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3**JAST (Journal of Animal Science and Technology) TITLE PAGE**

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<b>ARTICLE INFORMATION</b>	<b>Fill in information in each box below</b>
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<b>Article Title (within 20 words without abbreviations)</b>	A reduction in dietary crude protein with amino acid balance has no negative effects in pigs
<b>Running Title (within 10 words)</b>	Protein and amino acid level in pigs
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6

## Abstract

The aim of this experiment was to evaluate the effects of low crude protein (CP) level with essential amino acids (AA) addition on growth performance, nutrient digestibility, microbiota, and volatile fatty acid composition in growing pigs. A total of 160 growing pigs (LYD; average initial body weight  $16.68 \pm 0.12$  kg) were randomly allotted to one of the four treatments on the basis of initial BW. A randomized complete block design was used to conduct this experiment in the Research Center of Animal Life Sciences at Kangwon National University. There were ten pigs/replicate with four replicates in each treatment. The treatments include; CON (Control, 17.2% dietary CP level), low protein (LP)-1.10 (15.7% dietary CP level + 1.10% lysine level), LP-1.15 (15.7% dietary CP level + 1.15% lysine level), LP1.2 (15.7% dietary CP level + 1.20% lysine level). The pigs fed CON and LP-1.2 diet showed greater final body weight than that of LP-1.1 diet ( $p < 0.05$ ). Although average daily gain, average daily feed intake, and feed efficiency did not show any difference in phase 2 and 3, average daily gain and feed efficiency was significantly greater in CON and LP-1.20 in phase 1. However, the average daily feed intake did not show any difference during the experimental period. Isobutyric acid and isovaleric acid composition of LP treatments were lower than CON treatment in phase 2. Total branched chain fatty acid composition was significantly lower in LP treatment in phases 1 and 2. However, there was no significant difference among treatments in phase 3. The results of this study underscore the importance of AA supplementation when implementing a low-protein diet during the early growth phase (16-50 kg) in pigs.

**Keywords:** pig, crude protein, amino acid, growth performance, volatile fatty acid

## Introduction

Reducing crude protein (CP) might be effective to mitigate environmental and economic problems in swine production. Swine producers are able to lower the level of dietary CP when the diets satisfy the pig requirement for total nitrogen and essential amino acids (AA) [1]. Soybean meal is commonly added to corn-soybean meal feed to increase the lysine content, as corn contains lower levels of lysine [2]. However, excessive protein intake can lead to undigested AA and nitrogen being excreted in feces, resulting in decreased nitrogen utilization and protein fermentation in the hindgut, which can negatively impact intestinal health. Therefore, there is a growing trend towards reducing dietary CP levels and supplementing synthetic AA to meet the pig's nutritional needs [3-6].

Recent studies have shown that reducing dietary CP levels by 4% with limited amino acids, such as lysine, tryptophan, threonine, and methionine, does not affect the growth performance of growing and finishing pigs [7-9]. However, other studies have shown that reducing dietary CP levels by 4% with limited amino acid supplementation can have a negative impact on growth performance, particularly in younger pigs [10]. Additionally, reducing dietary CP levels by 5% can impair the growth performance of growing pigs due to essential AA deficits [11]. Dietary CP levels have also been found to affect microbial communities, with low levels decreasing pathogen activity in the intestine without affecting beneficial bacteria. Moreover, reducing dietary CP levels and supplementing synthetic AA may decrease odor emission by reducing branched-chain volatile fatty acid metabolism in the hindgut [14]. In light of these findings, this study aimed to evaluate the effects of low CP levels with essential AA supplementation on the growth performance, nutrient digestibility, microbiota, and volatile fatty acid (VFA) composition of growing pigs.

## Materials and Methods

### Animals and Experimental Design

A total of 160 growing pigs (LYD; average initial body weight  $16.68 \pm 0.12$  kg) were randomly allotted to one of the four treatments on the basis of initial BW. A randomized complete block design was used to conduct this study in the Research Center of Animal Life Sciences at Kangwon National University. There were ten pigs per replicate, with four replicates for each treatment. The treatments

62 were CON (17.2% dietary CP level), LP-1.10 (15.7% dietary CP level + 1.10% Lys level), LP-1.15  
63 (15.7% dietary CP level + 1.15% Lys level), LP-1.20 (15.7% dietary CP level + 1.20% Lys level).  
64 The experimental diets were supplemented for 52 days in three phases; phase 1 (day 0–14), phase 2  
65 (day 15–28), phase 3 (day 29–42). The pigs were grouped in partially slatted concrete floor pens 2.80  
66 m × 5.00 m in size. All pens contained a self-feeder and nipple drinker to allow *ad libitum* access to  
67 feed and water. The diets were formulated to provide all the nutrients that met or exceeded the  
68 nutrient requirements listed in the NRC [15], with the exception of Ca (Table 1).

69

### 70 **Growth Performance**

71 The body weights of all the pigs were measured at the end of each phase. The amount of feed  
72 supplemented was measured throughout the experimental period to calculate the average daily feed  
73 intake. The average daily gain (ADG), average daily feed intake (ADFI), and gain-to-feed ratio (G/F)  
74 were calculated at the end of each phase.

75

### 76 **Nutrient Digestibility**

77 The effects of dietary CP and AA supplementation on nutrient digestibility were determined as  
78 follows: pigs were fed a diet containing 2.5 g Cr<sub>2</sub>O<sub>3</sub>/kg for seven days before sampling, and fecal  
79 samples were collected for four days before sampling. In this trial, we evaluated dry matter (DM),  
80 gross energy (GE), and CP digestibility. Prior to fecal sample collection, the floor was cleaned to  
81 avoid contamination, and fecal samples were retrieved and placed in vacuum-sealed plastic bags.  
82 Fecal samples were stored in a freezer at –20°C to preserve the state until analyzed. Samples were  
83 thawed, dried at 60 °C for 72 h in a forced-air oven, grounded in a 1-mm screen Wiley mill (Thomas  
84 Model 4 Wiley Mill; Thomas Scientific, Swedesboro, NJ, USA), and analyzed to calculate  
85 digestibility. Each fecal sample was analyzed in quadruplicate for DM (Method 930.15), CP (Method  
86 990.03) according to AOAC methodology [16]. A bomb calorimeter (Model 1261; Parr Instrument  
87 Co., Moline, IL, USA) was used to analyze gross energy.

88

### 89 **Fecal Microflora DNA**

90 At the end of each phase, pigs were selected based on their average body weight for each treatment,  
91 and samples were collected via gentle rectal massage. The samples were immediately kept in liquid  
92 nitrogen and moved to a deep freezer at  $-80^{\circ}\text{C}$  until analysis. DNA was extracted using the QIAamp  
93 Fast DNA Stool Mini Kit (cat. No. 51604 2016; QIAGEN, Hilden, Germany). The fecal sample (200  
94 mg) was weighed in a 2 mL centrifuge tube and kept on ice. To ensure the highest possible DNA  
95 concentration in the final eluate, 1 mL of InhibitEX Buffer was added to each sample and vortexed  
96 continuously for 1 min until the sample was thoroughly homogenized. The samples were incubated in  
97 a  $70^{\circ}\text{C}$  water bath for 5 min and vortexed for 15 s to achieve consistent lysis. The samples were then  
98 centrifuged at  $20,000 \times g$  and 14,000 rpm for 1 min to pellet the feces. Secondly, 25  $\mu\text{L}$  of proteinase  
99 K and 600  $\mu\text{L}$  of the first sample's supernatant were combined in a fresh 2 mL centrifuge tube. Next,  
100 600  $\mu\text{L}$  of Buffer AL was added and vortexed for 15 s to create a homogeneous solution that was  
101 incubated for 10 min at  $70^{\circ}\text{C}$  and centrifuged briefly to eliminate drops inside the tube lid. The lysate  
102 was mixed with 600  $\mu\text{L}$  of ethanol (96%) and vortexed. In the QIAamp spin column, the lysate (600  
103  $\mu\text{L}$ ) was added to a QIAamp spin column and centrifuged at  $20,000 \times g$  and 14,000 rpm for 1 min.  
104 The QIAamp spin column was moved to a new collection tube, and the former tube was removed.  
105 QIAamp spin column was carefully opened, and 500  $\mu\text{L}$  of Buffer AW1 was added and centrifuged at  
106  $20,000 \times g$  and 14,000 rpm for 1 min. The QIAamp spin column was stored in a new collection tube.  
107 Subsequently, 500  $\mu\text{L}$  of Buffer AW1 was added to the QIAamp spin column and centrifuged for 3  
108 min at  $20,000 \times g$  and 14,000 rpm. Finally, the QIAamp spin column was moved into a new 2 mL  
109 centrifuge tube, and 200  $\mu\text{L}$  of Buffer ATE was mixed directly onto the QIAamp membrane, kept for  
110 1 min at room temperature, then centrifuged at  $20,000 \times g$  and 14,000 rpm for 1 min to elute the DNA,  
111 which was then later quantified using a spectrophotometer. The levels of fecal microflora, such as  
112 *Lactobacillus* spp., *Bifidobacterium* spp., *Clostridium* spp., and *E. coli*., were estimated using the  
113 methodology of Habeeb et al. [17].

114

### 115 **Volatile fatty acids**

116 Samples from pigs that were chosen based on the average body weight of each pen to minimize  
117 errors were collected (d 14, 28, and 42) directly through rectal massage to estimate VFA  
118 concentrations in feces. Fecal samples were immediately stored in collection tubes and placed on ice.

119 VFA concentrations in the feces were estimated using gas chromatography (HP 6890 Plus; Hewlett  
120 Packard, Houston, TX, USA) according to the method of Jeon et al. [18].

121

## 122 **Statistical Analysis**

123 The collected data from this experiment were analyzed using the Analysis of Variance (ANOVA),  
124 implemented through the General Linear Model (GLM) procedure of SAS (version 9.2; SAS Institute  
125 Inc., Cary, NC). For assessing growth performance, the initial BW was employed as a covariate, but  
126 was omitted from the model if it proved insignificant. Each pig served as an experimental unit for  
127 parameters such as growth performance, feed consumption, nutrient digestibility, blood electrolyte  
128 equilibrium, and bone measurements. The Tukey mean comparison test was utilized for treatment  
129 mean separation, with a significance level set at  $p < 0.05$ . Any probability below 0.1 was  
130 recognized as a trend.

131

## 132 **Results**

### 133 **Growth performance**

134 The effects of dietary CP and AA levels on growth performance are shown in Table 2. Pigs fed the control and  
135 LP-1.20 diet showed greater final body weight than those fed the LP-1.1 diet. Although ADG, ADFI, and G/F  
136 did not differ in phases 2 and 3, ADG and G/F were significantly greater in the CON and LP-1.20 in phase 1.  
137 However, ADFI showed no difference during the experimental period.

138

### 139 **Nutrient digestibility**

140 The effects of dietary CP and AA levels on nutrient digestibility are presented in Table 3. DM, ME, and CP  
141 digestibility were calculated to evaluate the effects of crude protein and amino acid level in the diet. In phase 1,  
142 CP digestibility was higher in the LP-1.2 than LP-1.1. However, there was no significant difference between  
143 treatments. In this study, DM, ME, and CP digestibility showed no significant differences among the treatments  
144 in the phase 2 and 3.

145

### 146 **Microflora**

147 The effects of dietary CP and AA levels on the microflora quantity are shown in Table 4. The content of  
148 *lactobacillus* spp., *bifidobacterium* spp., *E.coli*, *Clostridium* in the feces were analyzed to evaluate the effects  
149 of crude protein and amino acid level in the diet. Although *lactobacillus* spp. was increased and *clostridium*  
150 was decreased in the LP-1.2 than LP-1.1 in phase 1, there was no significant difference among the treatments.  
151 In the phase 2 and 3, the content of microflora didn't show difference among the various treatments.

152

### 153 **Volatile fatty acids**

154 The effect of dietary CP and AA level on VFA is shown in Table 5. Isobutyric acid and isovaleric acid  
155 compositions of LP treatments were lower than those of CON in phase 2. The total branched-chain fatty acid  
156 composition was significantly lower in the LP treatment in phases 1 and 2. However, there were no significant  
157 differences among the treatments in phase 3.

158

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161

## 162 **Discussion**

163 The present study investigated the effects of varying levels of dietary CP and AA supplementation  
164 on the growth performance, nutrient digestibility, microbiota, and VFA concentrations in growing  
165 pigs. The results indicate that reducing the CP level by 1.5% with a balanced supply of essential AA  
166 did not significantly affect the growth performance of pigs, but it did lead to a significant reduction in  
167 some branched-chain fatty acids. According to Kerr et al. [3], a low-CP diet, reduced by 2-4%, with  
168 the addition of a limited amount of AA, such as lysine, threonine, methionine, and tryptophan, is a  
169 viable option for pigs. However, an excessive decrease in dietary CP may hinder pig growth  
170 performance [19-21]. In our study, pigs that consumed low-protein diets and received additional  
171 essential AA, such as Lys, Thr, and Met, demonstrated growth performance comparable to that of the  
172 CON group during the trial period.

173 This study found that pigs fed the LP diet had poorer ADG and feed efficiency in phase 1 compared  
174 to those fed the CON diet, but no significant differences were observed in phases 2 and 3. These  
175 results suggest that pigs require a higher nitrogen intake for protein deposition, and the requirements  
176 for the first five essential AA are less well-defined in the early growth phase compared to the later  
177 phases. This finding is consistent with previous studies that have shown pigs to be more sensitive to  
178 dietary CP levels during the growing phase than the finishing phase [22,23]. However, nitrogen  
179 retention was observed during the finishing phase [24]. The LP diet contained a higher proportion of  
180 corn than the CON diet, resulting in an increased availability of starch, which may explain the  
181 compensatory growth observed in the finishing phase. Starch is known to be more efficient for fat  
182 deposition than protein [25]. No significant differences in growth performance were observed  
183 between the dietary treatments in each growth phase. However, it is possible that the nutrient supply,  
184 particularly non-essential AA in LP diets, may be insufficient for protein deposition in rapidly  
185 growing pigs, resulting in a lower growth rate compared to pigs fed high-protein diets [26].

186 Although this study reduced the CP concentration in pig feed by decreasing the soybean meal  
187 content, nutrient digestibility was not affected when dietary CP was decreased to 1.5%. The main  
188 factors affecting protein digestibility are the levels and balance of essential AA and animal  
189 requirements [8]. In this study, the levels (%) of Lys, Met, and Thr were similar for different CP  
190 treatments, indicating that the levels of limited AAs were not affected by reducing CP levels. The

191 digestibility of CP remained unchanged despite the reduction in dietary CP levels. When the limiting  
192 AA levels are constant, a decrease in CP levels can potentially lead to a more balanced AA  
193 composition compared to elevated CP levels. Ball [27] discovered a reduction in energy digestibility  
194 as CP levels decreased in diets featuring 6.9 g/kg of readily available lysine. Zervas and Zijlstra [28]  
195 echoed this finding, attributing it to the diminished fiber content in diets rich in protein [29]. The  
196 influence of CP level on nutrient digestibility warrants consideration since a decrease in digestibility  
197 could lower the slurry DM concentration, subsequently resulting in a surge in slurry output [30].

198 Portune et al. [31] showed a significant correlation between gut microbiota and the metabolism of  
199 dietary proteins. Undigested proteins in the gut provide nitrogen for saccharolytic bacterial growth  
200 and AA for fermentation by asaccharolytic species. The mammalian intestine harbors a multitude of  
201 microbial strains, numbering over  $10^{14}$  microbial cells. These microorganisms play pivotal roles in the  
202 host's physiology and metabolism. The fermentation process of undigested dietary proteins can foster  
203 the growth of protein-fermenting bacteria, thereby suggesting that the origin, quality, and volume of  
204 dietary protein can have a bearing on microbial communities. Research indicates that the level of  
205 dietary CP exerts a more profound effect on the composition of gut microbiota compared to its origin  
206 [32]. In the case of weaned piglets, a decrease in dietary CP led to a reduction in the *Clostridium*  
207 count, however, it didn't affect the total bacteria, *Lactobacilli*, *Enterobacteria*, and *Bacteroides*.  
208 Nonetheless, alterations in the dietary CP content didn't significantly impact bacterial communities in  
209 any part of the intestine under normal physiological circumstances, as the microbiota possesses a  
210 certain degree of adaptability. The outcomes of studies examining the influence of dietary CP levels  
211 on the microbiota composition are inconsistent, possibly due to the limitations of conventional  
212 culture-dependent or low-throughput culture-independent methodologies.

213 Canh et al. [33] stated that fermentable non-starch polysaccharides are the primary dietary  
214 components that affect the VFA concentrations in manure. Most VFAs in manure consist of short  
215 straight-chain VFAs such as acetic, propionic, and butyric acids, which account for 91% of the total  
216 VFA content. This was consistent with the findings of Otto et al. [34] and Le et al. [35]. They  
217 proposed that branched-chain VFAs are only produced during protein metabolism, which could  
218 explain the slight increase in isobutyric and isopentanoic acid concentrations in manure as dietary CP  
219 levels increased from 12% to 18%, although these changes were not statistically significant.

220 In summary, a reduction of 1.5% in protein levels by 1.5% with a balance of essential AA did not  
221 significantly affect on the growth of pigs; however, it did result in a significant reduction in some  
222 branched-chain fatty acids. In the case of lower protein diets, supplementation with AA balance led to  
223 increased body weight gain during the 16–50 kg phase, whereas not supplementing with AA during  
224 the same phases led to reduced growth performance. In this study, the nutrient digestibility and  
225 microbiota of pigs fed diets with different levels of CP were not affected. These results imply that a  
226 low-protein diet may be a viable choice when the AA composition is well-balanced.

227

228 **Acknowledgments**

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## Tables and Figures

Table 1. Ingredient and calculated composition of experimental diets (as-fed diets).

CP, %	17.2		15.7	
Lysine, %	1.10	1.10	1.15	1.20
Ingredients composition, %				
Corn	56.72	60.55	60.52	60.51
Bakery by product	5.00	5.00	5.00	5.00
Molasses	2.00	2.00	2.00	2.00
Soybean meal	22.75	18.40	18.09	17.81
DDGS	7.00	7.00	7.00	7.00
Animal fat	3.65	3.68	3.71	3.74
Salt	0.40	0.40	0.40	0.40
TCP	0.90	0.94	0.95	0.95
Limestone	0.62	0.60	0.60	0.60
Lysine (78%)	0.34	0.49	0.56	0.64
Tryptophan (100%)	0.15	0.37	0.50	0.61
Threonine (98.5%)	0.09	0.15	0.20	0.24
Methionine (99.5%)	0.08	0.12	0.16	0.20
Choline chloride	0.05	0.05	0.05	0.05
Mineral premix <sup>1</sup>	0.10	0.10	0.10	0.10
Vitamin premix <sup>2</sup>	0.10	0.10	0.10	0.10
Phytase	0.05	0.05	0.05	0.05
Total	100.00	100.00	100.00	100.00
Calculated composition, %				
ME, kcal/kg	3,400	3,400	3,400	3,400
CP	17.20	15.70	15.70	15.70
EE	7.01	7.13	7.16	7.19
Lysine	1.10	1.10	1.15	1.20
Methionine + cysteine	0.59	0.59	0.63	0.66
Threonine	0.63	0.63	0.67	0.70
Tryptophan	0.18	0.18	0.19	0.20
Valine	0.73	0.66	0.65	0.65
Ca	0.63	0.62	0.62	0.62
P	0.59	0.58	0.58	0.58

<sup>1</sup>Supplied per kilogram of diet: 45 mg Fe, 0.25 mg Co, 50 mg Cu, 15 mg Mn, 25 mg Zn, 0.35 mg I, 0.13 mg Se.

<sup>2</sup>Supplied per kilogram of diet: 16,000 IU vitamin A, 3,000 IU vitamin D<sub>3</sub>, 40 IU vitamin E, 5.0 mg vitamin K<sub>3</sub>, 5.0 mg vitamin B<sub>1</sub>, 20 mg vitamin B<sub>2</sub>, 4 mg vitamin B<sub>6</sub>, 0.08 mg vitamin B<sub>12</sub>, 40 mg pantothenic acid, 75 mg niacin, 0.15 mg biotin, 0.65 mg folic acid.

Table 2. The effects of CP and AA level on growth performance in growing pigs.

CP, %	17.2	15.7			SEM	p-value
Lysine, %	1.10	1.10	1.15	1.20		
BW, kg						
Initial	16.78	16.71	16.69	16.69	0.11	0.827
Final	51.43 <sup>a</sup>	48.72 <sup>c</sup>	49.84 <sup>bc</sup>	50.75 <sup>ab</sup>	0.51	0.001
Phase 1 (d 0-14)						
ADG, kg	812 <sup>a</sup>	638 <sup>b</sup>	707 <sup>ab</sup>	763 <sup>a</sup>	38.31	0.004
ADFI, kg	1,552	1,433	1,488	1,452	64.75	0.316
G/F	0.524 <sup>a</sup>	0.445 <sup>b</sup>	0.475 <sup>ab</sup>	0.526 <sup>a</sup>	0.02	0.012
Phase 2 (d 15-28)						
ADG, kg	827	816	825	831	47.53	0.991
ADFI, kg	1,656	1,653	1,665	1,657	50.28	0.995
G/F	0.499	0.493	0.494	0.502	0.02	0.953
Phase 3 (d 28-42)						
ADG, kg	837	832	836	839	23.48	0.993
ADFI, kg	1,634	1,611	1,616	1,631	15.83	0.414
G/F	0.512	0.517	0.517	0.514	0.02	0.987
Overall (d 0-42)						
ADG, kg	825 <sup>a</sup>	762 <sup>c</sup>	789 <sup>bc</sup>	811 <sup>ab</sup>	10.64	<0.001
ADFI, kg	1,614	1,566	1,590	1,580	23.06	0.251
G/F	0.511 <sup>a</sup>	0.487 <sup>b</sup>	0.497 <sup>ab</sup>	0.513 <sup>a</sup>	0.01	0.010

CP, crude protein; AA, amino acid; SEM, standard error of means; BW, body weight; ADG, average daily gain; ADFI, average daily feed intake; G/F, feed efficiency.

<sup>a-b</sup>means different superscript letters indicate significant differences (p<0.05).

Table 3. The effects of CP and AA level on nutrient digestibility in growing pigs.

CP, %	17.2	15.7			SEM	p-value
Lysine, %	1.10	1.10	1.15	1.20		
Phase 1 (d 14)						
DM	84.50	85.00	83.98	84.85	0.85	0.652
ME	80.22	79.71	79.12	79.50	0.71	0.494
CP	80.07	79.51	78.44	80.48	1.41	0.524
Phase 2 (d 28)						
DM	84.36	84.24	83.39	84.21	0.65	0.443
ME	79.24	79.52	78.47	79.36	1.12	0.794
CP	79.48	78.21	79.17	79.29	1.64	0.867
Phase 3 (d 42)						
DM	83.98	83.52	83.07	83.79	0.69	0.589
ME	78.78	78.30	78.04	79.01	1.08	0.801
CP	78.42	78.39	78.10	78.40	1.54	0.996

CP, crude protein; AA, amino acid; SEM, standard error of means; DM, dry matter; ME, metabolizable energy; CP, crude protein.

Table 4. The effects of CP and AA level on microbiota in growing pigs.

CP, %	17.2	15.7			SEM	p-value
Lysine, %	1.10	1.10	1.15	1.20		
Phase 1 (d 14)						
<i>Lactobacillus</i> spp.	1.247	1.241	1.258	1.271	0.02	0.412
<i>Bifidobacterium</i> spp.	0.774	0.819	0.774	0.787	0.02	0.139
<i>E.coli</i>	0.320	0.283	0.292	0.302	0.02	0.217
<i>Clostridium</i>	0.539	0.505	0.498	0.482	0.03	0.204
Phase 2 (d 28)						
<i>Lactobacillus</i> spp.	1.236	1.271	1.291	1.301	0.03	0.244
<i>Bifidobacterium</i> spp.	0.795	0.806	0.826	0.817	0.01	0.212
<i>E.coli</i>	0.323	0.306	0.308	0.293	0.02	0.357
<i>Clostridium</i>	0.515	0.500	0.498	0.494	0.02	0.619
Phase 3 (d 42)						
<i>Lactobacillus</i> spp.	1.281	1.318	1.281	1.301	0.03	0.624
<i>Bifidobacterium</i> spp.	0.791	0.798	0.782	0.792	0.02	0.929
<i>E.coli</i>	0.278	0.288	0.306	0.298	0.02	0.361
<i>Clostridium</i>	0.508	0.504	0.510	0.490	0.03	0.884

CP, crude protein; AA, amino acid; SEM, standard error of means.

Table 5. The effects of CP and AA level on volatile fatty acid in growing pigs (g/kg)

CP, %	17.2	15.7			SEM	p-value
Lysine, %	1.10	1.10	1.15	1.20		
Phase 1 (d 14)						
Acetic acid	4.15	4.26	4.29	4.26	0.14	0.799
Propionic acid	1.46	1.45	1.50	1.51	0.04	0.485
Butyric acid	1.14	1.15	1.11	1.16	0.03	0.336
Isobutyric acid	0.68	0.61	0.62	0.64	0.03	0.166
Isovaleric acid	0.87	0.76	0.77	0.80	0.04	0.059
Total SCFA	6.75	6.86	6.90	6.92	0.17	0.724
Total BCFA	1.55 <sup>a</sup>	1.37 <sup>b</sup>	1.39 <sup>b</sup>	1.44 <sup>ab</sup>	0.05	0.009
Total VFA	8.29	8.23	8.29	8.36	0.17	0.880
Phase 2 (d 28)						
Acetic acid	4.27	4.22	4.22	4.39	0.13	0.537
Propionic acid	1.54	1.47	1.50	1.49	0.04	0.430
Butyric acid	1.17	1.10	1.11	1.08	0.07	0.602
Isobutyric acid	0.54 <sup>a</sup>	0.39 <sup>b</sup>	0.42 <sup>b</sup>	0.39 <sup>b</sup>	0.04	<0.007
Isovaleric acid	0.84 <sup>a</sup>	0.60 <sup>b</sup>	0.61 <sup>b</sup>	0.62 <sup>b</sup>	0.04	<0.001
Total SCFA	6.98	6.79	6.84	6.96	0.16	0.584
Total BCFA	1.39 <sup>a</sup>	0.99 <sup>b</sup>	1.04 <sup>b</sup>	1.00 <sup>b</sup>	0.05	<0.001
Total VFA	8.37	7.78	7.88	7.97	0.17	0.880
Phase 3 (d 42)						
Acetic acid	4.10	4.33	4.18	4.12	0.11	0.175
Propionic acid	1.48	1.49	1.50	1.48	0.05	0.977
Butyric acid	1.10	1.13	1.08	1.13	0.05	0.703
Isobutyric acid	0.55	0.53	0.53	0.52	0.08	0.984
Isovaleric acid	0.70	0.74	0.70	0.73	0.06	0.900
Total SCFA	6.68	6.95	6.76	6.72	0.15	0.316
Total BCFA	1.25	1.27	1.23	1.24	0.13	0.994
Total VFA	7.93	8.22	7.99	7.97	0.19	0.434

CP, crude protein; AA, amino acid; SEM, standard error of means; SCFA, short chain fatty acid; BCFA, branched chain fatty acid; VFA, volatile fatty acid.

<sup>a-b</sup>means different superscript letters indicate significant differences (p<0.05).