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1. Behaviour of half-opaque maize.

Eleven inbred lines have been selected for incorporation of the opaque-2 gene. After two backcrosses, the opaque kernels from the different lines were utilized in developing an opaque-2 synthetic. In the second generation of the synthetic opaque-2, several sectored kernels with normal (translucent) tissue were observed. These were classified into seven distinct types (S_2 to S_8). Diallele crosses were made among these including complete opaque (S_1). After two generations of selfing, certain crosses in the F_3 gave mainly half opaque kernels (S_5).

Table 1. Half opaque maize from certain diallele crosses

Cross	F ₁	F ₂	F ₃		
s ₂ x s ₄	s ₅	s ₅ , s ₅	s ₅		
s ₄ x s ₃	\$ ₅	s <u>*</u>	s ₅		
s ₄ x s ₇	s ₂	s ₅ , s*	s ₅		
$s_4 \times s_7$	s 6	s *	\$ ₅		
s ₄ x s ₇	s ₅	s ₅ , s*	s ₅		
s ₄ x s ₈	s ₅	s ₅ , s•	s ₅		
s ₅ x s ₆	s_8	s ₅ , s*	s ₅		
s ₆ x s ₄	s ₅	s ₅	s ₅		

Note: $S_5^* = 1/3 / 3/4 \underline{o}_2$ $S_5 = 1/2 \underline{o}_2$ The lysine content of the sectored seed has been analyzed and it was found to be similar in all the types including half opaque suggesting that the sector size may not alter the lysine content. The modified opaque may be more useful in breeding high lysine maize and also may be more acceptable for direct consumption.

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2. Electrophoretic studies of half opaque endosperm (S5).

One of the S_5 cobs from the cross S_4 x S_7 was selected for biochemical studies. The normal half, $S_5(\pm)$, and opaque half, $S_5(\underline{o}_2)$, were separated and the soluble proteins extracted with .Ol M sodium pyrophosphate buffer containing 10^{-4} M EDTA and 0.7 ml of mercaptoethanol. The $S_5(\pm)$ and $S_5(\underline{o}_2)$ tissues were also subjected to three consecutive extractions with water, 5% NaCl and 70% ethanol. All these protein extracts were subjected to electrophoretic separation in 7.5% polyacrylamide gels. The saline extract was dialysed before electrophoresis. Table 1 gives the pattern of protein bands in both translucent and opaque tissues of S_5 .

Tissue	1	2	3	4	5	6	7	8
S ₅ (+)	I	I	М	М	A	I	М	F
	A	M	A	M	M	M	M	M
s ₅ (+)	F	A	I	М	F	F	F	
s ₅ (0 ₂)	M	M	M	F	F	F	A	
s ₅ (+)	F	M	F	М				
s ₅ (0 ₂)	F	F	A	A				
S ₅ (+)	F	F	F					
s ₅ (₀₂)	A	A	A					
	s ₅ (+) s ₅ (o ₂) s ₅ (+) s ₅ (o ₂) s ₅ (+) s ₅ (o ₂) s ₅ (+)	S ₅ (+) I S ₅ (o ₂) A S ₅ (+) F S ₅ (o ₂) M S ₅ (+) F S ₅ (o ₂) F S ₅ (+) F	S ₅ (+) I I S ₅ (o ₂) A M S ₅ (+) F A S ₅ (o ₂) M M S ₅ (+) F M S ₅ (o ₂) F F S ₅ (+) F F	S ₅ (+) I I M S ₅ (o ₂) A M A S ₅ (+) F A I S ₅ (o ₂) M M M S ₅ (+) F M F S ₅ (o ₂) F F A S ₅ (+) F F F	S_{5} (+) I I M M S_{5} (o ₂) A M A M S_{5} (+) F A I M S_{5} (o ₂) M M M F S_{5} (o ₂) F M F M S_{5} (o ₂) F F A A S_{5} (+) F F F	S_{5} (+) I I M M A S_{5} (o ₂) A M A M M S_{5} (o ₂) A M A I M F S_{5} (o ₂) M M M F F S_{5} (o ₂) F M F M S_{5} (o ₂) F F A A S_{5} (o ₂) F F F F	S_{5} (+) I I M M A I S_{5} (o ₂) A M A M M M M S ₅ (+) F A I M F F S ₅ (o ₂) M M M F F F F S ₅ (+) F M F M S ₅ (o ₂) F F A A S ₅ (+) F F F	$S_{5}(+)$ I I M M A I M $S_{5}(o_{2})$ A M A M M M M M $S_{5}(+)$ F A I M F F F $S_{5}(o_{2})$ M M M F F F A $S_{5}(+)$ F M F M $S_{5}(o_{2})$ F F A A $S_{5}(+)$ F F F

Table 1. Endosperm protein pattern of $S_5 \pm and S_5 \underline{o}_2$

⁽I = Intense, M = Medium, F = Faint, A = Absent)