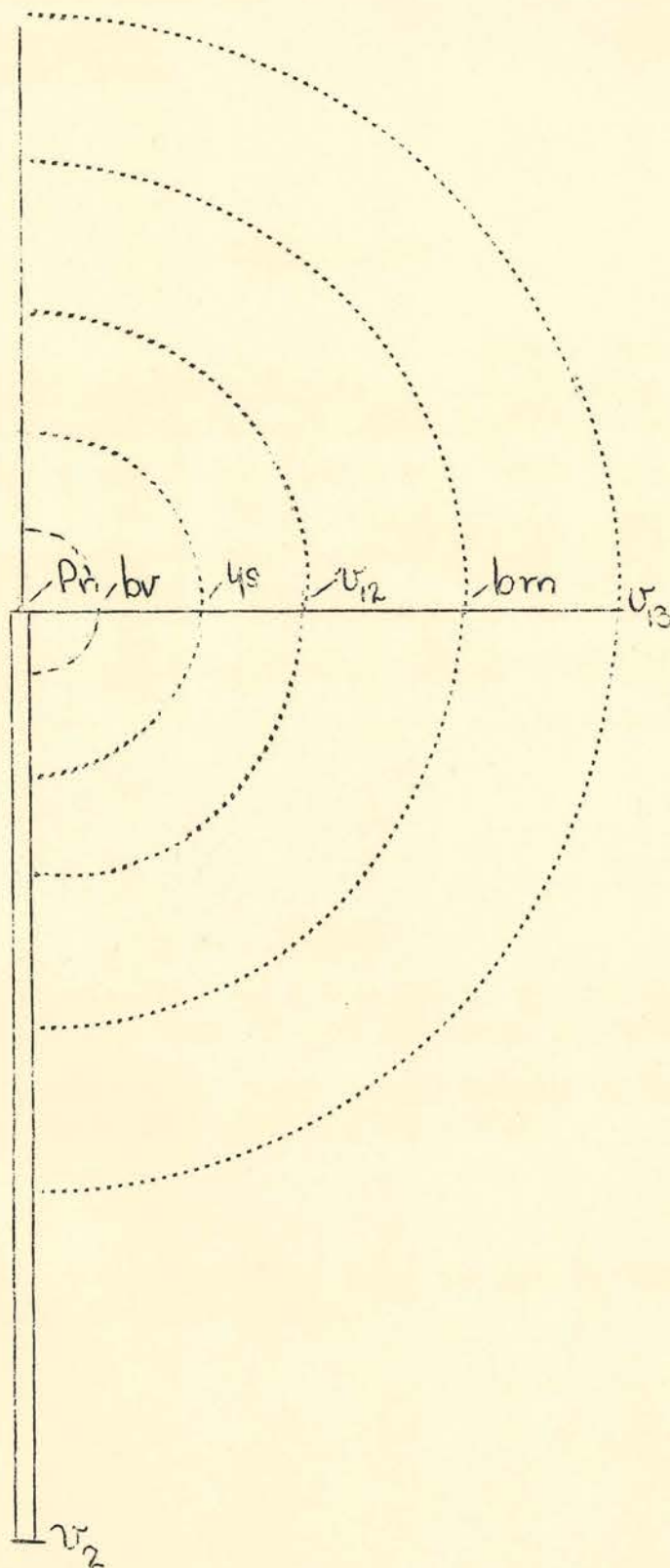
Notes

bm Eyster states that bm shows about 20 per cent recombinations with Pr but presents no data.

f₂ } Eyster states that these genes belong to the Pr
 sc₁ } linkage group but presents no data.
 tn }
 yg }

bv Li (Unpub.) has evidence that bv and Pr are relatively closely linked.

Pr v_2 Group



D₁ - PG₂ GROUP

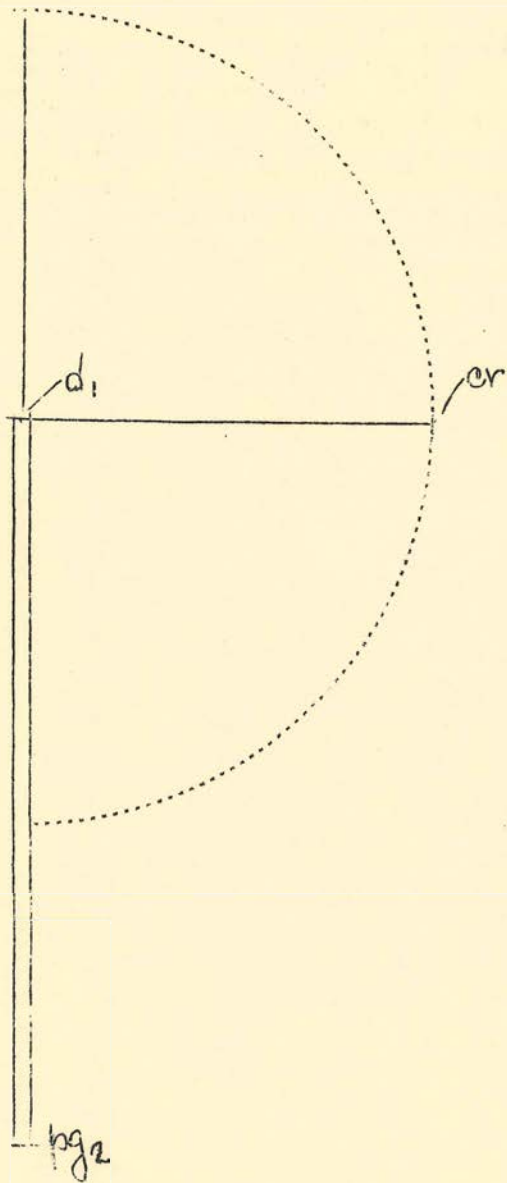
List of Genes

d ₁	Dwarf plant	Emerson 1912
pg ₂	Pale-green seedling	Demerec 1924
cr	Crinkly leaves	Emerson 1921

Linkage Data

Genes X Y	Link. phase	Number of individuals					Recombina- tions		Authority
		X Y	X y	x Y	x y	Total	No.	%	
D ₁ Pg ₂	R S	1364	584	580	65	2593		32	Demerec '24
D ₁ Cr	R Bc	15	53	48	15	131	30	22.9	Emerson
	C Bc	518	102	207	482	1209	209	17.3	Emerson
						1340	239	17.8	

d pg₂ Group



A-Ts₄ GROUP

List of Genes

A	Anthocyanin pigment	Emerson 1918
na	Nana - dwarf plant	Suttle (Unpub.)
ts ₄	Tassel-seed	Phipps 1928

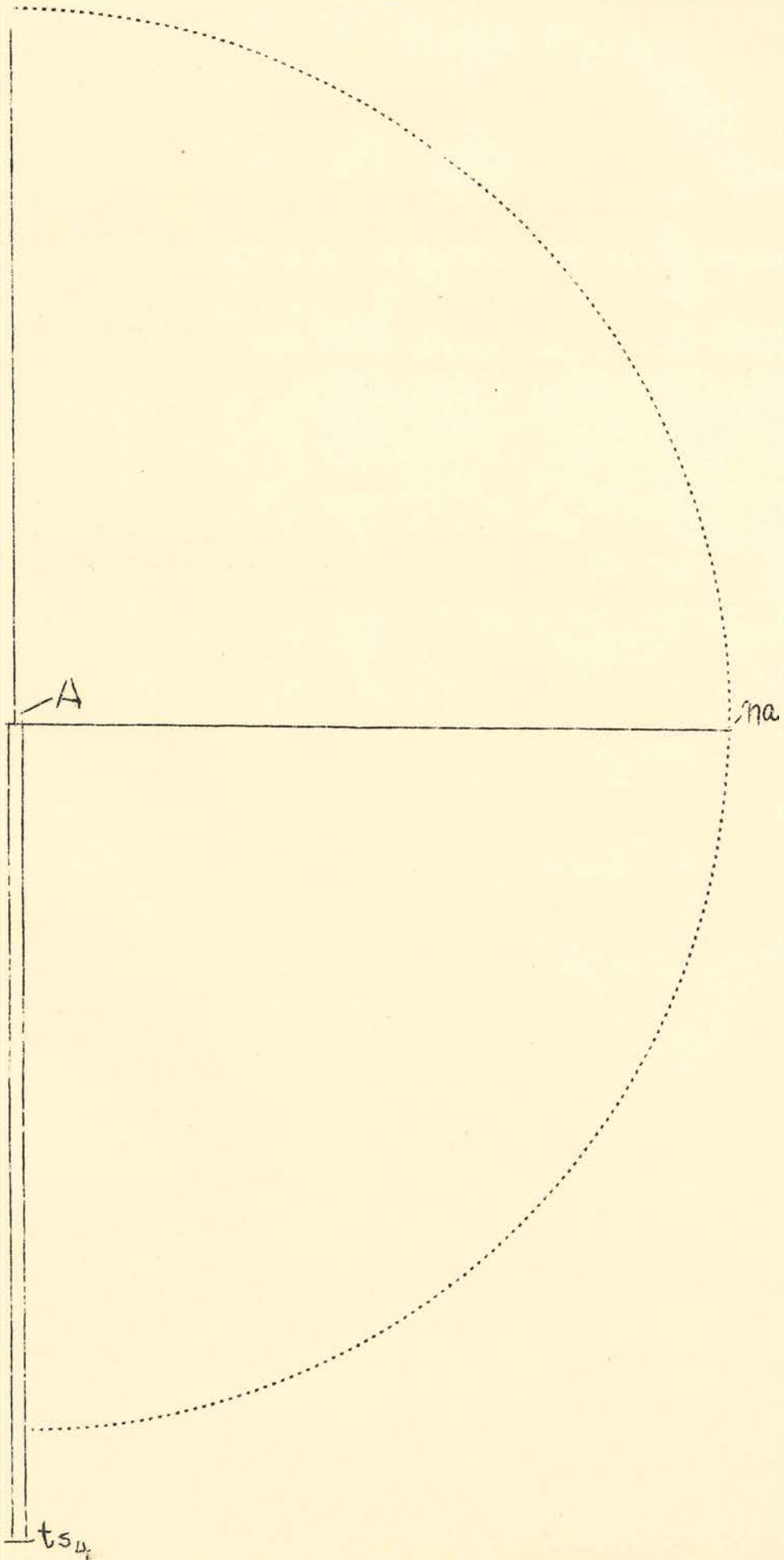
Linkage Data

Genes X Y	Link. phase	Number of individuals					Recombina- tions		Authority
		X Y	X y	x Y	x y	Total	No.	%	
A Ts ₄	C Bc	90	63	70	85	308	133	43.2	Phipps '28
	R Bc	262	351	372	333	1318	595	45.1	Phipps '28
						1626	728	44.8	

Notes

Li (Unpub.) has evidence that na is linked with A, showing about 40 per cent of recombinations.
Jones (Unpub.) also has evidence of this linkage.

A ts_4 Group



SUMMARY OF THREE-POINT LINEAGE TESTS IN MAIZE

Parent No. 1	Parental combina- tions		Recombinations			Total	Coin- ci- dence	Authori- ty
	No.1	No.2	Region 1	Region 2	Regions 1 & 2			
C sh Wx	2538 5246	2708	116 113 229 3.4%	601 626 1227 18.3%	4 2 6 0.12%	6708	0.14	Hutchis '22
I Sh wx	2215 4495	2280	121 139 260 4.3%	669 653 1322 21.7%	2 3 5 0.08%	6082	0.09	Hutchison '22
yg C Sh	54 51 105		7 9 16 12.7%	3 1 4 3.2%	1 0 1 0.8%	126		Jenkins '27
C sh ar	4678 8816	4138	259 192 451 3.6%	1248 1986 3234 25.8%	14 28 42 0.34%	12543	0.33	Eyster '22
Ts ₅ Su tu	163 113 276		9 12 21 5.6%	37 39 76 20.1%	2 3 5 1.3%	378	0.88	Emerson
Ts ₁ b Lg	111 71		24 17	48 35	6 3	315		Emerson
Ts ₁ B lg	57 57 296		20 21 82 15.3%	31 21 135 25.1%	7 8 24 4.5%	222 537	0.77	
Sk B Lg	148 131 279		13 8 21 5.1%	56 52 108 26.3%	0 2 2 0.5%	410	0.36	Anderson
Y Pl Sm	191 180		109 104	21 31	5 5			Anderson
Y pl Sm	436 377		165 206	45 50	5 1			'21
Y Pl sm	305 265		107 124	28 30	0 1			
Y pl sm	333 411 2498		183 152 1150 28.6%	66 59 330 8.2%	16 12 45 1.12%	4023	0.40	
Ts ₂ br F ₁	12 8 20		3 9 12 34.3%	0 3 3 8.6%	0 0 0 0.0%	35		Emerson
Bn Cl ₁ V ₅	83 98 181		22 23 45 18.7%	9 6 15 6.2%	0 0 0 0.0%	241		Kvakon '24

Summary of Data on the Independence of the Linkage Groups in Maize

	yg	CI	sh	wx	v ₁	au ₁	R	G ₁	nl	li ₁	v ₁₈	v ₂₀	Ts ₅	su	Tu	ts ₁	v ₄	sk	B	lg	Y	Pl	sm	w ₁	P	ts ₂	br	f ₁	Bn	gl ₁	v ₅	ra	d ₁	pg ₂	cr	Pr	v ₂	v ₃	v ₁₂	bv			
A ts ₄ na	99	<u>20</u>	<u>5</u>	48			99	<u>9</u>	6				<u>2</u>	<u>30</u>	<u>1</u>		<u>12</u>	<u>13</u>	<u>20</u>	<u>6</u>	<u>11</u>	<u>10</u>		10	<u>15</u>				<u>15</u>						<u>7</u>		<u>7</u>	<u>20</u>	<u>14</u>	<u>3</u>	<u>5</u>		
		<u>2</u>	<u>2</u>				5	<u>5</u>						<u>4</u>	<u>1</u>				<u>4</u>	<u>10</u>	<u>3</u>	<u>4</u>					5	<u>1</u>															
		<u>1</u>	<u>5</u>				<u>20</u>	<u>12</u>						<u>2</u>					<u>5</u>	<u>8</u>	<u>1</u>	<u>5</u>				<u>1</u>				<u>1</u>													
Pr v ₂ v ₃ v ₁₂ bv			<u>8</u>	<u>5</u>			<u>20</u>	<u>6</u>	<u>1</u>					<u>5</u>	<u>3</u>				<u>1</u>	<u>1</u>		<u>2</u>				?			<u>5</u>		<u>13</u>			<u>2</u>		<u>5</u>							
				<u>13</u>				<u>2</u>						<u>3</u>							<u>5</u>	<u>2</u>							<u>8</u>				<u>2</u>										
			<u>8</u>											<u>34</u>							<u>14</u>	<u>2</u>	<u>1</u>										<u>2</u>										
														<u>6</u>							<u>4</u>						4																
d ₁ pg ₂ cr		<u>4</u>			11		<u>5</u>	<u>2</u>						<u>8</u>	<u>5</u>		<u>27</u>		<u>3</u>		<u>3</u>	<u>3</u>				<u>2</u>	<u>5</u>														<u>5</u>		
		<u>7</u>	<u>4</u>	<u>4</u>			<u>7</u>							<u>6</u>							<u>26</u>																						
		<u>4</u>	<u>10</u>	<u>3</u>			<u>12</u>	<u>5</u>	<u>4</u>					<u>9</u>	<u>5</u>	<u>2</u>			<u>15</u>	<u>9</u>	<u>1</u>	<u>6</u>					<u>1</u>															<u>3</u>	
Bn gl ₁ v ₅ ra in		<u>26</u>	<u>14</u>	<u>14</u>	<u>9</u>		<u>6</u>							<u>29</u>	<u>2</u>					<u>5</u>	<u>41</u>	<u>6</u>				<u>3</u>																	
		<u>7</u>	<u>6</u>	<u>4</u>			<u>16</u>	<u>3</u>	<u>3</u>					<u>10</u>	<u>7</u>					<u>11</u>	<u>4</u>	<u>4</u>	<u>0</u>			<u>8</u>																	
		<u>45</u>																																									
P ts ₂ br f ₁		<u>8</u>	<u>3</u>				<u>12</u>	<u>5</u>	<u>6</u>					<u>7</u>	<u>6</u>	<u>1</u>			<u>30</u>	<u>15</u>	<u>18</u>	<u>23</u>	<u>9</u>																				
		<u>1</u>					5	<u>3</u>	<u>2</u>					<u>1</u>					<u>5</u>	<u>3</u>	<u>7</u>	<u>7</u>																					
		<u>1</u>							<u>2</u>						<u>7</u>		15		<u>2</u>	<u>6</u>	<u>1</u>	<u>3</u>																					
Y Pl sm		<u>9</u>	<u>8</u>	<u>12</u>		6	<u>15</u>	<u>4</u>	<u>2</u>	<u>10</u>	1	10		<u>13</u>	<u>1</u>		<u>14</u>		<u>14</u>	<u>10</u>																							
		<u>6</u>	<u>5</u>				<u>12</u>	<u>5</u>	<u>7</u>	<u>8</u>				<u>4</u>	<u>11</u>	<u>3</u>		<u>14</u>		<u>14</u>	<u>10</u>																						
ts ₁ v ₄ sk B lg								5																																			
		<u>12</u>		<u>7</u>			<u>12</u>							<u>8</u>																													
		<u>3</u>					<u>13</u>							<u>4</u>																													
		<u>33</u>	<u>11</u>				<u>46</u>	<u>1</u>	<u>7</u>	<u>7</u>				<u>13</u>	<u>19</u>																												
		<u>11</u>	<u>19</u>			15	<u>55</u>	<u>3</u>	<u>10</u>	<u>7</u>		10		<u>5</u>	<u>21</u>	<u>23</u>																											
Ts ₅ su Tu						2	<u>4</u>																																				
		<u>50</u>					<u>71</u>	<u>5</u>	<u>2</u>	<u>5</u>		3																															
		<u>9</u>					<u>9</u>	<u>5</u>	<u>3</u>																																		
R G ₁ nl li ₁ v ₁₈ v ₂₀		<u>17</u>	<u>23</u>	<u>6</u>																																							
		<u>1</u>	<u>3</u>	<u>3</u>																																							
			<u>3</u>	<u>3</u>																																							
			<u>8</u>	<u>2</u>																																							
			<u>2</u>	<u>2</u>																																							

Figures in table represent approximately the number of hundreds of individuals counted, the counts suggesting independent inheritance. Counts on backcross progenies are distinguished by an underscore from counts from self pollinations.

endence of the Linkage Groups in Maize

1	v ₄	sk	B	lg	Y	Pl	sm	w ₁	P	ts ₂	br	f ₁	Bn	gl ₁	v ₅	ra	d ₁	pg ₂	cr	Pr	v ₂	v ₃	v ₁₂	bv	
<u>22</u>	<u>13</u>	<u>20</u>	<u>€</u>	<u>11</u>	<u>10</u>			<u>10</u>	<u>15</u>				<u>15</u>			<u>7</u>			<u>7</u>		<u>20</u>	<u>14</u>	<u>3</u>	<u>5</u>	<u>5</u>
		<u>4</u>	<u>10</u>	<u>3</u>	<u>4</u>						<u>5</u>	<u>1</u>		<u>1</u>		<u>5</u>				<u>5</u>		<u>3</u>		<u>5</u>	
		<u>5</u>	<u>8</u>	<u>1</u>	<u>5</u>					<u>1</u>		<u>3</u>												<u>5</u>	
		<u>1</u>	<u>1</u>		<u>2</u>				<u>?</u>			<u>5</u>	<u>11</u>			<u>13</u>	<u>2</u>		<u>5</u>						
				<u>5</u>	<u>2</u>							<u>8</u>					<u>1</u>	<u>4</u>							
		<u>14</u>		<u>1</u>	<u>1</u>					<u>4</u>			<u>3</u>												
<u>27</u>		<u>3</u>		<u>3</u>	<u>3</u>				<u>2</u>	<u>5</u>														<u>5</u>	
		<u>15</u>	<u>9</u>	<u>1</u>	<u>6</u>					<u>1</u>		<u>1</u>												<u>3</u>	
			<u>5</u>	<u>41</u>	<u>6</u>				<u>3</u>			<u>7</u>													
		<u>14</u>	<u>4</u>	<u>4</u>	<u>6</u>				<u>8</u>			<u>7</u>													
		<u>30</u>	<u>15</u>	<u>18</u>	<u>23</u>			<u>9</u>																	
		<u>5</u>	<u>3</u>	<u>7</u>	<u>7</u>																				
			<u>16</u>	<u>1</u>	<u>1</u>																				
<u>15</u>		<u>2</u>	<u>6</u>	<u>2</u>	<u>3</u>																				
<u>14</u>		<u>14</u>	<u>10</u>																						
	<u>14</u>	<u>44</u>	<u>19</u>																						
		<u>1</u>																							

Figures in table represent approximately the number of hundreds of individuals counted, the counts suggesting independent inheritance. Counts on backcross progenies are distinguished by an underscore from counts from self pollinations.

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