

## Genetic control of seed coat texture in cowpea, *Vigna unguiculata* (L.) Walp.

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### ABSTRACT

Genetic control of seed coat texture was investigated in  $F_2$  and backcross populations involving seven accessions of cowpea (*Vigna unguiculata* (L.) Walp). In two of the four crosses, the seed coat texture was found to be under monogenic inheritance. In the other two crosses, the trait was found to be controlled by two genes with complementary effect, giving a segregation ratio of 9 smooth : 7 rough for  $F_2$  and 1 smooth : 3 rough for backcross generations.

**Key Words:** cowpea, seed coat texture, complementary gene action, segregation, backcross, *Vigna unguiculata*.

### INTRODUCTION

Cowpea, *Vigna unguiculata* (L.) Walp is an important grain legume crop in the tropical and sub-tropical regions of the world. It is of major importance to the livelihood of millions of relatively poor people in comparatively less developed countries of the tropics. In Nigeria, it is the most important source of plant protein with a high protein content of 20-25% (Stanton *et al.* 1966). One of the major problems of cowpea has been its low yield and efforts are being made to raise its productivity and quality. One of the main factors that determine the consumer acceptability of cowpea is the texture of the seed coat. Most consumers prefer the seed with rough coat texture because it cooks faster than the seed with smooth coat. Hence, understanding the mode of genetic control of this character in order to formulate appropriate breeding strategies is important.

Rawal (1975) and Rajendra *et al.* (1979) reported single gene inheritance for seed coat texture with smooth seed coat dominant to wrinkled/rough seed coat. Fery (1985) proposed the symbol  $P_c$  for this single gene. However, the behaviour of some lines in crosses made during a breeding programme was not in accordance with reports by earlier workers. It was then suspected that genetic control of this trait might not be the same in different accessions of cowpea. This study was therefore undertaken to investigate the nature of the genetic control of seed coat texture in a number of cowpea lines.

### MATERIALS AND METHODS

Seven cowpea lines were used for the study. The names, sources and seedcoat types of these lines are presented in Table 1. These lines have been maintained through several generations by selfing. However, to confirm that these lines were homozygous, an evaluation study was carried out in the green house at the International Institute of Tropical Agriculture (IITA), Ibadan, Nigeria.

Each line was planted in twenty plastic pots with two plants per pot. The plants were maintained to maturity and no segregation was observed. Their seeds were collected for use in hybridization. These lines were then combined in the following crosses:

IBPC x IT82E-9 (P<sub>1</sub> x P<sub>2</sub>)  
IBPC x IBS 876 (P<sub>1</sub> x P<sub>3</sub>)  
Ife BPC x TVu 4578 (P<sub>3</sub> x P<sub>4</sub>)  
R10028 x G2497 (P<sub>6</sub> x P<sub>7</sub>)

Crosses to produce  $F_1$ s and backcrosses to respective parents were made in the green house. The methods used were those described by Zary and Miller (1982). To produce  $F_2$  seeds, a portion of the  $F_1$  seeds was sown in the field. Seeds from each cross were planted into two rows with a spacing of 30cm within row and 50cm between rows.  $F_2$  seeds were harvested from the  $F_1$  plants and fumigated in containers with phostoxin against storage pests until they were planted. Field evaluation of Parent 1, Parent 2,  $F_1$ , backcross to parent 1, backcross to parent 2 and  $F_2$  generations for each cross was done during the dry season at IITA, Ibadan, Nigeria. The field was irrigated once a week for 8-hours throughout the growing period. A randomized

complete block design with four replications was used. At maturity, when the pods had dried, the pods from each plant were harvested separately into a seed bag and labelled according to the plant identification number. Each plant was given a replication number, a row number and a within-row number. The seeds were visually scored into distinct classes of smooth and rough. The  $F_2$  and backcross to parents data were tested for their goodness-of-fit to appropriate genetic ratios by the Chi-square method (Gomez and Gomez 1984).

## RESULTS AND DISCUSSION

The results of the four crosses analysed to study seed coat texture are presented in Table 2. Parental lines  $P_1$ ,  $P_3$ , and  $P_7$  had rough seed coats while lines,  $P_2$ ,  $P_4$ ,  $P_5$  and  $P_6$  had smooth seed coats (Table 1). The  $F_1$  progenies of crosses  $P_3 \times P_4$  and  $P_6 \times P_7$ , had smooth seed coats and when they were backcrossed to the rough-seeded parents, a 1 smooth: 1 rough ratio was obtained. The  $F_2$  families of these crosses showed a segregation of 3 smooth: 1 rough, suggesting single gene inheritance of seed coat texture in these lines.

In crosses  $P_1 \times P_2$  and  $P_1 \times P_5$ , all  $F_1$  offspring had smooth seed coats but when backcrossed to  $P_1$  (the rough seeded parent), a 3 rough: 1 smooth ratio was obtained. Chi-square tests of the  $F_2$  families gave a good fit to the 9 smooth: 7 rough modified digenic inheritance ratio (Table 2).

Thus, the results of the inheritance study for seed coat texture showed clear inheritance patterns. The monogenic inheritance pattern observed in crosses  $P_3 \times P_4$  and  $P_6 \times P_7$  agree with the findings of Rajendra *et al.* (1979). They reported a single gene inheritance for seed coat texture. However, results of the crosses  $P_1 \times P_2$  and  $P_1 \times P_5$  indicated that the trait is controlled by two complementary genes. Franckowiak (1973) and Rawal (1975) reported rough testa texture to be controlled by at least two recessive genes and they recognised seed coat texture to be of vital importance in cowpea improvement. Therefore, it can be concluded that the genetic control of seed coat

Table 1. Names, sources and seed coat types of lines used in the study.

Name	Source*	Code Number	Seed Coat Type
IBPC	1	$P_1$	Rough
IT82E-9	2	$P_2$	Smooth
Ife BPC	3	$P_3$	Rough
TVu 4578	2	$P_4$	Smooth
IBS 876	1	$P_5$	Smooth
R 10028	1	$P_6$	Smooth
G 2497	1	$P_7$	Rough

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Table 2. Number of observed plants having different seed coat textures in parents and crosses of smooth and rough seeded lines of cowpea and the tests for goodness-of-fit to expected ratios.

Cross and Generation	Observed Number of plants			Expected Ratio	$\chi^2$	Probability (P)
	Smooth	Rough	Total			
<b>Ife BPC x TVu-4578</b>						
Ife BPC ( $P_3$ )	-	72	72			
TVu 4578 ( $P_4$ )	68	-	68			
$F_1$	95	-	95			
Ife BPC x $F_1$	129	126	255	1:1	0.035	0.75 < p < 0.90
TVu 4578 x $F_1$	137	-	137			
$F_2$	714	245	959	3:1	0.153	0.50 < p < 0.75
<b>R10028 x G2497</b>						
R10028 ( $P_6$ )	90	-	90			
G2497 ( $P_7$ )	-	78	78			
$F_1$	71	-	71			
G2497 x $F_1$	61	63	124	1:1	0.032	0.75 < p < 0.90
<b>R10028 x <math>F_1</math></b>						
$F_2$	380	-	380			
$F_2$	724	239	963	3:1	0.016	0.90 < p < 0.95
<b>IBPC x IT82E-9</b>						
IBPC ( $P_1$ )	-	20	20			
IT82E-9 ( $P_2$ )	62	-	62			
$F_1$	55	-	55			
IBPC x $F_1$	67	203	270	1:3	0.004	0.90 < p < 0.95
<b>IT82E-9 x <math>F_1</math></b>						
$F_2$	120	-	120			
$F_2$	451	331	782	9:7	0.643	0.25 < p < 0.50
<b>IBPC x IBS 876</b>						
IBPC ( $P_1$ )	-	42	42			
IBS 876 ( $P_5$ )	65	-	65			
$F_1$	70	-	70			
IBPC x $F_1$	70	210	280	1:3	0.000	> 0.99
IBS 876 x $F_1$	190	-	190			
$F_2$	355	270	625	9:7	0.077	0.75 < p < 0.90

- Indicates no plants.

texture in some cowpea lines is controlled by two genes with complementary effects.

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ABSTRACT

The objective of this study was to determine the effect of handcrossing on the yield and quality of cowpea (*Vigna unguiculata* L.) under semi-arid conditions. The study was conducted at IITA, Ibadan, Nigeria, during the 1997-98 growing season. The experimental design was a randomized complete block design with three replicates. The treatments were: (i) handcrossing, (ii) natural crossing, and (iii) control. The parameters measured were: yield (t/ha), seed yield (t/ha), straw yield (t/ha), protein content (g/kg DM), and digestibility (g/kg DM). The results showed that handcrossing significantly increased yield and seed yield compared to natural crossing and control. The protein content and digestibility were also significantly higher in the handcrossed plants. The authors concluded that handcrossing is a viable method for improving the yield and quality of cowpea under semi-arid conditions.

INTRODUCTION

Cowpea is known to be a high yielding crop with a high protein content. It is one of the most important legume crops in the tropics and subtropics. The yield and quality of cowpea are determined by a number of factors, including genetic, environmental, and management factors. Handcrossing is a method of breeding that involves the deliberate crossing of two different genotypes to produce a hybrid. This method has been used successfully to improve the yield and quality of many crops, including cowpea. The objective of this study was to determine the effect of handcrossing on the yield and quality of cowpea under semi-arid conditions. The study was conducted at IITA, Ibadan, Nigeria, during the 1997-98 growing season. The experimental design was a randomized complete block design with three replicates. The treatments were: (i) handcrossing, (ii) natural crossing, and (iii) control. The parameters measured were: yield (t/ha), seed yield (t/ha), straw yield (t/ha), protein content (g/kg DM), and digestibility (g/kg DM). The results showed that handcrossing significantly increased yield and seed yield compared to natural crossing and control. The protein content and digestibility were also significantly higher in the handcrossed plants. The authors concluded that handcrossing is a viable method for improving the yield and quality of cowpea under semi-arid conditions.

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