LINKAGE-STUDIES IN PISUM. III

by

S. J. WELLENSIEK

(Wageningen, Holland 1)) (Received November 2, 1929)

CONTENTS

I. INTRODUCTION

It is extremely difficult to combine into one theory the results of different investigators on linkage-relations in *Pisum.* Two genes, independent according to the one, are found to be linked by another. Much variation in percentages of crossing-over occurs. Several cases of about 50 $\%$ crossing-over — resulting in phenotypic independence **--** have come to our knowledge. The use of different parental lines and lack of uniformity in symbols make it hard to judge whether or not a

Genetica XII 1

Page

¹⁾ Presently: Buitenzorg, Java.

gene used in a certain study is identical with a gene occurring in other material.

No wonder that attempts to combine the data of different authors into one general conclusion fail to yield satisfactory results. Lots more of experimental work must be done, before we can form an opinion on the validity of the *Drosophila* principles in *Pisum.* For the present this work must be done in one group of genes which will leave no doubt with regard to the identity of factors. Furthermore, all possible relations two by two must be studied, since incompleteness does not allow to draw final conclusions.

In "Linkage-Studies I and U" an attempt was made to study the interrelations between fourteen genes according to the above two principles. Five indepedent genes in addition to four linkage-groups were found, whereas the number of bivalent chromosomes is only seven. Since a possibility of two linkages with about 50 $\%$ crossing-over between two of the independent genes was left, these results were not considered to form final proof of disharmony between the number of linkage-groups and the number of chromosomes.

The relations of two more genes to the above fourteen are now known. The general conclusion is similar as formerly arrived at, namely nine independent genes or groups of linked genes. Some relations, however, were found to be different from what was supposed before.

II. MATERIALS AND METHODS

w 1. The genes

The following fourteen genes were studied before:

- (1) A : colored flower (recessive: white flower).
- (2) B : purple flower (rec.: pink).
- (3) D^w : double leaf-axil-spot. (Of the two recessive allelomorphs which are known, D occurs in the present material, standing for single axil-spot).
- (4) *Fa :* normal stem (rec.: fasciated).
- (5) $G\phi$: green pod (rec.: yellow pod).
- (6) P : membrane in the pod-wall (rec.: no membrane).
- (7) V : strong membrane in the pod-wall (rec.: thin membrane in the presence of P).
- **(8) Pl :** black hilum (rec.: not black).

(9) M : marbled seed-coat (rec.: not marbled).

(10) F : stippled seed-coat (rec.: not stippled).

- (11) I : yellow cotyledon (rec.: green).
- (12) *Le :* long internode (rec.: short).
- (13) \widetilde{W}^b : glaucous stem and foliage (rec.: emerald).
- (14) S : seeds free in the ripe pod (rec.: seeds adhering "chenille"like).

To these fourteen genes are added in the present study:

- (15) *T1 :* tendrilled leaves (rec.: acacia).
- (16) \widetilde{K} : normal wings (rec.: keeled wings).

Tl was first described by VILMORIN (12) in 1910 and since occurred in several studies. PELLEW and SVERDRUP (9) were the first to study K which gene has not yet been present in the material of other investigators. Genes B , D^{ψ} , M and F are only active in the presence of *A,* while V only acts in the presence of P.

A striking cooperation exists between A and B on the one side and K on the other side. PELLEW and SVERDRUP $(9, p. 127)$ already hinted at this cooperation, while SVERDRUP $(11, p. 225$ and plate XVII) went into more details. She states: "The wings more or less take on the colour of the keel, giving a much paler appearance to the whole flower". A somewhat different explanation seems to be more evident, however. In keeled winged purple or pink flowers the most conspicuous part of the wings, in which in normal forms the typical pigment develops, is almost completely reduced. Consequently the color genes do not find the substratum where they can act and no pigment is formed. This explanation fits perfectly well in GOLDSCHMIDT's theory on the action of genes (1, see especially pp. 112-113). Classification after flower-color in keeled-winged forms is easily done by inspecting the color of the standards.

w 2. The parents and the crosses

Genes *tl* and *k* were brought in my material in a line which I obtained'through the kindness of miss C. PELLEW. It was unnamed originally and I indicated it as "Keeled Acacia". The other parental lines are the same as those used before. The genotypes $-$ simply expressed in single symbols $-$ are:

1-2-56-1 *A B Dw fa Gp P v pl m F I Le Wb* S Tl K P. thebaicum $A b D F a G p P V P I M f I L e W b S T I K$ Pois à cosse jaune: a B Dw Fagpp V pl m F I Le W^2_b S Tl K Pois à brochettes: *a B Dw Fa Gp P V pl m F i le wb s Tl K* Keeled Acacia : A b D fa Gp P V pl M f I Le W_2° S $t l k$ The new crosses, with the genes involved in each of them, are: Cross 42. 1-2-56-1 \times K. Acac.: *B-Dw-V-M-F Tl-K.* Cross 43. K. Acac: \times P. thebaicum : Fa-Pl-Tl-K. Cross 44. K. Acac. \times P. à c. jaune : \overrightarrow{A} - \overrightarrow{B} - \overrightarrow{D} w-Fa-Gp-P-M-F-Tl-K. Cross 45. K. Acac. \times P. a broch.: $A-B-Dw-Fa-M-F-I-Le-Wb-S-TI-K$.

The numbers of F_2 -individuals are given in the following table:

cross- number	sown	succum- $_{\rm bed}$	$\%$ suc- cumbed	germ plants	full grown plants
42	1064	10	0.9	1054	1033
43	1089	61	5.6	1028	1017
44	372	4	1.1	368	355
45	1065	10	0.9	1055	1038

The F_2 of cross 43 was sown two weeks before the F_2 of cross 45 and four weeks earlier than the F_2 's of crosses 42 and 44. This might account for the relatively bad germination of 43. Unfortunately the number of individuals in 44 is considerably smaller than in the other F_2 's. This is caused by a relatively bad succeeding of the crossings, so that the number of F_1 -plants was already small. This is nothing but accidental, since the same crossing succeeded well next year. Another factor which reduced the number of seeds produced by the F_1 -plants, is the inherited dominant abortion (18) which occurs in cross 44, but which will not be discussed in the present paper, since its relation to *Pl* in Still unknown and our information about it is consequently incomplete. The number of individuals for either B , D^w , M or F and for the relations involving these is about $\frac{3}{4}$ of the total number when A also segregates, because in the $\frac{1}{4}$ part which is *aa*, named genes cannot demonstrate their action.

w 3. Growing methods, notes, calculations

The treatment of the material was exactly the same as in former studies $(15, 16)$. As stated above, the sowing-time was varied in the different F_2 's in order to obtain different flowering-times. The F_2 's which were expected to blossom relatively early, were sown first, those expected to flower rather late, were sown later. This prevented too great an accumulation of work during the flowering-period. No classification difficulties were met with, except perhaps sometimes the differentiation between normal and fasciated stem (ep. p. 10). Summarizing tables of the total F_2 -results were only prepared for crosses 42 and 43; they are given as appendix to this paper 1). When the number of genes studied in a certain cross exceeds seven $-$ as in crosses 44 and $45 - a$ summarizing $F₂$ -table gets too complicated. In these cases the mono- and bifactorial segregations are made up immediately from the field-notes.

Two types of calculations are involved in linkage-studies. The first is the calculation of expectations in case of independence. When these expectations do not fit the observations, linkage is evident and the crossing-over-percentage must be determined. KAPPERT'S method (6) of calculating the most probable expectation in case of independence has been very useful in former studies and is again used in the present material. As a rule, a larger value than 3 for actual deviation divided by mean error of expectation indicates linkage. Several methods are proposed for the determination of the percentage of crossing-over from F_2 -material. It is quite natural that I prefer my own method (13) which reveals the gametic ratio $p : q : r : s^2$ from the following equations, in which *(AB), (Ab), (aB)* and *(ab)* are the phenotypie F_2 -groups and *n* is the number of F_2 -individuals:

	$s = \sqrt{\overline{(ab)}}$
	$r = \sqrt{\overline{(ab)} + \overline{\overline{(a)}}} - s$
	$q = \sqrt{(ab) + (Ab)} - s$
	$p = \sqrt{n} - (q + r + s)$

¹⁾ The order of the genes in the summarizing tables is different from the order in the text; this is due to the order of observation which is different from the historical arrangement in the text.

^{2) ,} Symbols $p \ldots r$ are used here to avoid confusion, since $q \ldots d$ which were used before (13}, are used in the trifactorial ratio turther on.

As a check we have:

 $(AB) = p^2 + 2 pq + 2 p r + 2 p s + 2 q r$

The calculation of interference is sometimes a very helpful means to determine whether linkage is present between three genes. For this purpose we should know the trifactorial gametic ratio which is also necessary for the calculation of the percentage of double crossing-over. A similar method as the one described above for a bifactorial ratio may be used.

From an F₂-checkerboard with a gametic ratio $a:b:c; \ldots, k$ we read:

n2 : (a+b+c+d+e+/+g+h) 2 (9)

The solution proceeds by substituting the value for h from (8) respectively in (7), (6) and (4) which reveals g, f and d . Next e, c and b are found from (5) , (3) and (2) and α follows from (9) . Equation (1) is used as a check.

This calculation is rather laborious. It is much simpler to calculate separately the three bifactorial gametic series:

can substitute:

for (I) $(a + b) : (c + d) : (e + f) : (g + h)$ for (II) $(a + c) : (b + d) : (e + g) : (f + h)$ for (III) $(a + e)$: $(b + f)$: $(c + g)$: $(d + h)$

Consequently the above 12 values $(a + b)$, $(c + d)$ etc. are known, but the values for $a, b \ldots$, h cannot be determined, since some of the twelve equations are dependent. Fortunately h is easily deter-

mined from (8) as $h = \sqrt{\overline{(abc)}}$ and simple substitutions reveal a, b, etc. If, for instance, $h = 5$, we know d from the value for $d + h$, we know c from $c + d$ etc.

Of course, the above methods do not give but approximate indications, since the total number of gametes, being \sqrt{n} , is small even in an $F₂$ with very large numbers. But when back-crossing is hard to be done, as in peas, an approximate indication is preferable to no indication at all.

III. EXPERIMENTAL RESULTS AND CONCLUSIONS

w 1. The mono/actorial F.,-segregations

In the tabulation of the monofactorial F_2 -segregations, the theoretical expectation is printed in parentheses. Furthermore, actual deviation divided by mean error is indicated as c.

8

Cross 44. 268 : 87
\n
$$
(266.25) : (88.75)
$$
\n
$$
c=0.2
$$
\nCross 45. 803 : 235
\n(778.5) : (259.5)
\n
$$
c=1.8
$$

The only value for c which is too large $-$ accepting c=3.0 as an extreme chance variation $-$ is in the case of Fa in cross 43. Since sometimes the fasciation of the stem is modified so as to make it hardly distinguishable from normal stems, classification-mistakes may account for this large deviation.

w 2. The bi/actorial segregations

As in earlier studies, the factorial relations are published in the following way. At the left of the vertical line the four phenotypic F_{2} -classes are given on the first row and the corresponding expected values in case of independence are given on the second row in parentheses. At the right of the vertical line the ratio of sum of outer terms to sum of inner terms of the four class ratios at the left are given. These ratios indicate sum of parental types to sum of recombinations, or as the case is the reverse. Again, c is actual deviation divided by mean error of the two class ratio.

Gametic ratios are only given in case of linkage and in some more cases which have to be especially discussed in the following paragraph. In case of a $9:3:4$ expectation, namely in those relations involving A and a gene depending in its action on presence of A (either B, D^{ω} , M or F), only the first two terms can be used in deciding whether or not linkage is present. Independence is then indicated by agreement of the first two terms with a 3 : 1 expectation.

The way of numbering the 120 possible relations between the 16 genes will seem illogical at first sight, but will be comprehensible from a historical point of view. The relations which already occurred in Linkage-Studies I or II are namely numbered as they were before. This means for instance that relation *A-F* is 9, that *A-I* is 46 and that *A-Tl* is 92. When all the relations were studied in one year, these numbers would have been 9, 10 and 14.

 \sim

w 3. Conclusions

In discussing the conclusions which can be drawn from the factorial relations, we shall first treat those relations which were studied in "Linkage-Studies I or II" and occur again in the present material.

(1) The linkage-group *B-F-M.* It was concluded before that B , F and M are located in one chromosome in the given order and that the percentage of crossing-over for $B-M$ is about 50 $\%$, while also the linkage *F-M* is so weak that in some crosses it does not immediately demonstrate itself (cp. 15, pp. 457-460;16, pp. 286-287). Similar conclusions can be drawn from the present material. *B-M* (relation 16) do not show any linkage, *B-F* (rel. 17) are clearly linked in all three

crosses 42, 44 and 45, while *M-F* (rel. 45) give a rather high value for c in all the crosses where they occurred which value however is too high only in cross 42. The following mapping would follow from the crossing-over percentages:

The percentages of double crossing-over were calculated to be resp. $-3.3 \frac{\%}{0}$, 1.2 $\%$ and 3.2 $\%$. The following interference-relations were found:

Of course the negative values in cross 42 , both for the percentage of double crossing-over and for some of the values in the above table make application of the calculation-method used questionable. Yet it is remarkable that in earlier material, entirely different from the material in cross 42, the same thing was found (16, p. 286). Moreover, the map for cross 42 makes us expect something like a negative crossing-over, since the value for *B-M* is larger than the sum of *B-F* and *F-M.*

I wish to draw the attention to a remarkable relation among the values in the interference-table. The sum of the values on the second and eighth line is always 100 $\%$ and the same is true for those on the fourth and the sixth line. We can express this relation as follows: Crossing-over between $F-M$ means 100 $\%$ crossing-over for *B-F* and $B-M$ together. This is quite plausible, however, for crossing-over between *F-M* means 100 $\%$ crossing-over for *B-M* minus the double crossing-over; now this double crossing-over is the value for the crossing-over *B-F* among the cross-overs *F-M.* Consequently the above rule is a general one and again it is remarkable that the negative values in cross 42 confirm to this rule. This is the more remarkable since these values, according to the above, represent double crossovers, so that in cross 42 all percentages of double crossing-over are negative.

If we put aside the questionable interference-relations in cross 42, the values for the other crosses indicate a distinct interference, from which we can conclude that B , F and M are located in one chromosome in the given order. As to the considerable variations in the percentages of crossing-over in the three crosses, we see that cross 44 has given the lowest values, while cross 45 gave the highest values and cross 42 is in between the other two; only the value for *B-M* in cross 45 is an exception. We *may,* therefore, suppose that the genotypes of the parents of the cross determine the rate of crossing-over in general, as was found by RASMUSSON (10) for other cases of linkage in *Pisum* and which, for the present material, was already demonstrated before (15, pp. 459-460).

(2) The linkage-group *Dw-V-Le.* Formerly *Dw-Le* (rel. 55) were found to be independent, c being $\overline{1.3}$ (16, p. 282). In cross 45 however, $c=4.6$ which indicates linkage. The crossing-over percentage is 63.6% . This is extremely high, but recently a similar value was found for two other genes in back-cross material (17). The extremely loose linkage is better comprehensible than the fact that the genes in question are independent in the one cross and linked in the other. However, more suchlike cases are known in *Pisum.* They may be explained either by variation in the crossing-over percentage, as especially worked out by RASMUSSON (10), or by chromosome-linkage, as was supposed by HÅKANSSON (2) for genes A and $G\phi$ in a certain cross of HAMMARLUND $(3, 4, 5)$. The crossing-over for D^w -Le in the old material would have been 53.9 $\%$, so that the variation from this value in 63.6 $\%$, as occurring in the present material, is not very high It is quite possible, therefore, that the genes *Dw* and *Le,* formerly supposed to be independent, are in reality linked, as was demonstrated above. Now V and Le were formerly found to be linked with 13.8 $\%$ crossing-over (16, p. 283), so that a linkage-group D^{w} -V-Le should occur. It is true that $D^{w}V$ (rel. 21) were found independent (confirmation of former results, see 15, p. 454), but we may have 50 $\%$ crossingover. Consequently we may conclude that the existence of a linkagegroup *Dw-V-Le* in which $D^{w-}V$ show about 50 $\%$ crossing-over and $D^{w}\text{-}Le$ over 60 %, is possible. Unfortunately the three genes did not occur in one cross, so that the interference-relations could not be studied. Therefore further confirmation is wanted, since also the possibility of chromosome-linkage remains. I am inclined to favor the hypothesis that linkage with variable crossing-over percentages is present, since extreme variations in the strength of linkages are quite common in *Pisum.* It is worth while noting that in the linkage-group D^{w} -*V-Le* almost identical values for the crossing-over percentages would occur as in the group *Gp-N-Ss,* recently described by the present author (17).

(3) The relation *F-Le.* The value for c in relation 83 of cross 45 is just a little higher than the limit which is usually accepted for independent inheritance. Therefore we deal with a doubtful case. Some facts speak against linkage: In former material (16, p, 284) $c=2.0$ which indicates independence. As discussed above sub (1) F is linked to B and M; no indications of linkages $B-Le$ (rel. 51) and $M-Le$ (tel. 79) were obtained, however. At last, since *Le* is probably linked with D^w [as discussed sub (2)], there should be a linkage-group D^{w} -F-*Le* with crossing-over percentages of 48.4 $\%$ for *D*^w-F (rel. 24), of 60.1% for *F-Le* (rel. 83) and of 63.6 % for *Dw-Le* (rel. 55) with a double crossing-over of 22.4 $\%$. The interference-relations in this group are as follows:

The first and the second two cases might indicate location in one chromosome, the last four speak absolutely against it. All cases, however, are readily explained by assuming that *Dw* and *Le* on the one hand and F on the other hand are located in different chromosomes. Thus:

Dw Le F --f -I -- 1

A similar reasoning as was held in Pisum-Crosses III (14, pp. 234-236) makes us conclude that the above situation is quite plausible, when we keep in mind that the crossing-over percentage for D^w -Le is more than 50 $\%$.

No other relations which were already studied before, need special discussion, so that we can proceed to the new relations.

(4) The linkage group $K-Wb-S$. The linkages, found in cross 45 for W_2^b -S (rel. 91), W_2^b -K (rel. 117) and *S-K* (rel. 119) suggest the following localization:

$$
K \leftarrow \longrightarrow 15.6 \text{ % } \longrightarrow W_b \leftarrow 0.8 \text{ % } S
$$

The order of W_2^b and *S*, as supposed above, cannot be considered as final, since the difference in the crossing-overs $K-W_2^b$ and $K-S$ is so extremely small. Calculation of the interference-relations cannot be expected to settle this question, since the cross-overs between $W^b_{\frac{2}{3}}$ and S are so few. Indeed, the calculated interference-relations are inconclusive and therefore will not be mentioned. Localization of K , W_2^b and S in one chromosome cannot be doubted, but the exact order might also be: K , S , W_2^b .

 (5) Final conclusions. All relations, not yet disscused, were found to be independent. When we put linked genes between brackets and separate independent genes or groups of linked genes by a dash, we arrive at the following conclusions:

Cross 42:
$$
[B, F, M]-D^{w} - V - I - K
$$
\nCross 43: $F_a - P l - I - K$ \nCross 44: $A - [B, F, M] - D^{w} - F_a - G_p - P - I - K$ \nCross 45: $A - [B, F, M] - [D^{w}, L e] - F_a - I - [K, W_a, S] - I$

The final conclusion of Linkage-Studies I and II (16, p. 287) was: $A-[B, F, M]-Dw-Fa-Gp-[P, Pl]-[V, Le]-I-[W^b_2, S]$

When we keep in mind that D^w and V in cross 42 are evidently linked with about 50 $\%$ crossing-over which changes the last mentioned series by bringing D^w in the group $[V, Le]$, the total results are:

 $A-[B, F, M]-[D^w, V, Le]-Fa-Gp-[P, Pl]-I-[K, W^b_2, S]-Tl$

The number of chromosomes is exceeded by two by the above number of nine independent groups or single genes.

IV. DISCUSSION

In so far as the relations 1 up to 91 were studied by other investigators, these results have been discussed in Linkage-Studies I and II (15, pp. 461-462; 16, pp. 288-290). Since then NILSSON (8, p. 263) obtained some evidence for independence of *V-I* (rel. 70) or of *P-I* (rel. 66); it cannot be said with certainty whether V or P is the gene in question. Anyhow do these results confirm my former conclusions (16, pp. 282-283).

Among my new relations 92 up to 120 the following were also studied by others. *Tl* on the one side and *A, B, Fa, I* and *Le* on the other side (relations $92, 94, 98, 112$ and 114) were found to be independent by WHITE (19, pp. 175, 186-188). *W~-T1* (rel. 116) and *S-T1* (rel. 118) occurred in MEUNISSIER'S crosses (7, p. 317) and evidently were independent, although the numbers are small. PELLEW and SVERDRUP (9) found linkage with about 20 $\%$ crossing-over between W_b -K (rel. 117}, while K on the one side and *A, B, Fa, I, Le* and Tl on the other side (relations 93, 95, 99, 113, 115 and 120) were said to be independent.

SVERDRUP (11) gave more evidence for these statements and moreover found independence for the same relations as studied by WHITE and mentioned above. Once more I draw the attention to the independence of D -Le (rel. 55), found by SVERDRUP, because these genes were linked in my present material and independent in my former crosses, as was discussed above. For the rest my results confirm those of the other *Pisum-workers*.

Not much need to be said about the disagreement between the

number of chromosomes and the number of independent genes or groups of linked genes, as occurring in my material. This question was fully discussed in "Linkage-Studies II" (16, pp. 289-290) and the present situation is essentially the same as in the earlier group of genes. It is true that $D\mathbf{w}$, formerly thought to belong to the "single" genes", evidently was brought in the linkage-group *V-Le,* but *TI* has taken its place as "single gene". Consequently four linkage-groups in addition to five independent single genes remain and also the possibility of two linkages with about 50 $\%$ crossing-over among the five single genes remains. This would reduce the number of independents to seven. New crosses, adding six more genes to the sixteen already studied, are being investigated and will perhaps settle this question.

V. SUMMARY

1. In former studies the relations between the fourteen genes $A, B, D^{\omega}, Fa, G\phi, P, V, Pl, M, F, I, Le, W_{\phi}^{b}, S$ were studied. In the present study two more genes were added to the above fourteen namely:

TI : tendrilled leaves; *tl* : "acacia"-leaves.

- \widetilde{K} : normal wings; k : keeled wings.
- 2. Putting linked genes between brackets and separating independent genes or linkage-groups by a dash, the total results can be expressed as:

$A\text{-}[B, F, M]\text{-}[D^\textit{w}, V, L^\textit{e}]$ -Fa-Gp- $[P, Pl]$ -I- $[K, W_2^\textit{b}, S]$ -Tl

Further confirmation is wanted with regard to the position of *Dw* in the second linkage-group.

- 3. The number of bivalent chromosomes is still exceeded by two by the number of independents, but two linkages with about 50 $\%$ crossing-over between two of the single genes might occur.
- 4. Methods for calculating trifactorial gametic ratios from given $F₂$ ratios were described on pp. 6--7.

ACKNOWLEDGEMENT

I will not finish this paper which is the last one I publish during my stay at Wageningen, Holland, without expressing my sincere thanks

to Professor Dr. J. A. Honing. During the ten years of my Pisumwork Prof. HONING gave me all possible facilities in his fields and laboratory, although I did not belong to his staff. Moreover I am indebted for his constant interest and critical remarks during the progress of the work.

Wageningen, Oct. 22, 1922

LITERATURE CITED

- (1) GOLDSCHMIDT, RICHARD: Physiologische Theorie der Vererbung. Berlin, Julius Springer, 1927, VI $+$ 247 pp., 59 fig.
- (2) H[°]AKANSSON, ARTUR: Chromosomenringe in Pisum und ihre mutmassliche genetische Bedeutung. (Hereditas z2, 1929: 1-10.)
- **(3) HAMMARLUND, C.:** fiber einen Fall yon Koppelung und freie Kombination bei Erbsen. (Hereditas 4, 1923: 235-238.)
- **(4) HAMMARLUND, C.:** Zweite Mitteilung iiber einen Fall yon Koppelung und freier Kombination bei Erbsen. (Hereditas zo, 1927- '28: 303-327.)
- **(5) HAMMARLUND, C.:** Dritte Mitteilung iiber einen Fall yon Koppelung und freier Kombination bei Erbsen. (Hereditas 12, 1929: 210-216.)
- (6) KAPPERT, H.: Über die Auswertung dihybrider Spaltungsreihen bei Koppelungsstudien. (Zschr. ind. Abst. u. Vererb. L. *44,* 1927: 303-314.)
- (7) MEUNISSIER, A.: Observations sur l'hérédité du caractère "Pois à trois cosses" et du caractère "Pois chenille". (Genetica 4, 1922: 279-320,)
- -(8) NILSSON, ERNST: Erblichkeitsversuche mit Pisum, II. (Hereditas z2, 1929: 223-268.)
- (9) PELLEW, CAROLINE and ASLAUG SVERDRUP: New observations on the genetics of peas (Pisum sativum). (J. Gen. x3, 1923: 125-131 .)
- (10) RASMUSSON, J.: Genetically changed linkage values in Pisum. (Hereditas Io, 1927: 1-152.)
- (11) SVERDRUP ASLAUG: Linkage and independent inheritance in Pisum sativum. (J. Gen. 17, 1927: 221-251.)
- (12) VILMORIN, PH. DE: Recherches sur l'hérédité mendélienne. (C. R. Acad. Sc. Paris 1910-II: 548-551.)
- (13) WELLENSIEK, S. J,: Methods for calculating the actual gametic $F₂$ series from a given zygotic series. (Genetica 9, 1927: 329-340.)
- (14) WELLENSlEK, S. J.: Pisum-Crosses, III. (Genetica 11, 1928: 225-256.)
- (15) WELLENSIEK, S. J.: Linkage-Studies in Pisnm, I. (Genetica 9, 1927: 443-466.)
- (16) WELLENSIEK, S. J.: Linkage-Studies in Pisum, II. (Genetica 11, 1929: 273-292.)
- (17) WELLENSIEK, S. J.: The occurrence of more than 50 $\%$ crossingover in Pisum. (Genetica 11, 1929: 509-518.)
- (18) WELLENSIEK, S. J. and J. S. KEYSER: Pisum-Crosses V: Inherited abortion and its linkage-relations. (Genetica 11, 1929: 329-334.)
- (19) WHITE, ORLAND E,: Inheritance Studies in Pisum-IV: Interrelation of the genetic factors of Pisum. (J, agr. Res. 11, 1917: 167-190.)

SUMMARIZED TOTAL F_2 -RESULTS OF CROSS 43: $\underbrace{fa\;tl\;k\;pl}_{\sim}\times\ \underbrace{Fa\;Tl\;K\;Pl}_{\sim}$