

# The effect of using solid-state fermented peeled and unpeeled cassava root tubers and limiting amino acid supplementation on metabolisable energy for meat-type cockerels

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1 **The effect of using solid-state fermented peeled and unpeeled cassava root tubers and**  
2 **limiting amino acid supplementation on metabolisable energy for meat-type cockerels**

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22 **ABSTRACT**

23 A preliminary *in-vitro* solid-state fermentation of peeled (PCRM) and unpeeled cassava root  
24 meal (UCRM) using *Aspergillus niger* was conducted followed by a force-feeding experiment to  
25 investigate the effect of processing, solid-state fermentation and limiting amino acid  
26 supplementation on metabolisable energy (ME) of peeled (PCRM) and unpeeled (UCRM)  
27 cassava root meal for meat-type cockerels. Forty eight, 84 d-old meat-type cockerels (Ross 308)  
28 were assigned to 8 treatments consisting of 6 birds per treatment laid out in a  $2 \times 2 \times 2$  factorial  
29 arrangement of treatment consisting of PCRM and UCRM subjected or not to solid-state  
30 fermentation and supplemented with and without limiting amino acids. Additional 6 cockerels  
31 were also used for endogenous study. Peeling of cassava root increased ( $P < 0.05$ ) gross energy  
32 content of the resultant cassava meal when compared with UCRM. Solid-state fermentation  
33 using *Aspergillus niger* increased ( $P < 0.05$ ) the crude ash, ether extract and arginine  
34 concentration of PCRM and UCRM. Solid-state fermented PCRM recorded the highest ( $P <$   
35  $0.05$ ) amylopectin, least ( $P < 0.05$ ) resistant starch and hydrocyanide concentration. Highest ( $P$   
36  $< 0.05$ ) apparent metabolisable energy (AME) and nitrogen corrected AME (AMEn) values were  
37 obtained for cockerels fed with solid-state fermented PCRM supplemented with or without  
38 amino acid. However, supplementation of solid-state fermented PCRM with amino acid resulted  
39 in highest ( $P < 0.05$ ) true metabolisable energy (TME) and nitrogen corrected TME (TMEn) for  
40 meat-type cockerels. Reduced ( $P < 0.05$ ) AME and AMEn values were recorded for UCRM,  
41 regardless of solid-state fermentation and amino acid supplementation. In conclusion, solid-state  
42 fermentation and amino acid supplementation of PCRM resulted in improved AME, AMEn,  
43 TME and TMEn values for meat-type cockerels. Amino acid supplementation had no  
44 improvement on AME, AMEn and TME values of UCRM for meat-type cockerels.

45 *Keywords:* Amino acid supplementation, Cassava root meal, Cockerels, Metabolisable energy,  
46 Solid-state fermentation

47

## 48 **1. INTRODUCTION**

49 Cassava (*Manihot esculenta*) root is a cheap and sustainable energy feedstuff with potential  
50 to replace most conventional cereal grains in the tropics (Oso et al., 2014). Cassava root is rich in  
51 digestible starch, gross energy content (El-sharkawy, 2012) and has been used to a limited extent  
52 in poultry nutrition (Eruvbetine et al., 2003; Oso et al., 2014). However, the presence of  
53 hydrocyanide (HCN) residues, reduced protein levels, poor protein quality and reduced  
54 concentration of sulphur containing amino acids in cassava root constituted the major constraints  
55 to its maximal utilization as energy feedstuffs in poultry nutrition (Banea-Mayambu et al., 1997).  
56 During cassava processing which convert cyanide to a less toxic thiocyanate, the enzyme  
57 ‘rhodnase’ contained in cassava root utilizes the constituent methionine and other sulphur  
58 containing amino acids as sulfur donor (Cardoso et al., 2005). Thus, sulphur amino acids become  
59 grossly deficient in cassava-based diets fed to poultry birds. Hence, to maximally harness the  
60 rich energy potential of cassava root in poultry nutrition, it is essential to supplement cassava  
61 root based diets with limiting amino acids.

62 Cassava peeling process is the removal of the topmost layer of cassava root prior utilization  
63 as food or feed. This processing methods helps to reduce the resultant hydrocyanide (HCN)  
64 content in cassava root product since the largest concentration of HCN in cassava root is located  
65 on the uppermost layer (Bruijn, 1973). Preliminary study showed improved growth performance

66 of broilers fed diet containing graded levels of peeled cassava root meal when compared with  
67 group fed diet containing unpeeled cassava root meal (Akapo et al., 2014).

68 Solid-state fermentation with fungal culture has been recognized as a means of nutritionally  
69 enriching and detoxifying few cassava products (Oboh and Akindahinsi, 2003). Filamentous  
70 fungi such as *Aspergillus niger* been widely used in food industries for commercial solid-state  
71 fermentation due to its ease of culturing and absence of pathogenic ability (Berka et al., 1992).  
72 *Aspergillus niger* has the capacity to produce extracellular enzymes (such as hemicellulases,  
73 hydrolases, pectinases, protease, amylase and lipases), degrade fibre and enrich its substrate  
74 (Mathivanan et al., 2006; Villena and Gutierrez-Cornea, 2007). The present study seeks to  
75 evaluate the effect of processing, solid-state fermentation and limiting amino acid  
76 supplementation on metabolisable energy of peeled and unpeeled cassava root meal for meat-  
77 type cockerels.

78

## 79 **2. MATERIALS AND METHODS**

### 80 *2.1. Processing of cassava root*

81 Freshly harvested cassava root tubers (TMS 30572) were washed with water and divided into  
82 two equal batches. One batch was manually chipped without prior peeling to obtain the whole  
83 cassava chips (WCC) while the other batch was peeled (removal of 0.5 cm uppermost thick  
84 layer) before chipping to yield the peeled cassava chips (PCC). Both WCC and PCC were dried  
85 (10–11 % moisture content) and milled (2.5 mm sieve) separately to yield the unpeeled (UCRM)  
86 and peeled cassava root meal (PCRM), respectively.

87 *2.2. Solid-state fermentation of cassava root meal*

88 Pure laboratory strain of *Aspergillus niger* (Chinese International Centre for Type Culture  
89 Collection; CICC, No. 41126) was used as inoculum. A total of 8 kg cassava meal (consisting of  
90 4 kg UCRM and 4 kg PCRM) were measured and used for this study. Twenty (20) sub-samples  
91 of UCRM and PCRM, each weighing 200 g were measured and placed into separate conical  
92 flasks. Thus, forty (40) conical flasks were used in all for the study (20 flasks for UCRM and 20  
93 flasks for PCRM group). All UCRM and PCRM samples contained in flasks were randomly  
94 assigned, each into 2 treatments consisting of solid-state fermented and unfermented group. Thus  
95 there were four treatments in all laid out in a  $2 \times 2$  factorial arrangement of peeled (PCRM) and  
96 unpeeled (UCRM) cassava root meal, each subjected or not to solid-state fermentation. Samples  
97 (contained in flasks) subjected to solid-state fermentation were moistened (250 g/kg Moisture  
98 content) each with nutrient solution (containing analytical grade of 80 g urea, 7 g  $\text{MgSO}_4 \cdot 2\text{H}_2\text{O}$ ,  
99 13 g  $\text{KH}_2\text{PO}_4$  and 20 g citric acid) and inoculated with  $2 \times 10^7$  fungal spore of *A. niger* per gram  
100 of sample. Each conical flask was air-sealed and the substrate incubated (30° C) for 6 days in a  
101 bed-packed incubator. At the end of incubation period, fermented samples (contained in each  
102 flask) were sterilized (120° C for 20 min) and used for subsequent chemical analysis.

103 *2.3 Chemical analysis of samples*

104 Fermented samples of UCRM ( $n = 10$ ) and PCRM ( $n = 10$ ) and respective unfermented samples  
105 were analyzed for dry matter (DM) by drying at 80°C for 24 h (AOAC; 925.10). Ash was  
106 measured in a muffle furnace (510° C for 18 h), crude protein ( $6.25 \times \text{N}$ ) was determined by  
107 LECO FP-200 Analyser (St Joseph, MI, USA), oil was extracted with petroleum spirit using the  
108 soxhlet method (AOAC, 1990). Gross energy (Adiabatic bomb calorimeter, Model 1261; Parr

109 Instrument Co., Moline, IL, USA), fibre fraction (Van Soest et al., 1991), tannin (Makkar et al.,  
110 1993) and hydrocyanide content (De Bruijn, 1971) of samples were determined following  
111 standard procedures. The amylopectin (Amylose/Amylopectin kit, Megazyme International Co.  
112 Ireland) and resistant starch content (KRSTAR 08/11 Test kit, Megazyme International Co.  
113 Ireland) of samples were determined using appropriate commercial kits. Mineral analysis (ICP–  
114 MS, Agilent 7500 cx, Agilent Technologies) and amino acid analysis (RP-HPLC; Agilent 1100,  
115 Palo Alto, CA, USA) of the samples were also determined. All laboratory analysis was done at  
116 the Key Laboratory of Agro-ecological Processes in Subtropical Region, Institute of Subtropical  
117 Agriculture, Chinese Academy of Sciences, Hunan Province, China.

#### 118 *2.4 Metabolisable energy determination using gavage method*

119 The experimental protocol used in this study was approved by the Institutional Animal Care and  
120 Welfare Committee of the Institute of Subtropical Agriculture (ISA), Chinese Academy of  
121 Sciences, P.P.R China (Approval No. ISA AEC 2013-014). A total of fifty four (54) meat-type  
122 cockerels (Ross 308, 12-weeks-old) of average weight  $2250\text{g} \pm 115$  were used in all for this  
123 experiment. Forty eight (48) cockerels were assigned to 8 treatments in a  $2 \times 2 \times 2$  factorial  
124 arrangement of treatment consisted of peeled (PCRM) and unpeeled (UCRM) cassava root  
125 meals, fermented or not with *A. niger* and supplemented with and without limiting amino acids.  
126 There were 6 replicates per treatment of 1 bird per replicate. The remaining 6 cockerels were  
127 used for endogenous study. Birds were kept in individual iron-type battery cages (each of  
128 dimension  $35 \times 35 \times 50$ ; LBH) and fed commercial diets prior the commencement of the  
129 experiment. The amino acids supplemented were as follows: L-lysine (0.75 g/100 g cassava  
130 meal), DL-methionine (1.5 0 g/100 g cassava meal), L-arginine (0.75 g/100 g cassava meal) and  
131 L-cysteine (0.75 g/100g cassava meal). Birds were orally gavaged 30 g of respective processed

132 cassava meal after 48 hr of starvation following the standard procedure outlined by Mc Nab and  
 133 Blair (1988). All birds had free access to drinking water while birds assigned to endogenous  
 134 group were dosed each with warm glucose solution (30 g of glucose/50 ml of warm water).  
 135 Excreta voided from each bird following the feeding procedure were collected quantitatively. All  
 136 the birds survived the experiment as no mortality was recorded throughout the study. Gross  
 137 energy of samples of excreta was measured while the following equations were used to calculate  
 138 apparent metabolisable energy (AME), nitrogen corrected apparent metabolisable energy  
 139 (AMEn), true metabolisable energy (TME), and nitrogen corrected true metabolisable energy  
 140 (TMEn) of test ingredient (Sibbald, 1989):

$$141 \quad \text{AME /g of feed} = [(F_i \times \text{GE}_f) - (E \times \text{GE}_e)]/F_i$$

142 Where  $F_i$  is the feed intake (g on dry matter basis),  $E$  is quantity of excreta output (g on dry  
 143 matter basis),  $\text{GE}_f$  is the gross energy (MJ/ kg) of feed, and is  $\text{GE}_e$  the gross energy (MJ/ kg) of  
 144 excreta.

$$145 \quad \text{AMEn /g of feed} = \frac{\{[(F_i \times \text{GE}_f) - (E \times \text{GE}_e)] - (\text{NR} \times 36.5)\}}{F_i}$$

147 where nitrogen retention ( $\text{NR}$ ) =  $(F_i \times N_f) - (E \times N_e)$ ,  $N_f$  is the nitrogen content (g/kg) of  
 148 feed,  $N_e$  is the nitrogen content (g/kg) of excreta.

$$149 \quad \text{TME /g of feed} = \frac{\{[(F_i \times \text{GE}_f) - (E \times \text{GE}_e)] + (\text{FEm} + \text{UE}_e)\}}{F_i}$$



151 where FEm is metabolic faecal energy (kJ) (calculated from gross energy of excreta from  
152 endogenous loss), and UEe is endogenous urinary energy (kJ) (This is assumed zero since urine  
153 and faeces are passed together).

$$154 \quad \text{TME}_n / \text{g of feed} = \frac{\{[(F_i \times \text{GE}_f) - (E \times \text{GE}_e)] - (\text{NR} \times K)\} + \{(F_{\text{Em}} + \text{UE}_e) + (\text{NR}_o \times 36.5)\}}{F_i}$$

155  $F_i$

156 Where NR and NR<sub>o</sub> are estimates of nitrogen retention for fed (experimental) and starved  
157 (control) birds, respectively.

### 158 *2.5 Statistical Analysis*

159 As regards data obtained from compositional chemical analysis of unfermented and solid-state  
160 fermented UCRM and PCRM, replicate units in conical flasks ( $n = 10$  per treatment) served as  
161 experimental units for statistical analysis. These data was analysed as a two factor model  
162 (cassava peeling  $\times$  solid-state fermentation) consisting of peeled and unpeeled cassava root,  
163 subjected or not to solid-state fermentation. For the analysis of data obtained from estimation of  
164 metabolisable energy using gavage method, individual bird was used as the experimental unit ( $n$   
165 = 6 per treatment). Data obtained from gavage studies were analysed as a three factor model  
166 (cassava peeling  $\times$  solid-state fermentation  $\times$  amino acid supplementation) consisting of peeled  
167 and unpeeled cassava root, subjected or not to solid-state fermentation and supplemented with or  
168 without amino acids. All data generated in this study were subjected to analysis of variance using  
169 the general linear models procedure of the SAS (SAS Institute, 2002) to determine the main  
170 effects and their respective interactions. Significant differences were considered at  $P < 0.05$ .

171

### 172 *2.6 Statistical Model*

173 For two factor model (cassava peeling  $\times$  solid-state fermentation) analysis of chemical  
174 composition of peeled and unpeeled cassava root, the model used is as follows:

$$175 Y_{ij} = \mu + A_i + B_j + AB_{ij} + \epsilon_{ijk}$$

176 Where  $Y_{ij}$  = Observed value of the dependent variable

177  $\mu$  = Population mean

178  $A_i$  = Main effect of cassava peeling (peeled, unpeeled)

179  $B_j$  = Main effect of solid state fermentation (fermented, unfermented)

180  $AB_{ij}$  = Interaction effect of cassava peeling and solid state fermentation

181  $\epsilon_{ijk}$  = Random residual error.

182

183 For three factor model (cassava peeling  $\times$  solid-state fermentation  $\times$  amino acid supplementation)  
184 analysis of metabolisable energy determination of peeled and unpeeled cassava root, the model  
185 used is as follows:

$$186 Y_{ijk} = \mu + A_i + B_j + C_k + AB_{ij} + BC_{jk} + AC_{ik} + ABC_{ijk} + \epsilon_{ijkl}$$

187 Where  $Y_{ijk}$  = Observed value of the dependent variable

188  $\mu$  = Population mean

189  $A_i$  = Main effect of cassava peeling

190  $B_j$  = Main effect of solid state fermentation

191  $C_k$  = Main effect of amino acid supplementation

192  $AB_{ij}$  = Interaction effect of cassava peeling and solid state fermentation

193  $BC_{jk}$  = Interaction effect of solid state fermentation and amino acid supplementation

194  $AC_{ik}$  = Interaction effect of cassava peeling and amino acid supplementation

195  $ABC_{ijk}$  = Interaction effect of cassava peeling, solid state fermentation and amino acid  
196 supplementation

197  $\epsilon_{ijkl}$  = Random residual error

198 **3. RESULTS AND DISCUSSION**

199 *3.1. Solid-state fermentation of peeled (PCRM) and unpeeled (UCRM) cassava root meal*

200 Solid-state fermentation of PCRM and UCRM with *A. niger* resulted in increased ( $P <$   
201  $0.05$ ) ether extract, crude ash and reduced ( $P < 0.05$ ) dry matter content (Table 1). Increased  
202 ether extract content of resultant meal (UCRM and PCRM) following solid-state fermentation  
203 could be attributed to the ability of *A. niger* to synthesize long chain fatty acids from acetyl co-  
204 enzymes A and other complex unsaturated lipids during fermentation (Iyayi and Aderolu, 2004).  
205 Increased ash content recorded for fermented UCRM and PCRM when compared with  
206 unfermented meal could be due to increased available mineral caused by metabolic activities of  
207 the fermenting organism. The highest ( $P < 0.05$ ) ash content obtained for fermented UCRM  
208 could be attributed to the rich mineral content of the outer cassava peel contained in UCRM  
209 coupled with the fermentation. The outer layer of cassava root (peel) has been reported to contain  
210 richer macro-minerals than the pulp (Akapo et al., 2014).

211 Peeling of cassava root subjected or not to solid-state fermentation using *A. niger* resulted  
212 in improved (Cassava processing  $\times$  Solid-state fermentation,  $P < 0.05$ ) gross energy content and  
213 reduced ( $P < 0.05$ ) hydrocyanide content (HCN) of the resultant meal when compared with the  
214 unpeeled cassava meal. UCRM contain fibrous outer peels which could lead to a dilution effect  
215 of the constituent energy hence reduced energy content. Cassava root peeling led to reduced  
216 HCN because the highest concentration of HCN in cassava root is located on the outer peel when  
217 compared with the inside pulp (Bruijn 1973). Hence, peeling of cassava root to yield UCRM will  
218 yield a product with reduced HCN content.

219 Solid-state fermentation of PCRM using *A. niger* resulted in a fermented product with  
220 reduced NDF ( $P < 0.01$ ) and ADF ( $P < 0.05$ ) content. Fermentation with *A. niger* thus resulted

221 in efficient breakdown of the constituent fibre. *A. niger* has been earlier reported to produce  
222 ligno-cellulolytic enzymes during fermentation which break down constituent fibre in cassava  
223 root (Mathivanan et al., 2006; Villena and Gutierrez-Cornea, 2007). Solid-state fermentation of  
224 PCRМ in the current study also showed reduced ( $P < 0.05$ ) resistant starch content and improved  
225 ( $P < 0.05$ ) amylopectin content suitable for products that required adhesion (Bergmann et al.,  
226 1988). *A. niger* has been reported to degrade starch granules for substrate enrichment (Soccol et  
227 al., 1994).

228         Solid-state fermentation of both PCRМ and UCRM showed reduced ( $P < 0.05$ ) Cu levels  
229 of the resultant fermented products. In fact, solid-state fermentation of PCRМ resulted in  
230 reduced (Cassava processing  $\times$  Solid-state fermentation,  $P < 0.05$ ) K and Zn content of the  
231 fermented cassava products (Table 1). The effect of solid-state fermentation on mineral profile of  
232 cassava products has not been extensively investigated in literatures. The reduced concentration  
233 of Cu noticed for solid-state fermented PCRМ and UCRM could be due to the adsorption ability  
234 of the fungi. *A. niger* is known to produce large quantities of organic acids such as citrate and  
235 gluconate, both of which are capable of leaching or precipitating metals out of a number of  
236 substrate by either adsorption to fungal cell wall components, or complexation of the metals  
237 (Bosshard et al., 1996).

238         Amino acid profile of PCRМ and UCRM subjected or not to solid-state fermentation is as  
239 shown in Table 2. Solid-state fermentation of PCRМ and UCRM increased ( $P < 0.05$ ) the  
240 arginine concentration of the resultant fermented products. The improved arginine concentration  
241 obtained in fermented PCRМ and UCRM when compared with the unfermented meals  
242 corroborated the earlier findings that fungal fermentation of cassava products improved the

243 resultant amino acid profile (Oboh and Akindahinsi, 2003). Arginine is noted for its role in  
244 protein synthesis and its consequence influence on growth of animals (Kidd et al., 2001).

245

### 246 3.2. Metabolisable energy determination of PCRМ and UCRM using gavage method

247 Metabolisable energy values of PCRМ and UCRM subjected or not to solid-state fermentation  
248 and supplemented with and without amino acids is as shown in Table 3. Solid-state fermentation  
249 of PCRМ supplemented or not with amino acid recorded the highest (Cassava processing ×  
250 Solid-state fermentation × amino acid supplementation,  $P < 0.05$ ) AME and AMEn for meat-  
251 type cockerels. Highest AME and AMEn values of fermented PCRМ recorded in this study  
252 regardless of amino acid supplementation could be due to improved gross energy content and  
253 reduced HCN content of PCRМ following cassava root peeling and solid state-fermentation.  
254 This improved AME and AMEn of fermented PCRМ could also be linked with the increased oil  
255 content produced by *A. niger* during solid-state fermentation (Iyayi and Aderolu, 2004).  
256 Mathivanan et al. (2006) reported that solid-state fermentation produce digestive enzymes which  
257 pre-digest substrates and thus foster increased nutrient availability, digestibility and energy  
258 metabolisability.

259 Reduced (Cassava processing × Solid-state fermentation × amino acid supplementation,  
260  $P < 0.05$ ) AME and AMEn values of UCRM (regardless of solid-state fermentation and amino  
261 acid supplementation) obtained in the present study for meat-type cockerels could be linked with  
262 high fibrous constituent of UCRM. Fibrous feedstuffs have been reported to reduce energy  
263 metabolisability of poultry birds (Janssen and Carré, 1985). Meanwhile, peeling of the outer  
264 layer of cassava root helps in reducing the constituent fibre and thus leads to increased available

265 energy of the resultant product (PCRM). Amino acid supplementation showed no positive  
266 contribution to AME and AMEn values of UCRM from this study.

267         Highest (Cassava processing × Solid-state fermentation × amino acid supplementation, *P*  
268 < 0.05) TME and TMEn values obtained for fermented and amino acid-supplemented PCRM  
269 obtained for meat-type cockerels in the present study underscores the importance of cassava  
270 peeling process, solid-state fermentation and amino acid supplementation in improving the TME  
271 and TMEn values of PCRM. However, amino acid supplementation showed no improvement on  
272 TME and TMEn values of unfermented UCRM. Although, slight improvement on TMEn values  
273 of UCRM was noticed following solid-state fermentation, however these TMEn values were  
274 lower than corresponding values obtained for cockerels fed with fermented and amino acid-  
275 supplemented PCRM.

276

#### 277 **4. CONCLUSION**

278         The present study provides background information on the possible utilization of peeled  
279 and unpeeled cassava root as energy feedstuffs in the nutrition of meat-type cockerels. It was  
280 concluded that solid-state fermentation and amino acid supplementation of peeled cassava root  
281 meal had the best metabolisable energy values (AME, AMEn, TME and TMEn) for meat-type  
282 cockerels. Although solid-state fermentation of unpeeled cassava root meal had little prospect for  
283 improved TMEn, amino acid supplementation of unpeeled cassava root meal had no  
284 improvement on AME and AMEn values for meat-type cockerels.

285

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290

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368 **Table 1.** Effect of solid-state fermentation on the chemical composition and energy content of  
 369 unpeeled and peeled cassava root meal

Cassava root processing (CRP)	Unpeeled		Peeled		Pooled SEM	<i>Level of significance</i>		
	Solid-state fermentation (SSF) No	Yes	No	Yes		CRP	SSF	CRP × SSF
<b>Measurements</b>								
Dry matter (g/kg)	907.1 <sup>a</sup>	719.2 <sup>b</sup>	910.25 <sup>a</sup>	722.50 <sup>b</sup>	40.22	NS	<0.05	<0.05
Crude ash (g/kg)	11.4 <sup>c</sup>	15.1 <sup>a</sup>	10.9 <sup>c</sup>	13.2 <sup>b</sup>	4.00	<0.05	<0.01	<0.05
Ether extract (g/kg)	12.5 <sup>b</sup>	17.5 <sup>a</sup>	12.2 <sup>b</sup>	18.3 <sup>a</sup>	3.99	NS	<0.05	<0.05
Crude protein (g/kg)	14.5	15.0	14.1	15.5	0.12	NS	NS	NS
Gross energy (MJ/kg)	14.10 <sup>b</sup>	14.21 <sup>b</sup>	16.98 <sup>a</sup>	17.25 <sup>a</sup>	4.02	<0.05	NS	<0.05
NDF (g/kg)	360.5 <sup>a</sup>	330.0 <sup>b</sup>	320.7 <sup>c</sup>	305.2 <sup>d</sup>	36.44	<0.01	<0.01	<0.01
ADF (g/kg)	250.2 <sup>a</sup>	227.5 <sup>b</sup>	225.7 <sup>b</sup>	200.7 <sup>c</sup>	32.55	<0.05	<0.05	<0.05
Amylopectin (g/kg)	809 <sup>c</sup>	834.9 <sup>b</sup>	830.9 <sup>b</sup>	874.5 <sup>a</sup>	72.10	<0.05	<0.05	<0.05
Resistant starch (g/kg)	98.50 <sup>a</sup>	48.0 <sup>c</sup>	70.50 <sup>b</sup>	35.0 <sup>d</sup>	12.75	<0.05	<0.05	<0.05
hydrocyanide (mg/kg)	30.4 <sup>a</sup>	30.20 <sup>a</sup>	23.6 <sup>b</sup>	22.50 <sup>b</sup>	4.74	<0.05	NS	<0.05
Tannin (%)	0.32	0.30	0.30	0.29	0.02	NS	NS	NS
Ca (mg/kg)	0.31	0.29	0.30	0.30	0.001	NS	NS	NS
P (mg/kg)	0.52	0.50	0.51	0.54	0.055	NS	NS	NS
Mg (mg/kg)	0.62	0.59	0.60	0.60	0.082	NS	NS	NS
Mn (mg/kg)	0.009	0.008	0.009	0.009	0.0001	NS	NS	NS
Cu (mg/kg)	0.009 <sup>a</sup>	0.003 <sup>b</sup>	0.009 <sup>a</sup>	0.002 <sup>b</sup>	0.0007	NS	<0.05	<0.05
Fe (mg/kg)	0.11	0.10	0.11	0.10	0.004	NS	NS	NS
K (mg/kg)	6.2 <sup>a</sup>	5.70 <sup>a</sup>	6.4 <sup>a</sup>	5.0 <sup>b</sup>	0.92	NS	<0.05	<0.05
Zn (mg/kg)	0.03 <sup>a</sup>	0.02 <sup>a</sup>	0.03 <sup>a</sup>	0.01 <sup>b</sup>	0.007	NS	<0.05	<0.05

370 <sup>a, b</sup> Mean with different superscripts in each row are significantly different (P<0.05)

371 **Table 2.** Effect of solid-state fermentation on amino acid profile of unpeeled and peeled cassava  
 372 root meal

Cassava processing (CRP)	root	Unpeeled		Peeled		Pooled SEM	<i>Level of significance</i>		
		No	Yes	No	Yes		CRP	SSF	CRP × SSF
<i>Measurements (g/100g protein)</i>									
	Asparagine	0.14	0.15	0.15	0.14	0.002	NS	NS	NS
	Threonine	0.04	0.05	0.05	0.04	0.002	NS	NS	NS
	Serine	0.07	0.07	0.08	0.07	0.003	NS	NS	NS
	Glutamine	0.40	0.42	0.41	0.40	0.001	NS	NS	NS
	Glycine	0.10	0.10	0.11	0.10	0.003	NS	NS	NS
	Alanine	0.14	0.14	0.12	0.14	0.005	NS	NS	NS
	Cysteine	0.04	0.05	0.05	0.04	0.010	NS	NS	NS
	Valine	0.09	0.10	0.09	0.10	0.02	NS	NS	NS
	Methionine	0.01	0.02	0.10	0.10	0.002	NS	NS	NS
	Isoleucine	0.05	0.06	0.05	0.06	0.001	NS	NS	NS
	Leucine	0.15	0.15	0.15	0.16	0.001	NS	NS	NS
	Tyrosine	0.04	0.05	0.05	0.05	0.002	NS	NS	NS
	Phenylalanine	0.07	0.06	0.06	0.07	0.001	NS	NS	NS
	Lysine	0.02	0.01	0.01	0.01	0.002	NS	NS	NS
	Histidine	0.03	0.03	0.02	0.02	0.001	NS	NS	NS
	Arginine	0.08 <sup>b</sup>	0.15 <sup>a</sup>	0.09 <sup>b</sup>	0.17 <sup>a</sup>	0.012	NS	<0.05	<0.05
	Proline	0.15	0.16	0.15	0.16	0.004	NS	NS	NS

373 <sup>a, b</sup> Mean with different superscripts in each row are significantly different (P<0.05)

374 NS= Not significant

375 **Table 3.** Metabolisable energy values of peeled and unpeeled cassava root meal subjected to  
 376 solid-state fermentation and supplemented with or without amino acids for meat-type cockerels

Attributes			AME	AMEn	TME	TMEn
Cassava root processing	Solid-state fermentation	Amino acid supplementation				
Unpeeled	No	No	11.62 <sup>b</sup>	11.85 <sup>b</sup>	12.01 <sup>c</sup>	12.33 <sup>c</sup>
Unpeeled	No	Yes	11.70 <sup>b</sup>	11.90 <sup>b</sup>	12.15 <sup>c</sup>	12.44 <sup>c</sup>
Unpeeled	Yes	No	11.92 <sup>b</sup>	12.20 <sup>b</sup>	12.42 <sup>bc</sup>	12.64 <sup>b</sup>
Unpeeled	Yes	Yes	11.99 <sup>b</sup>	12.35 <sup>b</sup>	12.50 <sup>bc</sup>	12.80 <sup>b</sup>
Peeled	No	No	12.22 <sup>b</sup>	12.32 <sup>b</sup>	12.51 <sup>bc</sup>	12.59 <sup>b</sup>
Peeled	No	Yes	12.10 <sup>b</sup>	12.15 <sup>b</sup>	12.49 <sup>bc</sup>	12.60 <sup>b</sup>
Peeled	Yes	No	12.85 <sup>a</sup>	12.90 <sup>a</sup>	13.29 <sup>b</sup>	13.34 <sup>b</sup>
Peeled	Yes	Yes	12.75 <sup>a</sup>	12.84 <sup>a</sup>	13.67 <sup>a</sup>	13.76 <sup>a</sup>
Pooled SEM			2.22	2.07	2.10	2.05
<b>Significance</b>						
Cassava root peeling			NS	<0.05	<0.05	NS
Solid state fermentation			NS	<0.05	<0.05	NS
Amino acid supplementation			NS	NS	NS	NS
Cassava root peeling × Solid state fermentation			<0.05	<0.05	<0.05	<0.01
Cassava root peeling × Amino acid supplementation			NS	<0.05	<0.05	<0.05
Solid state fermentation × Amino acid supplementation			NS	<0.05	<0.01	<0.05
Cassava root peeling × Solid state fermentation × Amino acid supplementation			<0.05	<0.05	<0.05	<0.05

377 <sup>a,b,c,d</sup> Values in the same column not sharing a common superscript are significantly different at  $P$

378  $<0.05$ .

379 NS= Not significant