2. Striate-2 (sr_2).

This character has been reported under the name waseca stripe in the Corn News Letter 27 (page 66) and 29 (page 54). Further linkage data on it were obtained in 1955. These confirm its location on chromosome 10. The genetic constitution of the striate stock with respect to the aleurone color series was found to be $A_1A_1A_2A_2A_3A_3$ cc rr ii (not a_3a_3 as was previously stated).

The following data were obtained from backcrosses, F_2 's and combined backcross and F_2 data, and originated from ears segregating for one color factor (R) only. Heterogeneity tests were applied to the different groups of data and indicated that the data were homogeneous. The following table shows the source of the data, number of plants classified and the percent crossing over. All crosses were in the coupling phase.

golden₁ vs r.						
Source	N	% c.o.				
3 point backcross test F ₂ data (3:1)(1:1) data	436 231 1360	20.0 18.5 16.0				
striate ₂ vs r.						
3 point backcross test F ₂ data all backcrosses striate ₂ vs g	436 231 1796 olden	25.0 31.5 25.1				
	0 20.0					
3 point backcross test F ₂ data (3:1)(1:1)	436 231 1360	44.9 45.0 40.0				

The detailed data on the three point test are as follows:

grs	129	g++	38	gr+	49	g+s	3	
+++	118	+rs	42	++S	53	+r+	4	total 436.

Therefore the order of the genes is $golden_1 - R - striate_2$, as already reported. (Corn News Letter 29).

The striate stock had also been crossed on to nana-2. From three F_2 progenies it was concluded that nana-2 was independent of R and of striate-2. One ear segregated for Pr vs pr: 82 PrNa: 14 Prna: 13 prNa: 27 prna. The data suggest linkage between nana-2 and pr.

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