

In general, these data agree with the $v_3 - \underline{ae} - \underline{pr}$ data presented above. There seems to be a high frequency of associated crossovers in these data. It appears that whenever a cross over occurs between \underline{bv} and \underline{ae} there tends also to be one between \underline{ae} and \underline{pr} . This would result in a reduced recombination percentage between \underline{bv} and \underline{pr} as is shown by the 16% recombination.

These data indicate (1) an increase in the total number of double crossovers, as well as (2) an increase in the number of two-strand double crossovers.

c. $\underline{bm}_1 - \underline{ae} - \underline{pr}$ linkages.

A cross of $\frac{+ \underline{ae} +}{\underline{bm}_1 + \underline{pr}} \times \frac{\underline{bm}_1 \underline{ae} \underline{pr}}{\underline{bm}_1 \underline{ae} \underline{pr}}$ gave the following results:

+++	+ ae +	++ pr	+ ae pr	bm ++	bm ae +	bm + pr	bm ae pr
1	124	27	11	16	10	141	3

Recombination %

$\underline{bm}_1 - \underline{ae}$	12.3
$\underline{ae} - \underline{pr}$	9.3
$\underline{bm}_1 - \underline{pr}$	19.2

Individual Gene Segregation

\underline{Ae} 185:	\underline{ae} 148*
\underline{Bm}_1 163:	\underline{bm}_1 170
\underline{Pr} 151:	\underline{pr} 182

* Significant deviation from 1:1 ratio.

These recombination values appear to be normal in that the $\underline{bm}_1 - \underline{pr}$ recombination is what one expects from the $\underline{bm}_1 - \underline{ae}$, $\underline{ae} - \underline{pr}$ recombination observed. However, two things are aberrant in these data: (1) the recombination values are smaller than expected when they are compared with the $v_3 - \underline{ae} - \underline{pr}$ and $\underline{bv} - \underline{ae} - \underline{pr}$ data presented above, and (2) the \underline{Ae} vs \underline{ae} segregation deviates from a 1:1 ratio.

These data indicate differential transmission of the \underline{ae} and \underline{Ae} alleles through the female. A comparison of the reciprocal crossover classes supports differential transmission rather than misclassification of \underline{ae} .

Differential transmission would reduce the measurable recombination between the genes and could account for the observed differences between these recombination values and those mentioned above for $v_3 - \underline{ae} - \underline{pr}$ and $\underline{bv} - \underline{ae} - \underline{pr}$.

-- J. N. Jenkins

2. A new locus for studying the fine structure of the gene.

A method for studying the fine structure of the \underline{ae} locus has been developed. By overstaining with an excess of iodine and destaining with 25% alcohol and slight heating, one is able to differentiate $\underline{wx} \underline{ae}$ from $\underline{wx} +$ pollen. The $\underline{wx} \underline{ae}$ pollen stains black and the $\underline{wx} +$ pollen stains red. This technique allows a study of recombination at the \underline{ae} locus in a \underline{wx} background when different sources of \underline{ae} are crossed and the F_1 pollen is observed.

-- Roy G. Creech