



differs from  $\underline{R}^{st}$  in time of pigment formation. Three kinds of nonparental kernel phenotypes were observed among 68,883 seeds from  $\underline{g} \underline{R}^{st} \underline{M}^{st}/+ \underline{R}^{nj} +$  o X  $\underline{g} \underline{r} +$  ♂ matings: colorless, self-colored, and Navajo-stippled compound ( $\underline{R}^{nj}:st$ ). The colorless type occurred infrequently and each proved to be either  $\underline{R}^{nj}$  or  $\underline{R}^{st}$  in progeny tests. As self-colored kernels were frequent, only those from a portion of the total population were chosen for progeny analysis. The results of testing 25 for germinal transmission and linked marker composition are given in the first two lines of the table below:

Phenotype Kernel      Progeny		Effective kernel population	Linked marker composition			
			Parental		Recombinant	
			$\underline{g} \underline{M}^{st}$	$++$	$\underline{g} +$	$+ \underline{M}^{st}$
$\underline{R}^{sc}$	$\underline{R}^{st}$	14,035	11	0	0	1
$\underline{R}^{sc}$	$\underline{R}^{sc}$	14,035	6	0	4	3
$\underline{R}^{nj}:st$	$\underline{R}^{nj}:st$	62,815	0	0	0	15

The fraction of self-colored kernel selections which proved germinal, about one half, and the frequency of this class,  $18.5 \times 10^{-4}$  (based on the number of  $\underline{R}^{st}$  gametes), agree with the corresponding values obtained for  $\underline{R}^{st} \underline{R}^{st}$ . The distribution of linked markers also conforms to that predicted from the findings with  $\underline{R}^{st} \underline{R}^{st}$ . Four  $\underline{R}^{sc}$  mutants were found to carry the  $\underline{g} +$  combination whereas expectation based on independence of mutation from recombination predicts less than one. To permit interpretation of these four as separation of the two components postulated for  $\underline{R}^{st}$  by single exchanges, the location of the basic seed pigmentation element should be proximal to the element suppressing its action.

The third nonparental phenotype,  $\underline{R}^{nj}:st$ , strongly resembles the case described by Brink (M.G.C.N.L. 34:122). The typical stippled pattern is confined to those regions of the kernel normally pigmented by  $\underline{R}^{nj}$ . Each of the 15 progeny groups recovered from 16 initial  $\underline{R}^{nj}:st$  kernel selections yielded the compound phenotype and are found to correspond to  $\underline{R}^{nj}$ , rather than  $\underline{R}^{st}$ , in time of pigment formation. They therefore appear to combine the basic  $\underline{R}^{nj}$  pigmentation property with the instability of stippled. Each proved to carry the proximal marker of the  $\underline{R}^{nj}$  chromosome and the distal marker of the  $\underline{R}^{st}$  homologue. That this marker combination is complementary to the principal recombinant class of  $\underline{R}^{sc}$  mutants suggests that the two originate as reciprocal products of a single kind of recombinational event. Neither  $\underline{R}^{sc}$  nor  $\underline{R}^{st}:nj$  expresses the characteristic plant color effects of  $\underline{R}^{nj}$ , however, so either the two are not strictly complementary, or plant pigmentation is suppressed when brought into cis position with the inhibiting component of  $\underline{R}^{st}$ .

J. L. Kermicle