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| <b>Article Type</b>  | Research article  |
| <b>Article Title (within 20 words without abbreviations)</b>   | Effects of emulsifier and $\beta$ -mannanase supplementation on growth performance and nutrient utilization in grower pigs fed diets with different energy levels   |
| <b>Running Title (within 10 words)</b>   | Emulsifier and $\beta$ -Mannanase Supplementation in Grower Pigs  |
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## Abstract

This study aimed to evaluate the effects of dietary energy level and supplementation with  $\beta$ -mannanase ( $\beta$ M) and an emulsifier (EM) on growth performance, nutrient digestibility, and fecal bacterial populations in grower pigs. A total of 192 pigs (Landrace  $\times$  Yorkshire  $\times$  Duroc) with an average initial body weight of  $22.78 \pm 1.06$  kg were randomly assigned to 8 treatments with 6 replicates (4 pigs per pen). The experiment lasted 35 days (phase 1, d 1–21; phase 2, d 22–35) and followed a  $2 \times 2 \times 2$  factorial design with metabolizable energy levels (3,350 and 3,250 kcal/kg),  $\beta$ M supplementation (- and +0.05%), and EM supplementation (- and +0.05%). There was a tendency toward an interaction between EM and  $\beta$ M supplementation during phase 1 on fecal bacterial populations, by an increase in *Bifidobacterium* abundance ( $p = 0.083$ ) and a decrease in *Salmonella* counts ( $p = 0.081$ ). The higher energy level increased ( $p < 0.05$ ) final body weight, average daily gain, and feed efficiency. The dietary inclusion of EM tended to increase final body weight ( $p = 0.081$ ) and average daily gain ( $p = 0.082$ ). The supplementation of EM increased ( $p < 0.05$ ) the dry matter, gross energy, and ether extract digestibility in phase 1, and enhanced ( $p < 0.05$ ) dry matter, gross energy, organic matter, and ether extract digestibility in phase 2. The fecal *Escherichia coli* population tended to be lower while *Lactobacillus* increased ( $p = 0.084$ ) in the EM-supplemented group ( $p = 0.074$ ). In conclusion, EM supplementation improved growth performance and nutrient digestibility in grower pigs.  $\beta$ M supplementation showed limited effects but tended to interact with EM to help regulate intestinal *Salmonella* populations.

**Keywords:**  $\beta$ -mannanase, Emulsifier, Grower pigs, Growth performance, Nutrient digestibility.

## INTRODUCTION

Corn and soybean meal are the predominant ingredients in swine diets, serving as the main energy and protein sources. Lipids are crucial energy source in swine diets, providing high caloric density in small amounts [1]. Increasing dietary lipid inclusion not only enhances the energy concentration of the diet but also reduces the need for low digestible fibrous plant-based components, thereby improving feed efficiency [2]. However, practical swine diets based on corn and soybean meal inevitably contain substantial amounts of plant-derived non-starch polysaccharides (NSP), which can influence nutrient utilization, including lipid digestion.

Plant-based diets are rich in NSP including  $\beta$ -mannans, glucomannans, and galactomannans, with approximately 150-370 g/kg of the total NSP content [3,4]. Among these,  $\beta$ -mannans are known to increase intestinal digesta viscosity [5], slowing the passage rate and impeding enzymatic digestion and nutrient absorption [6]. Young pigs lack endogenous enzymes capable of hydrolyzing  $\beta$ -1,4-mannosyl and  $\alpha$ -1,6-galactosyl linkages, leading to incomplete degradation of  $\beta$ -mannans [7]. Although partial microbial fermentation occurs in the hindgut, the limited fermentative capacity of pigs often results in reduced nutrient digestibility and impaired growth performance [8]. Furthermore, undigested substrates can foster undesirable microbial fermentation, predisposing the gut environment to pathogenic bacterial proliferation [9,10]. To overcome these limitations, supplementation with exogenous  $\beta$ -mannanase ( $\beta$ M) has been proposed as a promising strategy.  $\beta$ M hydrolyzes  $\beta$ -mannans into manno-oligosaccharides (MOS) and mannose, which can be absorbed as additional energy sources [11]. Several studies have reported that  $\beta$ M supplementation reduces digesta viscosity, thereby enhancing nutrient digestibility [9,12] and growth performance in pigs [6,13].

An increased digesta viscosity not only hinders carbohydrate and protein digestion but also interferes with lipid digestion [14]. A viscous intestinal environment restricts lipase access and impairs fat emulsification [15]. Fat digestion depends largely on bile salts and pancreatic lipase, yet dietary fats exhibit inherently low digestibility due to their hydrophobic nature [16]. The use of exogenous emulsifiers (EM), which possess both hydrophilic and hydrophobic groups, has been shown to improve fat digestibility by enhancing emulsification and micelle formation [17-19]. Improved lipid utilization allows diets to be formulated with reduced energy density while maintaining overall nutrient balance and animal performance [20]. Moreover, enhanced fat digestion can promote the absorption of fat-soluble vitamins and improve feed efficiency [21,22]. Both  $\beta$ M and EM are recognized as effective feed additives for improving nutrient utilization and growth in pigs. Nevertheless, there is limited research exploring their combined or interactive effects, particularly under different dietary energy levels. Since  $\beta$ M may indirectly enhance lipid digestibility by reducing digesta viscosity, and EM directly promotes

lipid utilization, their concurrent use may yield synergistic effects, especially in low-energy diets in grower pig. Therefore, the objective of this study was to evaluate the effects of dietary energy level and supplementation with  $\beta$ -mannanase and an emulsifier, individually and in combination, on growth performance, nutrient digestibility, and fecal bacterial populations in grower pigs.

## **MATERIAL AND METHODS**

The Institutional Animal Care and Use Committee of Kangwon National University approved the animal care and experimental techniques utilized in this study (Ethical code: KW-240722-1).

### **Additive information**

The  $\beta$ M (800,000 U of  $\beta$ M/kg) was obtained from a commercial feed company (CTC Bio, Inc., Seoul, Republic of Korea) and it was produced by *Bacillus subtilis*. The EM produced by Molimen in Spain (Phospholipid + Lysophospholipid) was obtained from a commercial feed company (CTC Bio, Inc., Seoul, Republic of Korea).

### **Animals, experimental designs, and procedures**

The study was conducted on a commercial farm in Haman, Gyeongsangnam-do, Republic of Korea. A total of 192 pigs (Landrace  $\times$  Yorkshire  $\times$  Duroc; approximately 9 weeks of age) with an initial body weight (BW) of  $22.78 \pm 1.06$  kg were randomly assigned to 8 treatments with 6 replicates (2 barrows and 2 gilts per pen). Pigs were balanced across treatments according to initial body weight, age, and sex. The experiment was conducted for 35 days (phase 1, d 1-21; phase 2, d 22-35). The treatment groups were designed as a  $2 \times 2 \times 2$  factorial arrangement with energy levels (3,350 and 3,250 kcal/kg);  $\beta$ M supplementation (0 and 0.05%), and EM supplementation (0 and 0.05%). To ensure accurate inclusion of the low-level additives (0.05%), a pre-mix was prepared and thoroughly mixed with the basal diet using a precision batch mixer. Farm management, feeding, and animal health procedures followed the standard operating protocols established by the research facility, including twice-daily feeding, daily health checks, and strict biosecurity and sanitation measures. Diet formulations and chemical compositions are presented in Table 1, and all diets were formulated to meet or exceed NRC [23] nutrient recommendations. Experimental feeds were provided in mashed form and both feed and water were available *ad libitum*. No antibiotics, antimicrobial agents, or growth promoters were administered throughout the experimental period.

### **Experimental procedures and sample collection**

The experimental pig's BW was measured at the beginning and end of every period for average daily gain (ADG) calculation. Feed intake data were obtained by documenting the amount of feed provided

and the residual feed remaining in the feeders on a daily basis, with phase totals used to determine average daily feed intake (ADFI). Feed efficiency (G:F) was calculated based on ADG and ADFI. Mortality was 0%, and no pigs were removed during the experimental period. Each treatment consisted of 6 replicate pens with 4 pigs per pen, which were maintained until the end of the study.

#### **Nutrient digestibility**

To measure nutrient digestibility, a chromic oxide ( $\text{Cr}_2\text{O}_3$ ) indicator (0.25%) was added to all diets 7 days prior to sampling. On d 21 and 35, fecal samples were collected from at least one random pig per pen via rectal massage, ensuring uniform representation across all replicates. The collected feces and corresponding feed samples were analyzed for dry matter (DM), organic matter (OM), gross energy (GE), crude protein (CP), ether extract (EE), and crude fiber (CF). Samples were dried in a forced-air oven at 60°C for 72 h and ground through a 1 mm screen (Thomas Wiley Mill, Model 4). Digestibility markers in feed and feces were quantified spectrophotometrically using the method described by Jagger et al. (1992), and nutrient digestibility was calculated using the indicator method as follows:

$$\text{Digestibility (\%)} = 100 - [100 \times (\text{marker in feed} / \text{marker in feces}) \times (\text{nutrient in feces} / \text{nutrient in feed})]$$

Analytical procedures for DM, CP, EE, and CF followed AOAC International (2007) methods 930.15, 990.03, 960.39, and 978.10, respectively. OM was calculated as DM minus ash content. GE was determined using a bomb calorimeter (Model 1261, Parr Instrument Co., Moline, IL).

#### **Blood vitamin**

On d 21 and 35, blood samples were collected from all pigs in each pen to evaluate the retinol (Vitamin A), 25(OH) $\text{D}_3$  (vitamin D),  $\alpha$ -tocopherol (vitamin E), and menadione (vitamin K) properties in blood. The collected blood from the pig vena cava vein was centrifuged and the plasma was separated. The separated plasma sample (1ml) was transferred to a 5ml tube, protein was denatured with 500 $\mu$ l of pure ethanol, mixed by vortexer for 15 seconds, and extracted twice with 500 $\mu$ l of n-hexane. The extracted sample was collected from the organic layer and dried using nitrogen. The dried sample was vortexed with 100 $\mu$ l of methanol and centrifuged at 1,380 x g for 30 seconds. The precipitate 35 $\mu$ l was analyzed using an HPLC (high-performance liquid chromatography, Agilent Technologies Inc, Santa Clara, CA, USA) device according to the method of Yang et al. [24].

#### **Fecal bacterial populations**

On d 21 and 35, fecal samples were collected from all experimental groups for microbial analysis. Two pigs per pen (1 barrow and 1 gilt) were selected based on their BW to ensure that pigs were similar condition. Feces were directly collected from the rectum using sterile gloves and placed into individually labeled sterile plastic tubes under aseptic conditions. Immediately after collection, samples were snap-frozen in liquid nitrogen, transported to the laboratory, and stored at -80°C until analysis.

For DNA extraction pretreatment, the QIAamp Fast DNA stool Mini Kit (cat. no. 51604/2016) was used, and the procedure is as follows: 1) After weighing 200 mg of fecal samples, they were placed in a 2 ml centrifuge tube and kept on ice. After adding 1 ml InhibitEX buffer to the centrifuge tube, it was vortexed for 1 minute to maximize the DNA concentration and ensure uniform mixing. The samples were centrifuged at  $14,000 \times g$  for 1 min to separate pellet particles. 2) 25  $\mu$ l of proteinase K and 600  $\mu$ l of the supernatant from step 1 were transferred into a new 2 ml centrifuge tube and vortexed. Then, 600  $\mu$ l of ethanol was added to the centrifuge tube and vortexed again. The 600  $\mu$ l supernatant was transferred into a QIAamp spin column and centrifuged for one minute at 14,000 rpm. The QIAamp spin column was placed in a new 2 ml centrifuge tube, and the existing tube was discarded. The QIAamp spin column was opened, 500  $\mu$ l Buffer AW1 was added, and it was centrifuged under the same conditions. After transferring the QIAamp spin column to a new tube, 500  $\mu$ l of Buffer AW2 was added and it was centrifuged for three minutes. Finally, DNA was extracted by treating the QIAamp spin column with the same process.

Real-time polymerase chain reaction (qPCR) was employed for quantifying *Lactobacillus* spp., *Bifidobacterium* spp., *Clostridium* spp., *Salmonella* spp., *Escherichia coli* (*E. coli*), and  $\beta$ -actin (reference gene). To quantify microorganisms, 1 $\times$  universal SsoAdvancedTMR®Green Supermix, 2.5 ng/ $\mu$ l of primers, and 10 ng of DNA were added to a volume of 10  $\mu$ l [25]. Primer sequences are shown in Table 2. After 40 cycles at 95°C for 15 seconds for enzyme activation, SYBR green fluorescent signals were recorded at 72°C, and PCR results were derived after diluting 10 times for microbial quantification. For microbial quantification, qPCR Rotor-Gene Qiagen 2 plex program (Serial Number 0312272, Corvett Research) was used by Tajudeen et al. [26].

### Statistical analysis

Data generated in the present study were statistically analyzed using three-way factorial analysis of variance (ANOVA) based on the General Linear Model (GLM) procedure of SAS software (version 9.4; SAS Institute Inc., Cary, NC, USA) for growth performance, nutrient digestibility, blood vitamin and fecal bacterial populations parameters. The statistical model included dietary energy level,  $\beta$ M, and EM supplementation, and their interactions as fixed effects. The pen was considered the experimental unit for growth performance, whereas the individual pig was considered the experimental unit for nutrient digestibility, blood vitamin and fecal bacterial populations parameters. Significant differences among treatment means were determined using Tukey's honest significant difference (HSD) test. Data are presented as means  $\pm$  standard error of the mean (SEM). Differences were considered statistically significant at  $p < 0.05$ , and considered a tendency when  $0.05 \leq p < 0.10$ .

## RESULTS

## **Growth performance**

Pigs fed the high-energy diet had higher ( $p < 0.05$ ) final BW and showed a tendency ( $p = 0.081$ ) for increased BW in the EM-supplemented group (Table 3). During phase 1, there were no significant differences among treatments in ADG, ADFI, or G:F. In phase 2, there was no change in ADG and ADFI, however, pigs fed the high-energy diet tended ( $p = 0.091$ ). Overall (d 1–35), both ADG and G:F were increased ( $p < 0.05$ ) by the high-energy diet, and ADG tended ( $p = 0.082$ ) to be higher in pigs fed EM supplementation.

## **Nutrient digestibility**

There were no significant interaction effects among treatments were observed for nutrient digestibility during the experimental period (Table 4). In phase 1, EM supplementation increased ( $p < 0.05$ ) the digestibility of DM, GE, and EE, whereas CP and CF digestibility were unaffected. Neither dietary energy level nor  $\beta$ M supplementation influenced the digestibility of DM, OM, GE, CP, EE, or CF. In phase 2, EM supplementation increased ( $p < 0.05$ ) the digestibility of DM, OM, GE, and EE, while dietary energy level and  $\beta$ M supplementation had no effects.

## **Blood vitamin concentration**

There were no significant interaction effects among treatments were observed for blood vitamin concentrations in either phase (Figure 1). In phase 1, there was a tendency ( $p = 0.087$ ) for higher blood retinol levels in pigs fed the high-energy diet, without any significant effects on 25(OH)D3,  $\alpha$ -tocopherol, and menadione concentrations. The concentration of retinol, 25(OH)D3,  $\alpha$ -tocopherol, and menadione in blood was not affected with supplementation of  $\beta$ M and EM. In phase 2, none of the vitamin concentrations differed among treatment groups (Figure 2). Additionally, no interaction effects among treatments were observed in either phase.

## **Fecal bacterial populations**

An interaction tendency between EM and  $\beta$ M supplementation was observed in phase 1, resulting in increased *Bifidobacterium* counts ( $p = 0.083$ ) and decreased *Salmonella* counts ( $p = 0.081$ ) (Table 5). In phase 2, a similar interaction tendency ( $p = 0.084$ ) was observed, showing increased *Lactobacillus* abundance with the combined supplementation of  $\beta$ M and EM. Apart from these tendencies, no significant effects of dietary energy level,  $\beta$ M, or EM supplementation were detected on the populations of *Lactobacillus*, *Bifidobacterium*, *Clostridium*, *Salmonella*, or *E. coli* during either phase. However, EM supplementation tended to decrease *E. coli* counts ( $p = 0.074$ ) in phase 2.

## **DISCUSSION**



Energy sources account for approximately 70% of compound feed in swine production [27]. Consequently, strategies to improve energy utilization have received considerable attention [6,28]. In corn–soybean-based diets, NSP from plant cell walls are indigestible by endogenous enzymes, which can increase digesta viscosity, impair nutrient absorption, and impose metabolic burdens on pigs [29–31]. Therefore, exogenous enzymes such as  $\beta$ M have been widely studied for their ability to hydrolyze  $\beta$ -mannans, reduce viscosity, and enhance nutrient availability [4,32]. Likewise, the inclusion of exogenous EM in pig diets has been proposed to enhance lipid utilization. EM promotes emulsification by breaking down large fat globules into smaller micelles, thereby increasing lipase accessibility and improving energy digestibility [17–19].

In the present study, the high-energy diet improved growth performance, as reflected by a higher G:F despite no change in ADFI. This finding supports the established concept that increasing dietary energy density enhances feed efficiency. Similarly, EM supplementation tended to increase final body weight and ADG, which can be attributed to the observed increases in DM, GE, EE digestibility, reflecting improved lipid emulsification and energy utilization [16,17]. These findings align with the positive role of dietary energy and emulsifiers in grower pigs. In contrast,  $\beta$ M supplementation did not affect growth performance. This absence of effect is likely attributable to the low  $\beta$ -mannan concentration of the experimental diets (0.27–0.28%), which is lower than the  $\beta$ -mannan levels typically present in diets where consistent responses to  $\beta$ M supplementation have been reported in previous studies [31,33]. Mechanistically,  $\beta$ M is known to reduce intestinal viscosity and mitigate immune activation by degrading  $\beta$ -mannans that can mimic pathogen-associated molecular patterns [6]. Reduced immune stimulation may conserve metabolic energy for growth [13]. However, given the relatively low fiber and mannan levels in the current study, neither  $\beta$ M nor its interaction with EM produced measurable improvements in growth performance.

The EM supplementation increased the DM, OM, GE, and EE digestibility, consistent with previous studies demonstrating the positive effects of lysophospholipids and other emulsifiers on nutrient digestibility [19,34,35]. The tendency for higher growth performance in EM-supplemented treatments could be associated with enhanced digestibility. In our study, GE digestibility in phase 1 was higher in the low-energy diet when supplemented with EM compared to the high-energy diet. This suggests that EM can compensate for reduced dietary energy levels by improving fat emulsification and energy extraction efficiency. In contrast, neither dietary energy level nor  $\beta$ M supplementation significantly affected nutrient digestibility. While some studies have reported enhanced digestibility with  $\beta$ M supplementation in high energy diets [36,37], such responses are typically observed in diets with higher NSP or fiber levels [3,15]. The absence of effects on crude fiber digestibility in the present study likely

reflects the limited  $\beta$ -mannan content of the diet. Furthermore, no synergistic interaction was detected between  $\beta$ M and EM, suggesting that adequate substrate availability, either  $\beta$ -mannan or dietary fat, is essential to elicit a complementary effect.

The high-energy diet tended to increase the blood retinol concentration in grower pigs compared to the low-energy diet. This result may be explained by the higher fat content in the high-energy diet (6.55%) than in the 3,250 kcal/kg diet (4.59%). It is well known that a low-fat content in feed limits the absorption of fat-soluble vitamins [22]. However, other fat-soluble vitamins, such as 25(OH)D<sub>3</sub>,  $\alpha$ -tocopherol, and menadione, did not show significant differences. Currently, there is limited research on how the addition of EM and  $\beta$ M at different dietary energy levels affects blood vitamin levels in grower pigs, indicating a need for further investigation in this area.

Although statistical significance was not detected among treatment groups, there was a tendency for lower *E. coli* colonization with the supplementation of EM in phase 2. Furthermore, the interaction effects between the supplementation of EM and  $\beta$ M tended to increase *Bifidobacterium* and *Lactobacillus* levels while decreasing *Salmonella* levels. These tendencies may be associated with a reduction in undigested nutrients resulting from improved nutrient digestibility, rather than indicating a direct causal relationship. The fecal bacteria count serves as an indirect indicator of the intestinal microbiota, with the distribution of beneficial and harmful bacteria known to impact gut health, digestibility, and growth performance [38]. Factors influencing fecal microorganisms include nutritional elements such as feed composition and nutrients, as well as environmental factors like temperature and stress [39,40]. Some pathogenic microbes, such as *E. coli* and *Salmonella*, can utilize undigested nutrients for fermentation, including protein fractions, under certain dietary conditions and potentially produce harmful metabolites [9]. Among these, nutritional factors have the most significant impact on the gut microbiota, as undigested nutrients interact with gut microorganisms, affecting the microbial community [10,41,42].

## CONCLUSION

In conclusion, regardless of dietary energy level, EM supplementation improved growth performance and nutrient digestibility in grower pigs and was reduced in fecal *Escherichia coli* populations.  $\beta$ M supplementation alone showed limited effects; however, a tendency for interaction between  $\beta$ M and EM was observed in fecal bacterial populations during the early growth phase. The reduction of dietary energy level to 3,250 kcal/kg is not recommended due to compromised growth performance, and further studies are warranted to optimize the combined use of  $\beta$ M and EM under practical feeding conditions.

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ACCEPTED

## Tables

Table 1. Experimental basal diet

|                                       |        |        |
|---------------------------------------|--------|--------|
| Metabolizable energy, kcal/kg         | 3,350  | 3,250  |
| Ingredients, %                        |        |        |
| Corn                                  | 59.46  | 61.62  |
| Soybean meal                          | 16.31  | 15.90  |
| Dried distiller's grains with soluble | 10.00  | 10.00  |
| Wheat bran                            | 4.00   | 4.00   |
| Animal fat                            | 4.03   | 2.00   |
| Molasses                              | 3.00   | 3.00   |
| L-Lysine (78%)                        | 0.56   | 0.57   |
| DL-Methionine (99%)                   | 0.13   | 0.13   |
| L-Threonine (99%)                     | 0.15   | 0.15   |
| L-Tryptophan (100%)                   | 0.31   | 0.58   |
| Limestone                             | 1.01   | 1.02   |
| Dicalcium phosphate                   | 0.34   | 0.33   |
| Salt                                  | 0.30   | 0.30   |
| Vitamin premix <sup>1</sup>           | 0.15   | 0.15   |
| Mineral premix <sup>2</sup>           | 0.15   | 0.15   |
| Choline chloride                      | 0.05   | 0.05   |
| Phytase <sup>3</sup>                  | 0.05   | 0.05   |
| Total                                 | 100.00 | 100.00 |
| Chemical composition, %               |        |        |
| ME, kcal/kg                           | 3,350  | 3,250  |
| Crude Protein                         | 16.00  | 16.00  |
| Crude fat                             | 6.55   | 4.59   |
| Crude fiber                           | 3.57   | 3.59   |
| Ca                                    | 0.66   | 0.66   |
| Na                                    | 0.10   | 0.10   |
| STTD P                                | 0.31   | 0.31   |
| SID Lysine                            | 0.98   | 0.98   |
| SID Threonine                         | 0.59   | 0.59   |
| SID Met + Cys                         | 0.55   | 0.55   |

<sup>1</sup>Supplied per kg of diet: 16,000 IU vitamin A (palmitate), 2.00mg vitamin B<sub>1</sub> (thiamin), 5.00mg vitamin B<sub>2</sub> (riboflavin), 2.00mg vitamin B<sub>6</sub> (pyridoxine), 0.03 mg vitamin B<sub>12</sub> (cyanocobalamin), 25.00 mg niacin, 0.40 mg folic acid, 0.05 mg biotin, 5.00 mg ethoxyquin, 2,000 IU vitamin D<sub>3</sub> (cholecalciferol), 75.00 mg vitamin E (dl- $\alpha$ -tocopheryl acetate), 2.00 mg vitamin K<sub>3</sub> (menadione).

<sup>2</sup>Supplied per kg of diet: 100 mg Fe, 6 mg Cu, 4 mg Mn, 0.3 mg Se, 0.14 mg I, 0.25 mg Co.

<sup>3</sup>6-phytase from *E. coli* was added at 500 FTU/kg. Assigned nutrient release: 0.10% digestible P, 0.02% Ca.

Table 2. Primer sequence

| Item                        | Primer sequence                                      | Reference             |
|-----------------------------|--|-----------------------|
| <i>Lactobacillus spp.</i>   | F: AGCAGTAGGGAATCTTCCA<br>R: CACCGCTACACATGGAG       | Walter et al., 2001   |
| <i>Bifidobacterium spp.</i> | F: TCGCGTCYGGTGTGAAAG<br>R: CCACATCCAGCRTCCAC        | Rinttila et al., 2004 |
| <i>Clostridium spp.</i>     | F: GGCGGCYTRCTGGGCTTT<br>R: CCAGGTGGATWACTTATTGTGTAA | Omar et al., 2013     |
| <i>Salmonella spp.</i>      | F: TCGTCATTCCATTACCTACC<br>R: AAACGTTGAAAACTGAGGA    | Idrus et al., 2021    |
| <i>Escherichia coli</i>     | F: AAAACGGCAAGAAAAAGCAG<br>R: GCGTGGTTACAGTCTTGCG    | Amofa et al., 2022    |
| $\beta$ -Actin              | F: CTCCTTCTTGGGCATGGA<br>R: CGCACTTCATGATCGAGTTGA    | Leng et al., 2007     |



Table 3. The effects of  $\beta$ -mannanase ( $\beta$ M) and emulsifier (EM) supplementation in different dietary energy levels (EN) on growth performance of grower pigs.

| EN, kcal/kg       | 3,350 |       |       |       | 3,250 |       |       |       | <i>p</i> -value |       |       |       |               |               |               |                  |
|-------------------|-------|-------|-------|-------|-------|-------|-------|-------|-----------------|-------|-------|-------|---------------|---------------|---------------|------------------|
| βM <sup>1</sup>   | -     |       | +     |       | -     |       | +     |       | SEM             | EN    | EM    | βM    | EN<br>×<br>EM | EN<br>×<br>βM | EM<br>×<br>βM | EN×<br>EM×<br>βM |
| EM <sup>2</sup>   | -     | +     | -     | +     | -     | +     | -     | +     |                 |       |       |       |               |               |               |                  |
| BW, kg            |       |       |       |       |       |       |       |       |                 |       |       |       |               |               |               |                  |
| Initial           | 22.78 | 22.76 | 22.79 | 22.79 | 22.76 | 22.78 | 22.77 | 22.80 | 0.002           | 0.470 | 0.885 | 0.750 | 0.750         | 0.543         | 0.543         | 0.885            |
| Final             | 44.01 | 44.38 | 44.05 | 44.36 | 43.03 | 43.83 | 43.46 | 43.88 | 0.174           | 0.019 | 0.081 | 0.646 | 0.611         | 0.667         | 0.681         | 0.768            |
| Phase 1 (d 0-21)  |       |       |       |       |       |       |       |       |                 |       |       |       |               |               |               |                  |
| ADG, g/d          | 751   | 762   | 751   | 758   | 712   | 746   | 734   | 745   | 9.537           | 0.269 | 0.405 | 0.825 | 0.716         | 0.744         | 0.716         | 0.824            |
| ADFI, g/d         | 1,343 | 1,340 | 1,335 | 1,343 | 1,346 | 1,350 | 1,350 | 1,348 | 5.837           | 0.488 | 0.852 | 0.956 | 0.954         | 0.907         | 0.923         | 0.730            |
| G:F, g/g          | 0.56  | 0.57  | 0.56  | 0.56  | 0.53  | 0.55  | 0.55  | 0.55  | 0.007           | 0.156 | 0.451 | 0.789 | 0.729         | 0.747         | 0.635         | 0.891            |
| Phase 2 (d 22-35) |       |       |       |       |       |       |       |       |                 |       |       |       |               |               |               |                  |
| ADG, g/d          | 765   | 780   | 767   | 783   | 734   | 757   | 743   | 762   | 8.152           | 0.185 | 0.334 | 0.802 | 0.879         | 0.909         | 0.970         | 0.939            |
| ADFI, g/d         | 1,510 | 1,517 | 1,500 | 1,514 | 1,551 | 1,522 | 1,531 | 1,522 | 8.775           | 0.235 | 0.826 | 0.633 | 0.403         | 0.935         | 0.707         | 0.854            |
| G:F, g/g          | 0.51  | 0.52  | 0.51  | 0.52  | 0.47  | 0.50  | 0.48  | 0.50  | 0.007           | 0.091 | 0.336 | 0.735 | 0.679         | 0.878         | 0.867         | 0.954            |
| Overall (d 0-35)  |       |       |       |       |       |       |       |       |                 |       |       |       |               |               |               |                  |
| ADG, g/d          | 607   | 617   | 608   | 617   | 579   | 602   | 591   | 603   | 3.802           | 0.020 | 0.082 | 0.644 | 0.609         | 0.662         | 0.688         | 0.767            |
| ADFI, g/d         | 1,410 | 1,411 | 1,401 | 1,412 | 1,428 | 1,419 | 1,422 | 1,418 | 5.071           | 0.195 | 0.981 | 0.713 | 0.535         | 0.981         | 0.744         | 0.911            |
| G:F, g/g          | 0.43  | 0.44  | 0.43  | 0.44  | 0.41  | 0.42  | 0.42  | 0.43  | 0.003           | 0.009 | 0.134 | 0.579 | 0.478         | 0.723         | 0.598         | 0.877            |

<sup>1,2</sup> -, without supplementation; +, 0.05% supplemented.

Abbreviation: SEM, standard error of means; BW, body weight; ADG, average daily weight gain; ADFI, average daily feed intake; G:F, feed efficiency. Each value represents the mean of 6 replicate pens (2 barrow and 2 gilts per pen).

Table 4. The effects of  $\beta$ -mannanase ( $\beta$ M) and emulsifier (EM) supplementation in different dietary energy levels (EN) on nutrient digestibility of grower pigs.

| EN, kcal/kg     | 3,350 |       |       |       | 3,250 |       |       |       | <i>p</i> -value <sup>3</sup> |       |        |       |               |               |               |                  |
|-----------------|-------|-------|-------|-------|-------|-------|-------|-------|------------------------------|-------|--------|-------|---------------|---------------|---------------|------------------|
| βM <sup>1</sup> | -     |       | +     |       | -     |       | +     |       | SEM                          | EN    | EM     | βM    | EN<br>×<br>EM | EN<br>×<br>βM | EM<br>×<br>βM | EN×<br>EM×<br>βM |
| EM <sup>2</sup> | -     | +     | -     | +     | -     | +     | -     | +     |                              |       |        |       |               |               |               |                  |
| Phase 1 (d 21)  |       |       |       |       |       |       |       |       |                              |       |        |       |               |               |               |                  |
| DM, %           | 78.09 | 78.19 | 78.81 | 79.46 | 78.88 | 78.95 | 78.75 | 79.53 | 0.121                        | 0.117 | 0.016  | 0.104 | 0.121         | 0.917         | 0.197         | 0.874            |
| OM, %           | 79.10 | 79.20 | 79.55 | 80.08 | 79.51 | 79.64 | 79.69 | 80.14 | 0.190                        | 0.494 | 0.197  | 0.430 | 0.670         | 0.967         | 0.625         | 0.941            |
| GE, %           | 82.95 | 83.34 | 83.36 | 83.83 | 83.07 | 83.52 | 83.27 | 83.64 | 0.097                        | 0.989 | 0.037  | 0.122 | 0.957         | 0.460         | 0.997         | 0.845            |
| CP, %           | 79.20 | 79.38 | 79.55 | 79.56 | 79.28 | 79.37 | 79.41 | 79.99 | 0.198                        | 0.820 | 0.427  | 0.589 | 0.888         | 0.764         | 0.832         | 0.678            |
| EE, %           | 70.19 | 70.76 | 74.53 | 75.43 | 69.59 | 70.75 | 74.33 | 74.89 | 0.253                        | 0.507 | <0.001 | 0.123 | 0.947         | 0.907         | 0.893         | 0.646            |
| CF, %           | 41.59 | 42.41 | 42.27 | 42.70 | 42.95 | 42.02 | 41.55 | 42.83 | 0.193                        | 0.813 | 0.311  | 0.809 | 0.558         | 0.318         | 0.246         | 0.100            |
| Phase 2 (d 35)  |       |       |       |       |       |       |       |       |                              |       |        |       |               |               |               |                  |
| DM, %           | 77.18 | 77.35 | 77.76 | 78.52 | 77.90 | 77.94 | 78.15 | 78.56 | 0.130                        | 0.100 | 0.015  | 0.191 | 0.407         | 0.652         | 0.364         | 0.835            |
| OM, %           | 78.61 | 78.79 | 79.07 | 79.19 | 78.94 | 79.00 | 79.39 | 79.46 | 0.110                        | 0.204 | 0.049  | 0.621 | 0.957         | 0.855         | 0.948         | 0.939            |
| GE, %           | 81.96 | 82.61 | 82.28 | 82.87 | 82.06 | 82.67 | 82.16 | 82.85 | 0.120                        | 0.976 | 0.012  | 0.380 | 0.944         | 0.761         | 0.978         | 0.885            |
| CP, %           | 78.34 | 78.43 | 78.55 | 78.61 | 78.35 | 78.38 | 78.45 | 78.88 | 0.196                        | 0.932 | 0.533  | 0.698 | 0.905         | 0.845         | 0.813         | 0.787            |
| EE, %           | 69.84 | 70.17 | 74.02 | 74.85 | 69.03 | 70.39 | 73.53 | 74.38 | 0.253                        | 0.453 | <0.001 | 0.104 | 0.854         | 0.608         | 0.994         | 0.625            |
| CF, %           | 42.30 | 42.57 | 42.89 | 42.85 | 42.35 | 42.45 | 42.67 | 42.75 | 0.173                        | 0.775 | 0.773  | 0.284 | 0.974         | 0.859         | 0.818         | 0.839            |

<sup>1, 2</sup> -, without supplementation; +, 0.05% supplemented.

Abbreviation: SEM, standard error of means; DM, dry matter; OM, organic matter; GE, gross energy; CP, crude protein; EE, ether extract.

Each value represents the mean of 6 replicate pens (2 barrow and 2 gilts per pen).

Table 5. The effects of  $\beta$ -mannanase ( $\beta$ M) and emulsifier (EM) supplementation in different dietary energy levels (EN) on fecal bacterial populations of grower pigs

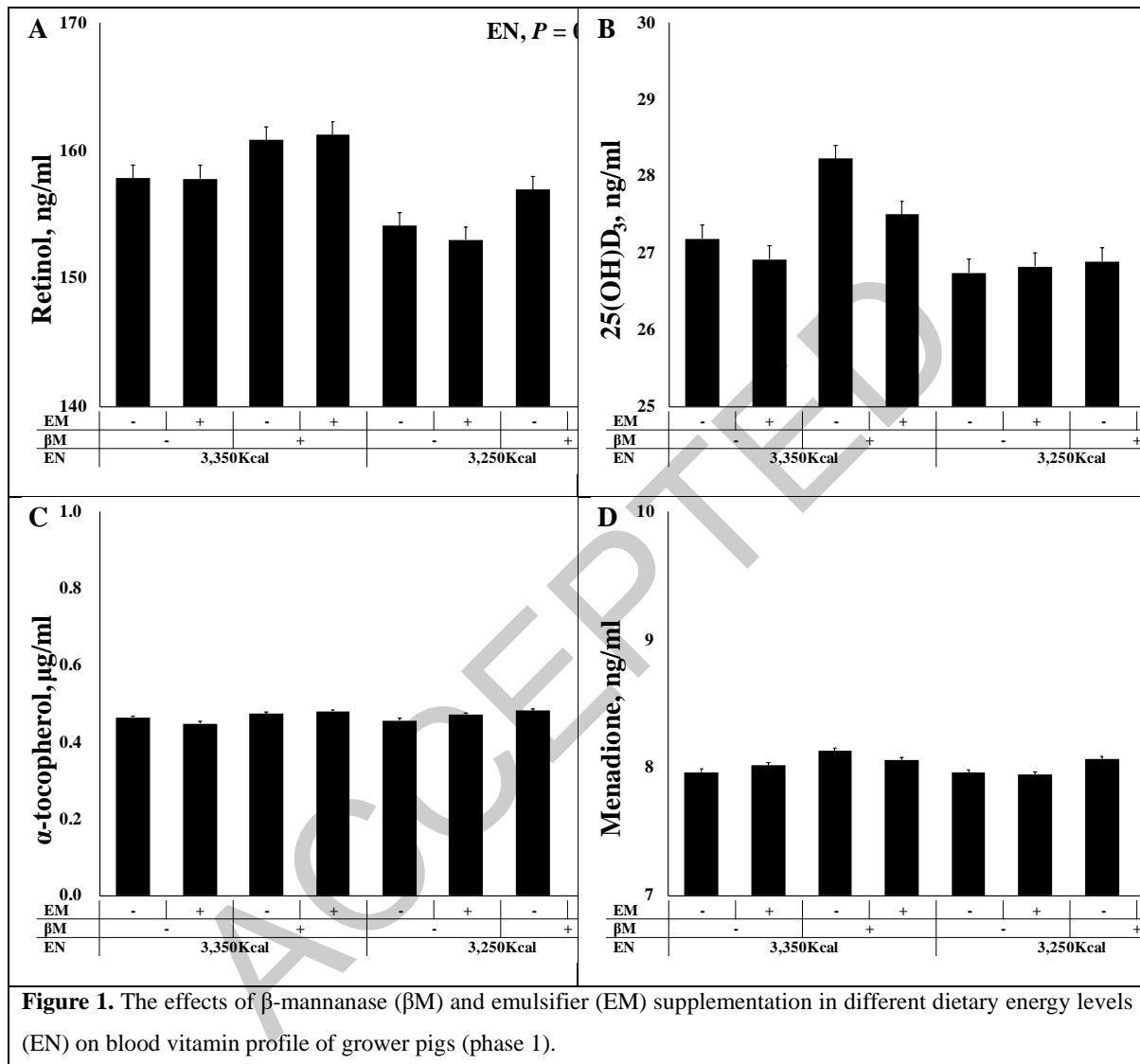
| EN, kcal/kg            | 3,350 |      |      |      | 3,250 |      |      |      | SEM   | <i>p</i> -value <sup>3</sup> |       |       |               |               |               |                  |
|------------------------|-------|------|------|------|-------|------|------|------|-------|------------------------------|-------|-------|---------------|---------------|---------------|------------------|
| βM <sup>1</sup>        | -     |      | +    |      | -     |      | +    |      |       | EN                           | EM    | βM    | EN<br>×<br>EM | EN<br>×<br>βM | EM<br>×<br>βM | EN×<br>EM×<br>βM |
| EM <sup>2</sup>        | -     | +    | -    | +    | -     | +    | -    | +    |       |                              |       |       |               |               |               |                  |
|                        |       |      |      |      |       |      |      |      |       |                              |       |       |               |               |               |                  |
| Phase 1 (d 21)         |       |      |      |      |       |      |      |      |       |                              |       |       |               |               |               |                  |
| <i>Lactobacillus</i>   | 1.42  | 1.56 | 1.54 | 1.58 | 1.49  | 1.48 | 1.46 | 1.40 | 0.100 | 0.394                        | 0.687 | 0.925 | 0.418         | 0.431         | 0.615         | 0.873            |
| <i>Bifidobacterium</i> | 1.83  | 2.00 | 1.77 | 1.78 | 1.92  | 1.99 | 2.10 | 1.71 | 0.130 | 0.321                        | 0.684 | 0.278 | 0.155         | 0.630         | 0.083         | 0.388            |
| <i>Clostridium</i>     | 0.54  | 0.52 | 0.53 | 0.43 | 0.57  | 0.48 | 0.60 | 0.73 | 0.120 | 0.259                        | 0.799 | 0.565 | 0.608         | 0.243         | 0.637         | 0.366            |
| <i>Salmonella</i>      | 0.79  | 0.56 | 0.55 | 0.63 | 0.94  | 0.78 | 0.56 | 0.86 | 0.120 | 0.192                        | 0.983 | 0.156 | 0.332         | 0.740         | 0.081         | 0.839            |
| <i>E. coli</i>         | 0.61  | 0.51 | 0.56 | 0.63 | 0.59  | 0.48 | 0.45 | 0.57 | 0.090 | 0.421                        | 0.933 | 0.885 | 0.875         | 0.664         | 0.156         | 0.847            |
| Phase 2 (d 35)         |       |      |      |      |       |      |      |      |       |                              |       |       |               |               |               |                  |
| <i>Lactobacillus</i>   | 1.51  | 1.40 | 1.44 | 1.54 | 1.65  | 1.57 | 1.41 | 1.65 | 0.110 | 0.211                        | 0.603 | 0.773 | 0.543         | 0.453         | 0.084         | 0.698            |
| <i>Bifidobacterium</i> | 1.86  | 1.96 | 1.87 | 1.90 | 2.03  | 1.91 | 1.82 | 1.86 | 0.110 | 0.912                        | 0.888 | 0.367 | 0.495         | 0.520         | 0.770         | 0.476            |
| <i>Clostridium</i>     | 0.68  | 0.47 | 0.54 | 0.44 | 0.58  | 0.47 | 0.57 | 0.51 | 0.100 | 0.923                        | 0.100 | 0.593 | 0.626         | 0.501         | 0.593         | 0.834            |
| <i>Salmonella</i>      | 0.63  | 0.76 | 0.56 | 0.74 | 0.68  | 0.67 | 0.56 | 0.69 | 0.130 | 0.782                        | 0.243 | 0.612 | 0.599         | 0.972         | 0.612         | 0.803            |
| <i>E. coli</i>         | 0.64  | 0.45 | 0.51 | 0.53 | 0.75  | 0.45 | 0.59 | 0.51 | 0.100 | 0.586                        | 0.074 | 0.625 | 0.484         | 0.867         | 0.166         | 0.991            |

<sup>1, 2</sup> -, without supplementation; +, 0.05% supplemented.

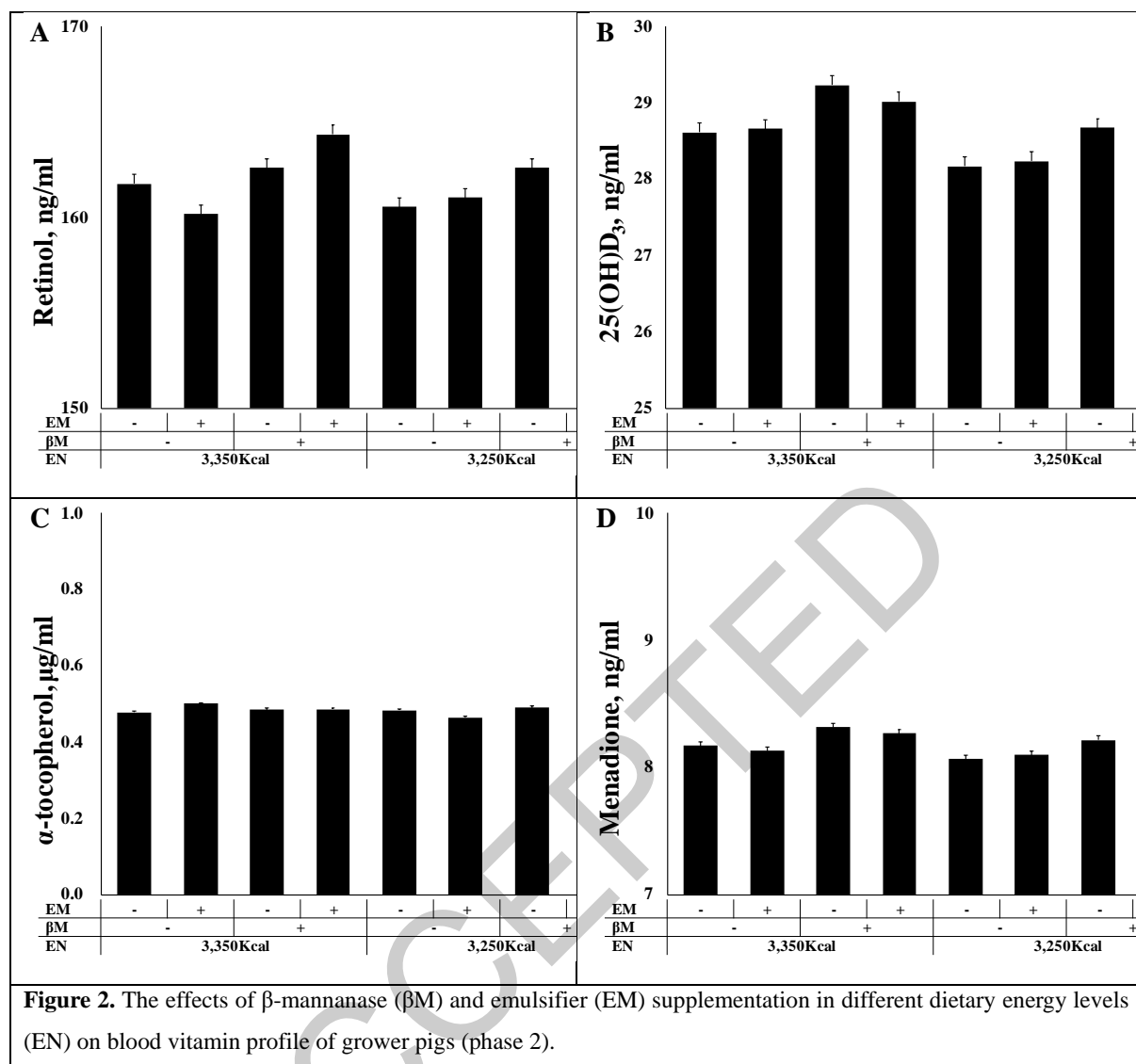
Abbreviation: SEM, standard error of means.

Each value represents the mean of 6 replicate pens (2 barrow and 2 gilts per pen).

## Figure legends



**Figure 1.** The effects of  $\beta$ -mannanase ( $\beta$ M) and emulsifier (EM) supplementation in different dietary energy levels (EN) on blood vitamin profile of grower pigs (phase 1).



**Figure 2.** The effects of β-mannanase (βM) and emulsifier (EM) supplementation in different dietary energy levels (EN) on blood vitamin profile of grower pigs (phase 2).