



Predicting the Digestible Energy of Rapeseed Meal from Its Chemical Composition in Growing-finishing Pigs

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ABSTRACT : Two experiments were conducted to establish a digestible energy (DE) content prediction model of rapeseed meal for growing-finishing pig based on rapeseed meal's chemical composition. In experiment 1, observed linear relationships between the determined DE content of 22 rapeseed meal calibration samples and proximate nutrients, gross energy (GE) and neutral detergent fiber (NDF) were used to develop the DE prediction model. In experiment 2, 4 samples of rapeseed meal selected at random from the primary rapeseed growing regions of China were used for testing the accuracy of DE prediction models. The results indicated that the DE was negatively correlated with NDF ($r = -0.86$) and acid detergent fiber (ADF) ($r = -0.73$) contents, and moderately correlated with gross energy (GE; $r = 0.56$) content in rapeseed meal calibration samples. In contrast, no significant correlations were found for crude protein, ether extract, crude fiber and ash contents. According to the regression analysis, NDF or both NDF and GE were found to be useful for the DE prediction models. Two prediction models: $DE = 16.775 - 0.147 \times NDF$ ($R^2 = 0.73$) and $DE = 11.848 - 0.131 \times NDF + 0.231 \times GE$ ($R^2 = 0.76$) were obtained. The maximum absolute difference between the *in vivo* DE determinations and the predicted DE values was 0.62 MJ/kg and the relative difference was 5.21%. Therefore, it was concluded that, for growing-finishing pigs, these two prediction models could be used to predict the DE content of rapeseed meal with acceptable accuracy. (**Key Words :** Pig, Rapeseed Meal, Digestible Energy)

INTRODUCTION

Rapeseed meal (RSM) is the second most widely traded protein ingredient after soybean meal representing a 12.40% of the world protein meal production, reaching 207 million metric tons in 2004 and 2005 (Ash and Dohlman, 2006). Rapeseed meal is readily available and could present an alternative and economical source of dietary protein for feed. After oil extraction, rapeseed meal has a high content of crude protein (35-40%, Näsi and Siljander-Rasi, 1991; 30-40%, Roth-Maier et al., 2004) and a higher neutral detergent fiber (NDF) content than soybean meal because hulls are not eliminated and represent about 30% of the meal; this causes a lower digestible energy content than that of soybean meal (2.6 vs. 3.4 Mcal/kg; Bell, 1993).

There is very little data on the digestible energy (DE) or metabolizable energy (ME) value of RSM, but RSM is widely used in pig diets. Therefore, it is important to estimate precisely the energy value of RSM, both for least

cost formulation purposes and for adapting the feed supply to the energy requirements of animals. *In vivo* digestibility studies with pigs to estimate the nutritive values of feed ingredients are a time-, cost-, and labor-intensive process; hence, the prediction of DE content from chemical composition, which can be determined rapidly *in vitro*, can be a useful tool for addressing DE variation and for accurate diet formation. Several factors can affect the accuracy of DE prediction models, which subsequently influence their successful use. One such factor is the sample size for regression analysis; another is the representative value of samples for the feedstuff as a whole (Zhao et al., 2008). Prediction models from smaller sample sizes may have greater R^2 and less residual standard deviation (RSD), but may not be as accurate as other models developed with a greater number of samples. On the other hand, the range of DE and chemical composition contents of samples obviously affect the accuracy of the prediction model. For example, low variation in DE and chemical composition contents of samples might provide an incorrect prediction model (Carré, 1990). Therefore, this study utilized 22 calibration samples comprising rapeseed meals from different regions or different production styles, containing a

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Table 1. Growth location and chemical composition of RSM (DM basis %, MJ/kg), experiment 1

Item ¹	Growth location	CP	CF	EE	Ash	NDF	ADF	GE	Ca	P
RSM-HUB	Hubei	43.05	13.70	0.64	10.99	45.48	27.78	19.19	0.91	0.98
RSM-NM	Neimenggu	42.28	13.91	2.35	7.84	37.80	26.42	20.02	0.76	0.95
RSM-HUN	Hunan	41.36	14.88	1.18	8.73	48.64	23.68	19.59	0.83	0.81
RSM-GZ	Guizhou	38.04	13.95	0.43	12.92	50.05	32.20	15.70	0.71	0.66
RSM-WP	Hubei	40.87	19.06	3.34	8.73	45.57	26.88	20.46	0.79	0.84
RSM-WPN	Hubei	59.02	14.63	0.76	10.28	49.90	15.44	20.91	0.93	0.36
RSM-WP1	Hubei	35.96	17.79	2.32	5.89	63.30	44.05	21.08	1.14	0.39

¹ Mean of three determinations per sample. CF = Crude fiber; ADF = Acid detergent fiber; NDF = Neutral detergent fiber; GE = Gross energy; EE = Ether extract.

large range of different chemical compositions, to establish DE or ME prediction models for growing-finishing pigs.

MATERIALS AND METHODS

Swine DE and ME assay all procedures were approved by the Institutional Animal Care and Use Committee at China Agricultural University, Beijing.

Difference method

The difference method is recommended when high levels of raw materials cannot be used because of toxicity or lack of palatability (Knabe et al., 1989). Due to its high level of fiber (Bell, 1993; Mińkowski, 2002) and its negative effect on feed intake, RSM is not recommended as the sole protein source in pigs when used to estimate its DE and ME. For this reason, the difference method was used. Preliminary observations were conducted on 10 pigs (35±1.1 kg of BW) fed graded quantities (10%, 15%, 20% and 25%) of RSM indicated that the inclusion level of RSM at 15% would maintain normal feed intake. In the experiment 1 and 2, the control diet was a corn-soybean meal diet and the experiment diets replaced 15% of the corn-soybean meal with RSM.

Diets

The objective of experiment 1 was to determine the relationship between DE (or ME) and the chemical composition of 22 RSM samples to develop a prediction model for DE or ME that could be utilized for the formulation of diets for growing-finishing pigs. Seven samples of prepress-solvent extracted RSM from 4 main rape production areas in China were collected from July to September 2010 (Table 1). The rapeseed meal of RSM-WP, RSM-WP1 and the protein-rich rapeseed meal production of RSM-WPN were produced by the Hubei Weipu Biological Technology Company. The 22 rapeseed meal calibration samples (Table 2) were made by combining different percentages of rapeseed meal to provide a wide range of proximate nutrient compositions and a big sample size. The composition and nutrient content of the diets used

in experiments 1 and 2 is presented in Table 3. In experiment 2, four prepress-solvent extracted RSM from different regions of China were used to test accuracy of the prediction models obtained in experiment 1 (Table 4).

Animals and experimental design

Experiments 1 and 2 were done simultaneously. Thirty Duroc×(Landrace×Large White) castrated male pigs weighing 30.5±2.1 kg were used in five incomplete Latin squares (6×6). Each diet was measured with 6 pigs. The control diet was fed to two pigs in each experiment period, meaning the control diet was measured with 12 pigs. The pigs were placed individually in metabolism cages provided

Table 2. The composition of calibration RSM, experiment 1

Calibration RSM sample	RSM sample ingredients (%)						
	HUB	NM	HUN	GZ	WP1	WP	WPN
1	-	-	-	-	7	91	2
2	-	-	-	-	14	82	4
3	-	-	-	-	23	70	7
4	-	-	-	-	31	60	9
5	-	-	-	-	39	49	12
6	-	-	-	-	47	39	14
7	-	-	-	-	20	80	-
8	-	-	-	-	27	59	14
9	-	-	-	-	31	48	21
10	-	-	-	-	35	36	29
11	-	-	-	-	39	25	36
12	-	-	-	-	72	28	-
13	-	-	-	-	52	48	-
14	40	-	-	-	-	60	-
15	-	-	-	-	-	100	-
16	-	-	35	-	-	65	-
17	-	-	74	-	-	26	-
18	-	-	-	28	-	72	-
19	-	-	-	37	-	63	-
20	-	-	-	47	-	53	-
21	-	-	-	66	-	34	-
22	-	-	-	85	-	15	-

Table 3. Ingredient composition (% , as-fed basis) of control and experimental diets

Ingredient	Control diet	Experimental diets	
	(n = 2)	(experiment 1: n = 22; experiment 2: n = 4)	
Corn	77.3	65.705	
Soybean meal	18.6	15.81	
Rapeseed meal	0	14.4	
L-lysine	0.1	0.085	
Dicalcium phosphate	0.9	0.9	
Calcium carbonate	0.9	0.9	
Carrier-medical stone	0.9	0.9	
Salt	0.3	0.3	
Minerals and vitamins premix ¹	1	1	
Total	100	100	

Analyzed nutrient content		Mean	Range
DM	87.71	88.21	87.78 to 89.31
CP	16.43	18.82	17.38 to 19.76
CF	2.73	4.75	3.44 to 5.61
EE	1.66	1.58	1.50 to 1.67
Ash	5.29	6.34	5.93 to 7.29
NDF	13.31	17.34	13.13 to 19.68
ADF	5.29	8.95	7.37 to 9.73

¹ Supplied per kilogram of diet: vitamin A (retinyl acetate), 4,500 IU; vitamin D₃, 1,400 IU; vitamin E (DL- α -tocopheryl acetate), 13.5 IU; vitamin K₃, 2.7 mg; thiamin, 0.9 mg; riboflavin, 2.7 mg; pyridoxine, 1.4 mg; vitamin B₁₂ (cobalamin), 9 μ g; pantothenate, 11 mg; folate, 0.60 mg; biotin, 0.04 mg; choline chloride, 350 mg; copper (CuSO₄·5H₂O), 18 mg; iron (FeSO₄·7H₂O), 75 mg; zinc (ZnSO₄), 70 mg; manganese (MnSO₄·H₂O), 20 mg; selenium (Na₂SeO₃), 0.3 mg; iodine (KI), 0.35 mg.

with a feeder and a nipple watering device set in a room with controlled temperature (19±2°C). Pigs were adapted to the digestibility cage for a period of 10 d before the collection of feces and urine. The whole experiment would be divided into 6 phases of collection of feces and urine, each phase lasts 10 d. The first 5 d of each phase were for feed adaptation and the last 5 d were for separate and total collection of feces and urine. Feed quantity was increased gradually during the experiment period. This feeding level represents about 90% of the spontaneous feed intake of the cage-housed pigs. All the pigs received their diets twice daily (0830 and 1530 h) in two equal meals and had free access to water.

For each diet, a sample of feed was collected and measured for its DM content and subsequently used for chemical analyses. Each day, feces and acidified (with H₂SO₄ to reach a pH below 2.0) urine were collected. Daily urine and feces collection were cumulated and stored at

-20°C. Collected feces were then homogenized and subsampled for DM analysis and freeze-dried for further chemical analyses at the end of the collection period.

Chemical analyses

The AOAC (2000) methods were used for measuring moisture, ash, crude protein (N×6.25; Kjeltec, 2100), Weende crude fiber (CF), and crude fat (extracted with petroleum ether; Soxtec Avanti 2050; Foss, Höganäs, Sweden). The ADF and NDF contents of feedstuffs were determined according to the procedure described by Van Soest (1963) and Van Soest et al. (1991). Samples of rapeseed meal, diets, feces and urine were analyzed for gross energy (GE) via adiabatic oxygen bomb calorimeter (Parr Instruments, Moline, IL, UAS).

Calculations and statistical analyses

Gross energy consumed was calculated by multiplying

Table 4. Growth location and chemical composition of testing RSM (DM basis %, MJ/kg), experiment 2

Item	Growth location	CP	CF	EE	Ash	NDF	ADF	GE	DE	DE/GE	Ca	P
RSM-XJ	Xinjiang	43.38	9.60	1.92	7.62	35.74	16.68	19.86	11.23	0.57	0.83	0.79
RSM-HEN	Henan	40.19	14.28	3.54	11.86	42.30	23.09	19.39	11.19	0.58	0.78	0.97
RSM-FJ	Fujian	41.49	13.23	2.06	7.18	30.82	22.31	20.00	12.26	0.61	0.70	1.03
RSM-SC	Sichuan	44.75	15.52	0.65	6.10	37.29	23.81	19.87	11.91	0.60	0.25	0.96

Table 5. Correlation coefficients between chemical composition and DE (ME) of calibration samples, experiment 1

	CP	CF	EE	ASH	NDF	ADF	GE	DE	ME
CP	1								
CF	-0.15	1							
EE	-0.22	-0.24	1						
ASH	-0.14	-0.63	-0.07	1					
NDF	-0.37	0.30	0.06	0.04	1				
ADF	-0.33	0.14	-0.03	0.15	0.63	1			
GE	0.21	0.21	-0.02	-0.49	-0.48	-0.56	1		
DE	0.39	-0.38	-0.02	0.03	-0.86	-0.73	0.56	1	
ME	0.27	-0.11	0.11	-0.40	-0.29	-0.46	0.44	0.45	1

the GE value of the diet fed by feed intake over the 5 d collection period. Apparent DE values were calculated by subtracting fecal energy from intake energy. Apparent ME values were calculated by subtracting urinary energy from apparent DE.

Data were analyzed by ANOVA using the procedure of SAS. In all analyses, $p < 0.05$ was considered significant. Simple regression analyses were conducted to establish prediction equations.

RESULTS

The growth location and nutrients of RSM used in experiment 1 and 2 were presented in Table 1 and 4. Excluding the protein-rich rapeseed meal production of RSM-WPN, the range of RSM crude protein were 35.96 to 44.75% on a dry matter basis; residual ether extracts 0.43% to 3.54%; crude fiber 9.6 to 19.06%; ash 5.89 to 12.92%; calcium 0.25 to 1.14%; and phosphorus 0.39 to 1.03%. Crude protein of RSM-WPN was 59.02% on a dry matter basis; residual ether extracts 0.76%; crude fiber 14.36%.

The correlation coefficients between chemical composition and DE (ME) of calibration samples in experiment 1 were presented in Table 5. The results indicated that the DE was negatively correlated with NDF ($r = -0.86$) and ADF ($r = -0.73$) contents and moderately

Table 6. Prediction equations of the DE (MJ/kg of DM) values of rapeseed meal according to neutral detergent fiber (NDF) and gross energy (GE) contents (% DM basis), experiment 1

Prediction equation	R ²	RSD	p-value
DE = 16.775-0.147×NDF	0.73	0.85	<0.01
DE = 11.848-0.131×NDF+0.231×GE	0.76	0.82	<0.01

correlated with gross energy (GE; $r = 0.56$) content in rapeseed meal calibration samples. In contrast, no significant correlations were found for CP, ether extract, CF, and ash contents. There were no significant correlations between ME and the other chemical compositions. The absolute correlation coefficients between ME and the other chemical compositions were less than 0.5, so it is not necessary to build a ME prediction model.

Prediction equations to the DE values of RSM according to NDF and GE contents were presented in Table 6. Two prediction models: DE = 16.775-0.147×NDF ($R^2 = 0.73$) and DE = 11.848-0.131×NDF+0.231×GE ($R^2 = 0.76$) were obtained. To test the suitability of these prediction models to predict the DE content of RSM, the DE and ME content of four samples of RSM were measured by both the *in vivo* method and prediction models in experiment 2 (Table 7). The maximum absolute difference between the *in vivo* DE determinations and the predicted DE values was

Table 7. Comparison of ME contents in rapeseed meal determined by using the *in vivo* method and prediction model, experiment 2

Test RSM	Observed DE (MJ/kg)	Equation 1: DE = 16.775-0.147×NDF			Equation 2: DE = 11.848-0.131×NDF+0.231×GE		
		Predicted DE (MJ/kg)	Difference (MJ/kg)	Difference (%)	Predicted DE (MJ/kg)	Difference (MJ/kg)	Difference (%)
RSM-XJ	11.23	11.52	0.29	2.59	11.75	0.52	4.63
RSM-HEN	11.19	10.56	-0.63	-5.63	10.79	-0.4	-3.58
RSM-FJ	12.26	12.25	-0.01	-0.09	12.43	0.17	1.39
RSM-SC	11.91	11.29	-0.62	-5.21	11.55	-0.36	-3.03
Mean	11.65	11.41			11.63		
SD	0.53	0.70			0.67		
p-value			0.3677			0.9419	

0.62 MJ/kg and the relative different was 5.21%.

DISCUSSION

The increased cost of the inclusion of SBM as a protein supply for pig diets has promoted interest into alternative sources of protein. One such substitute source is RSM. The presence of anti-nutritional factors such as glucosinolates, tannins, sinapine and erucic acid are found to affect the nutritional value of RSM as a protein source for pigs (Mawson et al., 1993). The potential of RSM as a pig feed component is also limited by its high crude fiber content, almost double that of SBM (124 vs. 60 g/kg; Sauvants et al., 2004) and this in turn reduces the energy value and nutrient digestibility of the diet (Fenwick, 1982). Yong-Gang Liu (1994) reported that results were obtained from more than 200 RSM samples from various oil mills in China. For screw-pressed cake and prepress-solvent extracted meal, respectively, crude protein averaged 389 and 432 g/kg on a dry matter basis; residual lipids 106 and 19 g/kg; crude fiber 132 and 138 g/kg; ash 87 and 99 g/kg; calcium 8 and 9 g/kg; phosphorus 11 and 12 g/kg. In the current study, excluding the protein-rich rapeseed meal production of RSM-WPN, the range of RSM crude protein were 38.04 to 44.75% on a dry matter basis; and crude fiber 9.6 to 19.06%. The crude protein content of RSM-WPN was 59.02%, and crude fiber 14.63%. The large range of chemical compositions of RSM used in the current study was benefit to the accuracy of the prediction model.

Noblet and Perez (1993) proposed prediction equations for the digestibility of nutrients and energy values of pig mixed diets from chemical analysis. The results showed that the DE was negatively correlated with NDF ($r = -0.80$), ADF ($r = -0.72$) and CF ($r = -0.71$) contents. In the current study, the DE was negatively correlated with NDF ($r = -0.86$) and ADF ($r = -0.73$) similarly, while barely correlated with CF ($r = -0.38$). According to the high correlation between DE and fiber content, the single predictor was always the fiber estimate. Among the different fiber estimates, the predictions with the lowest RSD were obtained with NDF alone as CF or ADF alone were inferior predictors. A few studies comparing different fiber criteria (King and Taverner, 1975; Morgan et al., 1987; Noblet et al., 1989) also concluded that NDF was a more accurate predictor than ADF or CF.

The DE of Canola RSM (*B. campestris*) was found to be 14.15 MJ/kg dry matter and energy digestibility was 71% (Bell et al., 1981). Bourdon and Aumaitre (1990) reported that the DE of high-glucosinolate RSM was 13.42 MJ/kg and energy digestibility was 68.5% based on dry matter. In our experiment 1, the average DE content of 22 RSM calibrations was 10.17 MJ/kg and the energy digestibility was 56% (data not shown). In our experiment 2, the average

DE content of 4 testing RSM was 11.65 MJ/kg and the energy digestibility was 59%.

The DE content is relative with the chemical content, physical property, rapeseed cultivar and quality, etc. The high fiber content in RSM originates mainly from a high hull content of about 25 to 30%. Rapeseed hulls contain about 60% NDF, of which lignin constitutes almost 50% (Grala et al., 1998). Mitaru et al. (1984) reported that the fiber of rapeseed hulls decreases ileal digestibilities of all nutrients in pigs. Studies carried out by De Lange et al. (1990) and Grala et al. (1998), demonstrated that rapeseed fiber (NDF) considerably affects ileal losses of dietary and endogenous nitrogen in pigs.

The nutritional value of rapeseed meal may vary among samples in China. Seneviratne et al. (2010) showed that extraction of oil from rapeseed meal in solvent-extraction plants was more efficient than in expeller pressing plants, resulting in a lower DE and ME content in solvent-extracted rapeseed meal. The process used for oil extraction could affect the nutrition value of RSM by leaving different amounts of oil or by decreasing the fiber content through dehulling (Bourdon et al., 1982). In the current study, to exclude the effects of different production procedures, we chose ten rapeseed meals taken from solvent-extraction production. Bayley et al. (1969) demonstrated the DE content of *B. campestris* RSM was 11.58 MJ/kg when tested as 40% of a corn-soybean diet fed to 45 to 90 kg gilts. When pelleted and reground, the DE value increased to 12.68 MJ/kg.

Bell (1975) found low DE of 1,750 kcal/kg (7.32 MJ/kg) for Bronowski RSM and 2,520 kcal/kg (10.54 MJ/kg) for a commercial sample of *B. campestris* RSM. It is reported that the Bronowski cultivar was not adapted to the region where the test sample was grown and consequently, the seed harvested was immature and of poor quality. The DE values shown do not reflect the possibly higher value of *B. campestris* RSM (3,210 kcal/kg, DM) compared with *B. napus* RSM (3,370 kcal/kg, DM) (Bush et al., 1978; Sharma et al., 1980).

In the current study, the maximum absolute difference between the *in vivo* DE determinations and the predicted DE values was 0.62 MJ/kg and the relative different was 5.21%. In the 10th revised edition of Nutrition Requirements of Swine (Subcommittee on Swine Nutrition Committee on Animal Nutrition. 1998), the Canola RSM (DM, 90%) has 21.2% NDF and 2,885 kcal/kg DE content; on a DM basis, the DE value is 3,205 kcal/kg (13.46 MJ/kg). Using the prediction model, $DE = 16.775 - 0.147 \times NDF$, we calculate the DE value as 13.66 MJ/kg. The absolute difference between the DE value of Nutrition Requirements of Swine and the predicted DE values was 0.20 MJ/kg, and the relative difference was 1.49%. Therefore, the two prediction models in this article can be used to predict the

DE content of RSM for growing-finishing pigs with acceptable accuracy.

Some specific equations for prediction of DE value of raw materials such as cereals (Wiseman and Cole, 1980), wheat by-products (Batterham et al., 1980a), sunflower meal (Perez et al., 1986), and cassava meal (INRA, 1984), etc. were reported. The comparison of different prediction equations is difficult because they do not propose the same predictors; they were not established from comparable sets of diets, and the digestibility measurements were obtained under different physiological conditions (body weight of the pigs, feeding level, etc.) (Noblet and Perez, 1993). It would be logical to favor the equations with the lowest residual standard deviation, practically applicable (low inter-laboratory standard deviation and cost of analysis), and established with a large number of chemically variable diets, each diet being analyzed by different laboratories (Noblet and Henry, 1993).

The reasons for these differences between measured energy values and those estimated from equations are unclear. However, it is relatively well established that the digestive utilization of diets is increased when body weight of animals increased and (or) feeding level is markedly reduced (Cunningham et al., 1962; Everts et al., 1986). In addition, the results (Noblet and Shi, 1993) indicated significant interactions existed between chemical characteristics of the diet (or dietary energy density) and body weight, physiological stage, or feeding level: the effect of these factors on DE values was negligible for highly digestible diets and was most important for low-energy diets. The RSM contained high CF content, so the effect of these factors on DE value can not be ignored. Furthermore, the dietary fiber indicators (e.g., NDF, ADF) are not easy to measure or values obtained in different labs may be rather different. The equations proposed from our experiments should then be used with caution. Additional measurements are required to confirm our findings and increase the number of samples per category.

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