

## Original Research Article

### DIALLEL ANALYSIS OF SWEET POTATO [*IPOMOEA BATATUS* (L.) LAM] GENOTYPES FOR COMBINED BETA CAROTENE AND DRY MATTER CONTENT IN SOUTHERN GUINEA SAVANNA, NIGERIA

#### Abstract

Roots of orange-fleshed sweet potato varieties currently available in Nigeria contain high quantities of  $\beta$ -carotene or pro-vitamin A but have high moisture content. These varieties have been found to be a cheap and crucially important remedy for vitamin A deficiency. The cream or white-fleshed varieties, on the other hand, have a sweet taste with high dry matter content, giving a dry texture, a quality trait preferred in Nigeria. Development of sweet potato genotypes that can combine these two important quality traits is the objective of this breeding work. A diallel experiment using six parental sweet potato genotypes crossed in all possible combinations were carried out and thirty progenies were evaluated for beta carotene ( $\beta$ -carotene) and dry matter content in Landmark University, Omu Aran, Kwara State, Nigeria. The 30  $F_1$  progenies along with their parental lines were planted in the same field trial. The trial was laid out in 6 x 6 triple lattice in two replications. Highly significant ( $P \leq 0.01$ ) differences were observed among the genotypes for the traits. The average  $\beta$ -carotene content among the progenies was 2.86 (mg/100g.f.w) while the dry matter content had a mean value of 31.89%. The cross progenies 199024.2 x Excel had the highest beta carotene (14.37mg/100g.f.w) content with the highest dry matter content (40.10%) and are therefore recommended for further evaluation.

**Key words:** Diallel analysis; dry matter; Southern Guinea Savanna; sweet potato; Vitamin A,  $\beta$ -carotene.

## 27 INTRODUCTION

28 Sweet potato [*Ipomoea batatas* (L.) Lam] is the seventh most important crop in the world  
29 with an estimated 124 million metric tons produced annually. In the tropics, sweetpotato ranks  
30 fifth in terms of caloric contribution after rice, wheat, maize, and cassava [1,2]. In many  
31 developing countries, sweet potato is a staple because they are easy to propagate and maintain  
32 and yield well under a variety of adverse conditions, including drought. The potential of this crop  
33 as a food and a carbohydrate source is widely recognized [3].

34 Sweet potato is one of the most under exploited of the developing world's major crops  
35 [4] as evidenced by its breeding initiatives that are at relatively early stages compared to other  
36 crops. The need to identify local germplasm with desirable traits has long been recognised by  
37 breeders [4]. It has been long known that many sweetpotato traits are mainly quantitatively  
38 inherited [5]. To meet the quality needs there is a need to take into account the farmer and  
39 consumer preferences when developing and selecting sweetpotato varieties and in most cases,  
40 this can be addressed through participatory variety selection. Fortunately, the attributes  
41 considered most important by farmers and consumers were already identified and ranked by [4].  
42 Given the enormous genetic diversity of sweet potato in Uganda [6], the possibility for  
43 sweetpotato improvement to accommodate specific uses is expected to be rapid [4]. There is  
44 wide genetic variability for vitamin A occurring naturally in sweetpotato. This means  
45 conventional breeding techniques can be employed to combine  $\beta$ -carotene and dry matter into  
46 sweetpotato varieties.

47 Diallel mating designs have been widely used in genetic research to investigate the  
48 inheritance of important traits in a set of genotypes [7, 8]. Diallel mating designs were devised,

49 specifically to investigate the combining ability of the parental lines for the purpose of  
50 identification of superior parents for use in hybrid development programmes. A diallel cross is a  
51 set of  $p^2$  possible single crosses and selfs between  $p$  homozygous [9,10,11,12] or heterozygous  
52 [13] parents; it provides a powerful method for investigating the relative genetic properties of  
53 these parents. It is possible to partition treatment variation into components due to general  
54 combining ability (GCA) and specific combining ability (SCA) [14, 7, 16, 17]. General  
55 combining ability (GCA) is the average performance of a genotype in hybrid combination while  
56 Specific combining ability (SCA) are those cases in which certain combinations perform  
57 relatively better or worse than expected on the average [18]. The estimates of the relative  
58 magnitude of the variances of GCA and SCA indicate the type of gene action determining the  
59 traits. Variance due to GCA indicates the predominance of additive gene action while that of  
60 SCA indicates the predominance of non-additive gene action arising largely from dominance and  
61 epistatic deviations [19]. Evaluation of dry matter, starch and beta-carotene content in orange-  
62 fleshed sweet potato (*Ipomoea batatas* L.) genotype tested in three agro-ecological zones of  
63 Malawi has been described by Kathabwalika et al. [20]. Little, however, appears to have been  
64 reported about the relationship between the sensory characteristics and the physical and chemical  
65 constituents, such as carotenoids or dry matter. For example, for consumers in Africa, it has been  
66 reported that they prefer high dry matter varieties of sweet potato [21-23] . The present research  
67 examined the quantitative inheritance of important traits in sweet potato by means of a diallel  
68 analysis with a view to estimating the GCA and SCA components of genetic variance, and to  
69 determine the associated type of gene action controlling  $\beta$ -carotene content and root dry mass.

70

## 71 **2. Materials and Methods**

### 72 **2.1. Description of the Study Area**

73 A field experiment was conducted on six sweet potatoes genotypes (three orange flesh  
74 and three white flesh) at the Teaching and Research Farm of Landmark University, Omu Aran,  
75 Nigeria. The experimental site is located at the Southern Guinea Savanna agro-ecological zone of  
76 Nigeria with distinct wet and dry seasons. The land had been used continuously for the  
77 cultivation of arable crops like maize, melon, cowpea and vegetables for more than three years.  
78 Soil samples were collected from the trial site before cropping and were analyzed in the  
79 laboratory for physical and chemical properties (Table 2). The soil texture was loamy sand.

80

## 81 **2.2. Treatments and Experimental Design**

82 The parent's materials used for the experiment were obtained from the germplasm  
83 collection centre of the Department of Agronomy, the University of Ibadan which was originally  
84 from the listed sources in Table 1. The parents were selected on the basis of being cross-  
85 compatible. Hand crosses were carried out in a 6 x 6 full diallel, excluding selfs from 2010 to  
86 2011 at the Teaching and Research Farm of Landmark University, Omu Aran, Nigeria. Fruits  
87 were harvested between 30-50 days after pollination in the early morning to prevent scattering.  
88 The fruits were further air dried, shelled, put in a labelled envelope and kept in desiccators. The  
89 harvested seeds were soaked in water over night and planted into polythene bags filled with  
90 loamy soil. Once the plants were about 30cm tall, they were transplanted to well-prepared ridges  
91 for further growth and development. Twenty cuttings of a 25cm length of the sweet potato vines  
92 from F<sub>1</sub> progeny were made to represent each cross. The selected 30 F<sub>1</sub> progeny along with their  
93 parental lines were planted in the same field trial. The trial was laid out in 6 x 6 triple lattice in  
94 two replications. The plot size used was 3m x 1m in two rows. Each plot comprised the 20  
95 cuttings from each progeny of a cross. Each vine was inserted at a slant, with two-third buried  
96 below the soil surface. Weeding was done 4, 6 and 8 weeks after planting, using small hoes. No

97 herbicides or fertilizers were applied. Appropriate agronomic practices were followed to raise a  
98 good crop.

### 99 **2.3. Data Collection**

100 All data were recorded on an individual plant basis and then averaged across the 20  
101 progeny of each F<sub>1</sub> cross. The quantitative traits were evaluated as follows:  $\beta$ - carotene content  
102 expressed as mg 100 g<sup>-1</sup> and dry matter content (g) expressed as a percentage of root fresh mass  
103 (g).

### 104 **2.4 Statistical analysis of triple lattice**

105 Data collected on the two traits were subjected to diallel analysis using Griffing (1956)  
106 Method II (parents and crosses together), Mixed I (fixed effects). Both general and specific  
107 combining abilities were computed using [20] for the parent and crosses.

#### 108 Diallel analysis

109 To test the null hypothesis of no genotypic differences among parents and crosses, one-  
110 way analysis of variance was performed. Treatment sum of squares was partitioned into three  
111 components, parents (P), crosses (C), and parent vs. crosses (P. vs. C.). General Combining  
112 Ability and Specific Combining Ability variance components of the cross mean square was  
113 computed according to Griffing's (1956) fixed-effects model I. Reciprocals were defined as  
114 being below the diagonal, adopting [14] notation the following genetic statistical model for  
115 analysis within one environment.

## 116 **3. Results**

### 117 **3.1 Analysis of variance for $\beta$ -carotene and dry matter content**

118  $\beta$ -carotene content and dry matter content means squares were both significant ( $p < 0.01$ ) among  
119 the parents and their 30F<sub>1</sub> families, this shows that there is genetic variation among the parents  
120 and their crosses as shown in table 4. Crosses out-performing their parents can be attributed to  
121 transgressive segregation which is desirable for improving  $\beta$ -carotene content and dry matter

122 content. The results of average performances of some of the crosses presented in table 5 show  
123 that the performances of crosses are significantly higher than the two parents for the traits. Cross  
124 1 x 3 had the highest values in term of  $\beta$ -carotene and dry matter content with means of 14.37 mg  
125  $100\text{ g}^{-1}$  and 40.10% respectively followed by 1 x 4 for  $\beta$ -carotene content with means of 12.39  
126 mg  $100\text{ g}^{-1}$  and dry matter content with a mean of 30.05% while 2 x 4 had the least  $\beta$ -carotene  
127 content and dry matter content with a means square values of  $0.03\text{ mg }100\text{ g}^{-1}$  and 31.15%.

128

### 129 **3.2 General and specific combining ability analysis for $\beta$ -carotene content and dry matter** 130 **content**

131 General combining ability and Specific combining ability mean the sum of squares for  $\beta$ -  
132 carotene content and dry matter content were highly significant ( $p < 0.01$ ) across the parents,  
133 parent x cross and the crosses as presented in Table 6. The mean squares for reciprocals of  $\beta$ -  
134 carotene content were significant ( $p < 0.01$ ) whereas mean squares for the dry matter for the  
135 reciprocal is not significant.

### 137 **3.3 COMBINING ABILITY EFFECTS**

138

#### 139 **3.3.1 Beta-carotene content**

140 Table 7 presented estimates of GCA effects for  $\beta$ -carotene content and dry matter content of six  
141 sweet potato parents. The GCA effects for  $\beta$ -carotene content of parent 1, 2 and 3 were positively  
142 and highly significant ( $p < 0.01$ ). The GCA effects for parent 5 is significant ( $p < 0.01$ ) but  
143 negative. The GCA effect for parent 4 and 6 was negative and was not significant. The SCA  
144 effects of crosses 1 x 2, 1 x 4, 1 x 5, 1 x 6, 2 x 3, 2 x 5, 2 x 6 and 3 x 5 were positive and highly  
145 significant ( $p < 0.01$ ) (Table 8) whereas cross 3 x 6, 4 x 6 is also significant but negative. The rest  
146 of the crosses are positive and not significant, apart from 1 x 3 which was negative and is not  
147 significant ( $p < 0.01$ ). Four reciprocals (5 x 2, 5 x 4, 6 x 5 and 6 x 2) were not significant ( $p < 0.01$ )

148 and negative except cross 6 x 2 which is positive. Crosses 3 x 1 and 3 x 2 were highly significant  
149 although they were negative. The rest crosses were positive and highly significant ( $p < 0.01$ )  
150 (Table 8).

### 151 **3.3.2 Dry Matter Content**

152  
153 The GCA effects for parent 2, 4 and 6 were positively and highly significant ( $p < 0.01$ ). The GCA  
154 effects for parent 1 is also significant ( $p < 0.01$ ) but negative. The GCA effect for parent 5 was not  
155 significant but positive (Table 7). Crosses 1 x 2 and 3 x 5 were positive although not significant  
156 (Table 8). This is against crosses 2 x 3, 3 x 4, 4 x 5 and 4 x 6 which were negative and not  
157 significant ( $p < 0.01$ ). SCA effect for the rest of the crosses was significant ( $p < 0.01$ ) and positive  
158 (Table 8). For reciprocal, crosses 6 x 1 and 6 x 2 are the only crosses that were positively and  
159 highly significant ( $p < 0.01$ ).

## 160 **4 Discussion and conclusion**

### 161 **4.1 General and specific combining ability for $\beta$ -carotene content and dry matter content**

162 Both GCA and SCA variances were significantly (Table 5), this suggests that both additive and  
163 non-additive gene effects played a major role in the inheritance of  $\beta$ -carotene and dry matter  
164 content. The GCA and SCA mean squares for the  $\beta$ -carotene and dry matter content were  
165 significant ( $p < 0.01$ ). This implies that both additive and non-additive gene action were involved  
166 in their expression. This study indicates that additive gene action was relatively more  
167 predominant than non-additive gene action in controlling the expression  $\beta$ -carotene content and  
168 dry matter content. Hence, predicting progeny performance based on GCA for the traits will be  
169 largely successful. The highly significant ( $p < 0.01$ ) reciprocal mean squares for  $\beta$ -carotene and  
170 dry matter content indicates that maternal effects can play a major role in the inheritance of these  
171 traits and consequently the performance of a parent in a cross is dependent on whether it is used  
172 as a female or a male.

173 **4.2  $\beta$ - carotene content**

174 The GCA effects for parent 1 (1.33) and (1.12) were significant ( $p < 0.01$ ) and positive indicating  
175 that additive gene action contributed positively to the expression  $\beta$ -carotene content consequently,  
176 their cross 1 x 2 is positive (3.28) and significant ( $p < 0.01$ ) SCA effect. This means  
177 that the interaction between the parent for the non-additive gene action resulted in the cross  
178 performing above the expectation based on additive effects. The crosses that had positive and  
179 significant ( $p < 0.01$ ) SCA effects were 1 x 4, 1 x 5, 1 x 6, 2 x 3, 2 x 5, 2 x 6 and 3 x 5 indicating  
180 that the non-additive gene action arising from the interaction of the parents contributed positively  
181 to the expression of the trait. Parents 5 that had negative GCA effects (- 0.44) produced a cross  
182 with a positive (0.022) and highly significant ( $p < 0.01$ ) SCA effect (Table 6). This shows that  
183 parents cannot be disqualified solely on the basis of negative GCA effects. In other word, parents  
184 with high positive GCA effects did not necessarily produce crosses with the desired  
185 performance. The parents used in this study, as well as the crosses, generated exhibit different  
186 level of significant and desirable crosses were obtained from crossing parents with high GCA  
187 effects with parents with low GCA effects that is 1 x 5, 2 x 5 and 3 x 5.

188 **4.3 Dry Matter Content**

189 The GCA and SCA mean squares for dry matter content were significant ( $p < 0.01$ ), but the  
190 reciprocal mean square was not significant. For the specific combiners for dry matter content  
191 parent 3 had a positive GCA and their crosses with parent 1 and 2 given dry matter content of  
192 40.01% and 38.20%.

193 **5. Recommendation**

194 It is, therefore, recommend that:



- 195 1. The parent 1 and 3 identified to be good general and specific combiners of  $\beta$ - carotene  
 196 and dry matter content should be further intrigressed into other proven cultivated in the  
 197 improvement of  $\beta$ - carotene and dry matter content in sweet potato.
- 198 2. The identified crosses with the highest dry matter and  $\beta$ - carotene content could be  
 199 incorporated into an on-farm trial for proof.

200  
 201  
 202  
 203  
 204  
 205  
 206

**Table 1: Parental genotypes and their traits used in a 6x6 full diallel excluding selfs**

No	Genotype	Root flesh colour	Root Dry mass (%)	Source
1	199024.2	Orange	31.02	CIP Kenya
2	440034	Orange	26.92	CIP Kenya
3	Excel	Orange	28.53	South Africa
4	W-151	Yellow	34.29	CIP Kenya
5	TIS 87/0087	White	30.67	IITA Ibadan
6	440168	White	32.31	CIP Kenya

207

208 **Table 2. Physical and chemical characteristics of the experimental site soil at Landmark**  
 209 **University, Omu Aran.**

Physical characteristics	properties
Texture	Loamy sand
pH 1:1 (H <sub>2</sub> O)	5.4
Sand %	84.1
Clay %	8.02
Silt %	6.42
<b>Chemical characteristics</b>	

210  
 211

Exchangeable Ca <sup>2+</sup> (C. mol kg <sup>-1</sup> )	1.12
Exchangeable Mg <sup>2+</sup> (C. mol kg <sup>-1</sup> )	1.62
Exchangeable Na <sup>+</sup> (C. mol kg <sup>-1</sup> )	0.19
Exchangeable K <sup>+</sup> (C. mol kg <sup>-1</sup> )	0.01
Total acidity H <sup>+</sup> (C. mol kg <sup>-1</sup> )	0.05
Cation exchange capacity (C. mol kg <sup>-1</sup> )	2.83
% Organic Carbon	0.24
% Soil organic matter	1.03
% Total Nitrogen	0.24
Available Phosphate (mg kg <sup>-1</sup> )	20.31

212  
213  
214  
215  
216  
217  
218  
219  
220  
221  
222  
223  
224  
225

**Table 3 Analysis of variance for Griffing's (1956b) Model I, Method I and the expected mean squares for a full diallel.**

Source	Df	sum of squares	of mean squares	Expected squares	mean	F-ratio
GCA	p-1	S <sub>g</sub>	M <sub>g</sub>	$\delta^2 + 2p[1/p-1]\Sigma g_i^2$		M <sub>g</sub> /M <sub>e</sub>
SCA	p(p-1)/2	S <sub>s</sub>	M <sub>s</sub>	$\delta^2 + 1/p(p-1) \Sigma_i \Sigma_j s_{ij}^2$		M <sub>s</sub> /M <sub>e</sub>
Reciprocal effects	p(p-1)/2	S <sub>v</sub>	M <sub>v</sub>	$\delta^2 + 2[2/p(p-1)] \Sigma_i \Sigma_j r_{ij}^2$		M <sub>r</sub> /M <sub>e</sub>
Error	M	S <sub>e</sub>	M <sub>e</sub>	$\delta^2$		

226  
227  
228  
229  
230  
231

**Table 4: ANOVA for six sweet potato parents and their 30 F<sub>1</sub> families evaluated in a triple lattice design**

Source	Df	Mean squares	
		$\beta$ -carotene content(mg 100 g-1)	Dry Matter content (%)
Rep	1	0.75 <sup>ns</sup>	7.85 <sup>ns</sup>
Treatment	35	38.39**	34.28**
Block within reps	35	14.1	29.60
Intra-block error	70	0.32	5.10
Total	141		

232 \*, \*\* Significant at ( $p < 0.05$ ) and ( $p < 0.01$ ) ( $F$ -probability) respectively; ns=not significant  
233

234

235

236

237

238

239

240

241

242

243 **Table 5: Block corrected means for six sweet potato parents and their diallel evaluated**

Parents/Crosses	$\beta$ -carotene content (mg 100 g <sup>-1</sup> )	Dry Matter content (%)
1 x 2	1.5	33.00
1 x 3	14.37	40.10
1 x 4	12.39	30.5
1 x 5	1.32	31.88
1 x 6	3.37	28.43
2 x 3	5.49	38.67
2 x 4	0.03	31.15
2 x 5	0.03	27.38
2 x 6	1.74	27.27
3 x 4	1.5	29.49
3 x 5	0.12	31.67
3 x 6	1.38	35.71
4 x 5	0.02	37.04
4 x 6	0.03	38.89
5 x 6	1.38	34.15
Reciprocal		
2 x 1	11.03	33.82
3 x 1	4.92	31.86
4 x 1	4.41	29.73
5 x 1	0.12	34.00

6 x 1	1.38	34.78
3 x 2	4.92	25.86
4 x 2	0.13	35.00
5 x 2	1.66	32.69
6 x 2	1.50	34.72
4 x 3	6.12	32.56
5 x 3	4.92	28.30
6 x 3	0.03	33.33
5 x 4	1.38	24.49
6 x 4	1.5	33.94
6 x 5	0.03	27.47
Parent 1	13.38	36.25
Parent 2	0.15	32.00
Parent 3	5.49	26.47
Parent 4	0.00	25.86
Parent 5	0.03	29.73
Parent 6	0.12	28.30
Mean	2.86	31.89
s.e	0.39	5.38
CV (%)	15.1	7.01
LSD <sub>0.05</sub>	0.85	6.04

244

245 **Table 6: Combining ability ANOVA for  $\beta$ -carotene content and dry matter content**

246

Source	Df	Mean squares	
		$\beta$ -carotene content(mg 100 g <sup>-1</sup> )	Dry Matter content (%)
Rep	1	0.65**	7.85 <sup>ns</sup>
Parent	5	31.39**	134.28**
Parent x cross	1	11.1**	29.60**
Crosses	11	62.32**	25.10**
GCA	5	83.98**	54.76**
SCA	6	53.76**	10.80**
Reciprocal	12	54.23**	6.9 <sup>ns</sup>
Error	100	0.032	7.63
Total	141		

247 \*\* Significant at  $p < 0.01$  (by  $F$ -probability); ns=not significant; GCA=variation due to general  
 248 combining ability, SCA=variation due to specific combining ability, reciprocal=variation  
 249 between reciprocal

250

251

252

253 **Table 7 Estimates of GCA effects for  $\beta$ -carotene content and dry matter content of six**  
 254 **sweet potato parents**

Parent	$\beta$ -carotene content	Dry Matter Content
1	1.33**	-3.41**
2	1.12**	4.38**
3	0.50**	- 2.88**
4	-2.13ns	2.05**
5	-0.44**	0.12ns
6	-0.355ns	1.04**

255 \*\* Significant at  $p < 0.01$  (by  $F$ -probability); ns=not significant.  
 256  
 257  
 258  
 259  
 260  
 261  
 262  
 263  
 264  
 265  
 266  
 267  
 268  
 269

270 **Table 8: Estimates of SCA effects for the Diallel analysis for  $\beta$ -carotene content and dry**  
 271 **matter content**  
 272

Crosses	$\beta$ -carotene content (mg 100 g-1)	Dry content (%)	Matter
1 x 2	3.28**	1.64ns	
1 x 3	-2.14ns	3.10**	
1 x 4	5.16**	0.78**	
1 x 5	0.022**	1.88**	
1 x 6	3.37**	3.43**	
2 x 3	5.49**	-8.67ns	
2x 4	0.03ns	1.15**	
2 x 5	4.03**	2.38**	
2x 6	1.74**	2.92**	
3 x 4	1.27ns	-2.49ns	
3 x 5	0.12**	1.67ns	
3 x 6	-1.38**	3.76**	
4 x 5	-0.02ns	-3.04ns	
4 x 6	-7.03**	-3.89ns	
5 x 6	-1.38ns	3.15**	
Reciprocal			

2 x 1	11.03**	-3.82ns
3 x 1	-4.92**	-1.86**
4 x 1	-6.41**	2.73**
5 x 1	-3.12**	3.00ns
6 x 1	1.38**	3.73**
3 x 2	-4.92**	-2.06ns
4 x 2	0.13**	-3.00ns
5 x 2	-1.66ns	2.69Ns
6 x 2	5.50ns	3.72**
4 x 3	6.12**	-3.56
5 x 3	4.92**	2.30ns
6 x 3	0.02**	-3.33
5 x 4	-1.38ns	-2.49
6 x 4	1.5ns	-3.94ns
6 x 5	-5.03ns	2.47ns

\*, \*\* Significant at ( $p < 0.05$ ) and ( $p < 0.01$ ) ( $F$ -probability) respectively; ns=not significant

## References

1. FAO. The global potato economy. Int. Year Potato 2008, Trade and Markets Division, FAO, Rome, Italy. 2008; <http://www.fao.org/potato-2008/en/potato/IYP-3en.pdf>
2. Reddy UK, Bates GT, Ryan-Bohac J, Nimmakayala P. Sweetpotato. In: KOLE, C (ed.) Genome mapping and molecular breeding in plants. New York: Springer. 2007; pp. 237-239.
3. Jarret RL, Gawel N, and Whittemore A. Phylogenetic relationship of sweetpotato [*Ipomoea batatas* (L.) Lam.]. *J. Amer. Soc. Hort. Sci.* 1992; 117: 633-637.
4. Rees D, van Oirschot Q.E.A, Amour R, Rwiza E, Kapinga R, Carey T. Cultivar variation in keeping quality of sweet potatoes. *Postharvest Biol. Technol.*, 2003; 28: 313-325.
5. Jones A, Steinbauer CE, Pope DT. Quantitative inheritance of ten root traits in sweet potatoes. *Journal of the American Society for Horticultural Science* 1969; 94: 271-275.
6. Mukasa, SB, Rubaihayo, PR, Valkonen, JTP. Incidence of viruses and viruslike disease of sweetpotato in Uganda. *Plant Disease* 2003;87:329-335

- 300  
301 7. Collins WP. Analysis of growth in Kennebec with emphasis on the relationship  
302 between stem number and yield. *American Potato Journal* 1977; 54:33-40.  
303  
304 8. Mwanga ROM, Yencho GC, Moyer JW. Diallel analysis of sweetpotato for resistance  
305 to sweetpotato virus disease. *Euphytica* 2002; 128: 237-248.  
306  
307 9. Hayman BI. The analysis of variance of diallel table. *Biometrics* 1954a; 10: 235-244.  
308  
309 10. Hayman BI. The theory and analysis of diallel crosses. *Genetics* 1954b; 39: 789-809.  
310  
311 11. Hayman BI. The theory and analysis of diallel crosses. II. *Genetics* 1958; 43: 63-85.  
312  
313 12. Hayman BI. The theory and analysis of diallel crosses. III. *Genetics* 1960; 45: 155-172.  
314  
315 13. Dickinson AG, Jinks JL. A generalised analysis of diallel crosses. *Genetics* 1956; 41:  
316 65-78.  
317  
318 14. Griffing B. Concept of general and specific combining ability in relation to  
319 diallel crossing systems. *Aust. J. Biol. Sci.*, 1956; 9: 463-493  
320  
321 15. Mihovilovich E, Mendoza HA, Salazar LF. Combining ability for resistance to  
322 sweetpotato feathery mottle virus. *Hort. Science* 2000;35: 1319-1320.  
323  
324 16. Yan W, Hunt LA. Biplot analysis of diallel data. *Crop Science* 2000;42: 21-30.  
325  
326 17. Salami AE, Agbowuro GO. Gene Action and Heritability Estimates of Grain Yield  
327 and Disease Incidence Traits of Low-N Maize (*Zea mays* L.) Inbred lines *Agriculture  
328 And Biology Journal Of North America* 2016;Vol. 7 (2) pg 50-54,  
doi:10.5251/abjna.2016.7.2.50.54  
329  
330 18. Rojas BA, Sprague GF. A comparison of variance components in corn yield trials: III.  
331 General and specific combining ability and their interaction with locations and years.  
332 *Agron. J.* 1952; 44: 462–6.  
333  
334 19. SAS Institute. SAS/STAT user's guide. Version 6, 4th ed. 1995 Vol. I and II. SAS Inst.  
335 Inc. Cary N.C., U.S.A.  
336  
337 20. Kathabwalika, D. M., Chilembwe, E. H. C., & Mwale, V. M. (2016). Evaluation of dry  
338 matter, starch and beta-carotene content in orange-fleshed sweet potato (*Ipomoea*

- 329 *batatas* L.) genotypes tested in three agro-ecological zones of Malawi. *African Journal*  
330 *of Food Science*, 10(11), 320-326.
- 331 21. Tomlins, K., Owori, C., Bechoff, A., Menya, G., & Westby, A. (2012). Relationship  
332 among the carotenoid content, dry matter content and sensory attributes of sweet  
333 potato. *Food Chemistry*, 131(1), 14-21.
- 334 22. Burri, B. J. (2011). Evaluating sweet potato as an intervention food to prevent vitamin  
335 A deficiency. *Comprehensive Reviews in Food Science and Food Safety*, 10, 118–130.
- 336 23. Bechoff, A., Westby, A., Owori, C., Menya, G., Dhuique-Mayer, C., Dufour, D., et al.  
337 (2010). Effect of drying and storage on the degradation of carotenoids in orange-fleshed  
338 sweet potato varieties. *Journal of the Science of Food and Agriculture*, 90, 622–629.
- 339

UNDER PEER REVIEW