

Calibration of a nutrient flow model of energy utilization by growing pigs

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A computational framework to represent energy utilization for body protein and lipid accretion by growing pigs is presented. Nutrient and metabolite flows, and the biochemical and biological processes which transform these, are explicitly represented in this nutritional process model. A calibration procedure to adjust the marginal input–output response is described, and applied, using reported experimental results, to determine a complete set of parameters for representing energy utilization by growing pigs. A reasonable value for minimum basal energy requirements is also determined. Although model inputs and outputs need not at any time be converted to equivalent energy flows, to facilitate comparison of model response with that of conventional energy-based models, a simple means to estimate energy flows from model-predicted nutrient flows is described. The well-known hierarchy of marginal (biological) energetic efficiencies with which pigs use different classes of nutrients is predicted by the model, based only on simple biological and biochemical principles. The significance of independent diet and metabolic effects on both energetic efficiency and maintenance requirements is examined using model predictions from simulated experiments.

Modelling: Nutrient flows: Calibration: Growing pigs

A conceptual framework for representing nutrient utilization by animals was presented in Birkett & de Lange (2001*a*). Based on this approach a computational structure with parameters applicable to growing monogastric animals was developed (Birkett & de Lange, 2001*b*). This model explicitly represents the material flows and transformations of nutrients and derived metabolites in terms of basic and functionally distinct nutritional processes: (1) intake, the acquisition of absorbed nutrients; (2) metabolic, the conversion of absorbed nutrients to energy-yielding and anabolic substrates; (3) faecal excretion of non-digestible materials; (4) urinary excretion of non-metabolizable materials; (5) production, the synthesis, degradation, and retention of body protein and lipid; (6) basal, the residual nutrient requirements not explicitly represented in the other processes. Energy requirements to drive these processes are met by ATP generated from the metabolite pools, providing a single calibration parameter for each of the main processes to adjust its input–output response. This present article describes a logical procedure for calibrating the monogastric model for a particular species, and applies this to derive appropriate parameters for a fully calibrated nutrient flow model of energy utilization by growing pigs.

Calibration parameters

Nutrient response for the primary pathways (Fig. 1) can be adjusted with six calibration parameters (Table 1). ATPd determines energy requirements for intake, digestion, and absorption of nutrients from faecal digestible DM (fDM). Intake of non-digestible DM (xDM) and its excretion as waste faecal material requires energy expressed in terms of ATPx. Faecal digestible nutrient intake is characterized in terms of crude protein (CP), crude fat (CFA), starch (ST), sugars (SU), and NSP. Ileal digestible amino acid (iAA) intake represents enzymatically-digested AA, which can meet basal AA requirements, provide the substrate for body protein retention (Pd, g/d) or are catabolized. Energy requirements that are (incrementally) related to Pd are represented by ATPp, including energy for protein retention and any associated metabolic processes that are incrementally dependent on production level, but excluding those associated with nutrient intake and faecal and urinary excretion. The energy requirements for urinary excretion are expressed in terms of ATPu and represent the excretion of N from all sources, and other products in urine. Synthesized triacylglycerol from either newly synthesized fatty acids

Abbreviations: AA, amino acid; ACA, acetyl-CoA; BE, basal free energy expenditure; BW, body weight; CFA, crude fat; CP, crude protein; DE, digestible energy; f, faecal digestible; FA, Fatty acid; FS, fermentable substrate; GL, glucose; i, ileal digestible; Ld, lipid retained; ME, metabolizable energy; Pd, protein retained; REL, retained energy in lipid; REP, retained energy in protein; ST, starch; SU, sugar; uN, urinary nitrogen; VFA, volatile fatty acid; x, non-digestible.

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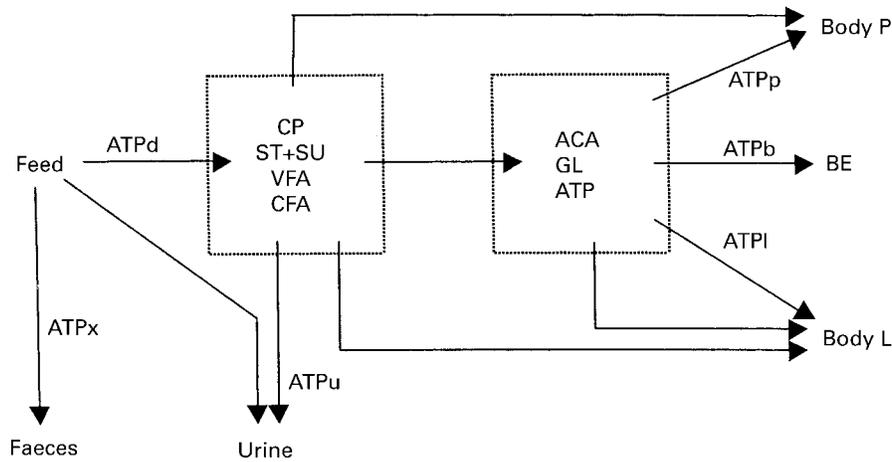


Fig. 1. Calibration parameters and main nutrient pathways. ATPd, intake, digestion, and absorption of nutrients; ATPx, processing of faecal non-digestible DM; ATPu, synthesis and excretion of urinary products; ATPi, lipid retention from both dietary and *de novo* sources; ATPp, protein retention; ATPb, basal ATP requirement; CP, crude protein; ST, starch; SU, sugar; VFA, volatile fatty acids; CFA, crude fat; ACA, acetyl-CoA; GL, glucose; P, protein; BE, basal free energy expenditure; L, lipid.

(FA) (assumed to be 18:1), or from dietary FA intake, is retained as body lipid (Ld, g/d) with an energy requirement expressed as ATPi (mol ATP/mol Ld), applied to Ld derived from either FA source. As for protein retention, ATPi includes any associated metabolic processes that are incrementally related to Ld and not separately explicitly represented. ST and SU are converted directly to glucose (GL), the primary metabolite in the model. A basal requirement for ATP, representing basal free energy (BE) expenditure, is expressed in terms of the parameter ATPb, calculated as the residual requirement after the requirements for intake, excretion, and production are taken into account. BE is assumed to be a function only of the animal's physical body state. Further details of the model structure can be found in Birkett & de Lange (2001a,b).

Simplifications and assumptions for growing pigs

The representation of nutrient utilization for monogastric animals described in Birkett & de Lange (2001b) can be simplified further when constructing a model for growing pigs. Various assumptions have been adopted for practical reasons, or because adequate information to support more complex concepts is not available:

1. Total tract digestibility of ST is usually found to be

very close to 100% in pigs (Bakker, 1996; Jorgenson *et al.* 1996). Furthermore, ST is almost completely digested at the terminal ileum, with ileal digestibilities of 96–100% measured for almost all ingredients. For some high-fibre diets both Bakker (1996) and Jorgenson *et al.* (1996) report slightly reduced ileal ST digestibilities, but never less than 95%. It is therefore not unreasonable to assume a 100% ileal digestibility for ST, unless results for a particular diet indicate that ST fermentation occurs to a significant extent.

2. The simplified representation of fermentation by an energy transformation from fermentable substrate (FS) to absorbed volatile fatty acids (VFA) is adopted, as described in the general model for monogastric animals (Birkett & de Lange, 2001b). Energetic efficiency from FS to VFA is fixed at 83% which is reasonable for pigs. Absorbed VFA are assumed to be acetic (C₂), propionic (C₃), and butyric (C₄) acids generated in the fixed molar ratios 70:25:5 (Bakker, 1996).
3. Ileal SU digestibility reported in Centraal Veevoeder Bureau (1998) is generally quite close to 100%, but, for some ingredients, values are given as low as 60%. It is reasonable to speculate that a lower energetic efficiency for some sugars (Noblet *et al.* 1989a, 1994)

Table 1. Calibration parameters for adjusting energetic response of the nutritional process model

Symbol	Units	Biological processes
ATPd	mol ATP/kg fDM intake	Intake of faecal digestible nutrients
ATPx	mol ATP/kg xDM excreted	Faecal excretion of non-digested materials
ATPu	mol ATP/mol N excreted	Urinary excretion of non-metabolizable materials
ATPp	mol ATP/g Pd	Protein retention and related metabolic processes
ATPi	mol ATP/mol Ld	Lipid retention and related metabolic processes
ATPb	mol ATP/d/kg body P	Basal energy requirement

fDM faecal digestible DM; xDM, non-digestible DM; Pd, protein retained; Ld, lipid retained; P, protein.

in some diets can be attributed to an increase in SU fermentation. This is represented in the pig model and requires that diet characterization includes separate fractions for ST and SU, and measures of ileal SU digestibilities. Alternatively, the fractions consisting of iST and iSU, and fermentable ST and SU intake combined, would also be sufficient.

4. Comprehensive, meaningful data are not available to characterize diets adequately and consistently with regard to FA, for true ileal digestibilities, endogenous losses, and net synthesis and utilization by intestinal microflora. Furthermore, ileal FA digestibilities are significantly and unpredictably affected by ingredient interactions and the FA profile (Jorgensen *et al.* 1992, 1993). Analytical methods, such as whether or not samples are hydrolysed prior to diethyl ether extraction, can also have a major impact on the measured digestibilities, as demonstrated by the two alternative sets of values reported in Noblet *et al.* (1989a). To reflect these considerations, the nutrient pathways related to CFA are simplified in the model as follows: (1) if reported faecal digestible crude fat (fCFA) is greater than iCFA, the difference, i.e. fermented fat, is ignored, and it is assumed that all fCFA is absorbed as enzymatically digested fatty acids (iFA); (2) if reported fCFA is less than the iCFA the implied net synthesis of fat is assumed to be derived on a one-to-one basis from fCFA, with no explicit energy requirement. These assumptions are accomplished by equating iCFA to fCFA and using fCFA to characterize the dietary available CFA intake.
5. Supported by the results for growing pigs reported in Birkett & de Lange (2001b), catabolism of FA is assumed to be zero, unless intake is so low that BE cannot be met from non-fat nutrient sources.

General calibration procedure

Eight distinct pathways are possible for the utilization of nutrients from the four main nutrient intake classes CP, ST+SU, NSP and CFA as energy substrates to support either Ld or Pd. Calibration of the marginal response with regard to these pathways consists of adjusting the five ATP parameters (ATPd, ATPx, ATPu, ATPp, and ATPl) until the incremental change in Ld and (or) Pd in response to a nutrient increment is consistent with observations from animal experiments. An implicit assumption of the nutritional process model is that the relative efficiency of utilizing energy derived from absorbed nutrients (CFA, ST, SU, AA, NSP) for synthesis of FA, as well as the generation of ATP, is determined by biochemical material conversions of metabolites: GL and acetyl-CoA (ACA). The use of absorbed nutrients to support Pd is represented by ATP requirements and material transformations of dietary AA to Pd. As a result the hierarchy of absorbed available nutrients (other than dietary AA) to support Pd is identical to their relative ability to generate ATP. Another important assumption is that the calibration parameters are constant, i.e. that the marginal energetic response to incremental

nutrient intake is independent of nutrient intake level. A careful procedure must be followed to avoid incorporating systematic errors into the model parameterization, for example because of inaccurate characterization of digestible nutrient intake or the observed retained protein and lipid in reported data. As far as possible model calibration should be based on clear experimental data which independently isolates the response pathways corresponding to each individual parameter determined.

As an alternative to calibrating nutrient response directly, an indirect approach can be used to take advantage of the extensive available literature data expressed in terms of energy flows. This requires that model variables be converted into equivalent energy flows, the product of nutrient material flow (g/d or mol/d) and nutrient energy density (kJ/g or kJ/mol). The marginal (dietary) energetic response is defined as the ratio of the increment in retained energy (in the form of either lipid or protein) to a corresponding increment in energy intake derived from (dietary) nutrient sources. This latter approach also allows for direct comparison to conventional models representing energy utilization.

For calculating marginal energetic response, energy intake is generally expressed in terms of either digestible energy (DE) or metabolizable energy (ME) intake. The marginal energetic response to ME intake is determined by partitioning ME between energy retained in protein (REP) and lipid (REL), and a residual term (ME_m):

$$ME = ME_m + (1/k_f) \times REL + (1/k_p) \times REP.$$

Reported values for the partial efficiencies k_f and k_p (Table 2) exhibit wide variation, and there are various conceptual and practical problems associated with obtaining an accurate and consistent measure of ME to characterize a diet (Birkett & de Lange, 2001a), particularly so if the incremental ME intake consists entirely of CP. These considerations might imply that solid experimental data to support the proposed calibration procedure are not available, making it difficult to achieve a robust model parameterization by simulating ME partitioning. However, the problems can be largely circumvented by using experimental data in which incremental intake of pure nutrients from non-protein sources is related to REL and REP. Reported marginal efficiencies are generally highly consistent in this case, in particular for utilizing (digestible) ST or CFA for REL.

The calibration procedure mimics the statistical regression of experimental data by constructing a simulated

Table 2. Reported marginal efficiencies of utilizing metabolizable energy for retained lipid (k_f) and protein (k_p)

k_f	k_p	Reference
0.80	0.60	Noblet <i>et al.</i> (1989b)
0.70	0.58	Close <i>et al.</i> (1973)
0.74	0.56	Agricultural Research Council (1981)
0.60	0.51	Tess (1981)
0.74	0.45	Pullar & Webster (1977)
0.75	0.44	Whittemore (1997)

response plane relating ME to REL and REP from partial linear relationships between ME and REL at fixed REP, and ME and REP at fixed REL. This procedure will be exact only when model-generated points are co-planar, i.e. the energetic response is linear with regard to both REP and REL as a function of ME intake. Consideration of the model calculation rules implies this linearity condition will generally be met exactly, or very closely, except for some situations in which the partitioning of nutrient intake between the main processes (Ld, Pd and ATP generation) does not change linearly with changes in nutrient intake level. This can occur, for instance, when AA intake is reduced below that required for Pd. In the case of such nonlinear response, constant k_f and k_p values are not possible.

Simulation of partial marginal responses requires pure increments in either Ld or Pd, something which can easily be forced in a model simulation, even though it is practically impossible to achieve with growing animals in experiments. For example, to simulate an incremental change in REL only, ME can be incremented while maintaining the predicted Pd at a fixed value by (artificially) manipulating the PdMax parameter, the maximum Pd according to animal constraints. Alternatively a pure REP increment can be simulated by adjusting REP using PdMax, then adjusting ME intake until REL returns to its initial value.

Calibration of marginal response proceeds as follows: (1) the two parameters ATPd and ATPi are determined simultaneously so that simulated marginal responses for increments of (digestible) GL and CFA are consistent with reported values; (2) a value for ATPx is determined from experimental data pertaining to the energetic cost of ingesting and excreting inert material; (3) reported observations on energy derived from protein sources are then taken into account to estimate a value for ATPu; (4) a value for ATPp is determined by adjusting it until simulated response with regard to both Pd and Ld is in agreement with observed response, i.e. simulated marginal efficiency from ME to REP is adjusted until it is close to the best estimate of k_p from literature values, using a simulated diet of similar composition to the experimental diets. Basal energy requirement is irrelevant when calibrating marginal response, provided the same BE value applies at both of the energy intake levels used to calculate the incremental ME intakes. Therefore any fixed value for ATPb can be maintained for simulations throughout the calibration procedure for marginal response.

The values reported here for the calibration parameters are intended only as reasonable estimates. Carefully controlled experiments directed at exploring the response for each nutrient pathway will be required to obtain more definitive parameter values (e.g. van Milgen *et al.* 2001).

Table 3. Energy densities of selected nutrients: calculated and literature values from various sources

	Energy density (kJ/g)				
	CP	CFA	ST	SU	NSP
Maize-soyabean meal base diet*	23.7	39.6			
100% Leucine†	31.7				
100% Threonine†	20.8				
Non-essential AA †‡	18.9				
Body protein and lipid§	23.7	39.6			
Whole-body protein	23.1				
Whole-body protein¶	23.3				
Protein and lipid**	23.5	39.3			
Protein and lipid††	23.8	39.6			
Protein and lipid‡‡	23.6	39.3			
Pure nutrients§§			17.35	15.6–16.5	15.6–26.3
Dietary nutrients	22.37	38.70	17.34	16.69	18.46
	(SE 0.72)	(SE 0.89)	(SE 0.21)	(SE 0.65)	(SE 0.46)
Digestible dietary nutrients	23.17	39.06	17.35	16.78	16.19
	(SE 0.49)	(SE 0.70)	(SE 0.14)	(0.40)	(SE 0.53)
Values used in model calibration	23.3	39.3	17.35	16.8	16.2

CP, crude protein; CFA; crude fat; ST, starch; SU, sugar; AA, amino acid.

* Values estimated from calculations based on AA and fatty acid contents of ingredients according to Centraal Veevoeder Bureau (1998), and heats of combustion of AA (Blaxter, 1989) and fatty acid (Critical Reviews in Chemistry, 1996). For details of composition of diet, see Table 4.

† Heats of combustion of free AA (Blaxter, 1989).

‡ Average heat of combustion of non-essential free AA (from Blaxter, 1989), weighted according to proportional content in typical maize-soyabean diet.

§ Agricultural Research Council (1981).

|| Estimated from whole-body AA composition (from Fuller, 1994), pig at 45 kg live weight, and free AA heats of combustion (from Blaxter, 1989).

¶ Estimated from whole body AA composition (from Kyriazakis *et al.* 1993; Moughan, 1999).

** Body and dietary protein and lipid according to Pullar & Webster (1977).

†† Body and dietary protein and lipid according to Kyriazakis & Emmans (1992) and Emmans (1994).

‡‡ Body and dietary protein and lipid according to Whittemore (1997).

§§ Obtained by bomb calorimetry of pure nutrient (Critical Reviews in Chemistry, 1996). SU values relate to constituent sugars; values for oligosaccharides are about 10% higher. Energy density of lignin is 26.3 kJ/g.

||| Calculated based on regression analysis of diet gross and digestible energy values (determined by bomb calorimetry) v. (digestible) nutrient composition of sixty-one experimental diets (Noblet *et al.* 1994).

Simulation of energy intake and retention

Simulation of digestible energy intake

For comparing the response of the nutritional process model to that of conventional models of energy utilization in growing pigs, energy intake and retention need to be simulated. Based on the intake and nutrient composition of the diet, fDM (kg/d) is converted to a DE intake (kJ/d) using fixed values for the energy densities (kJ/g) of the main nutrient classes, for which various values are reported in Table 3.

The energy content of free AA varies widely, from 31.7 kJ/g for leucine, to 20.8 kJ/g for threonine, and an average value of 18.9 kJ/g representing the weighted mean of non-essential AA. Thus one can expect variation in the energy density of CP according to AA composition, and particularly because of variation in the non-essential AA content of the CP. A value of 23.7 kJ/g CP is estimated for a typical pig diet, while a somewhat lower value of 23.3 kJ/g is obtained from regression analysis of DE contents for the sixty-one practical diets of Noblet *et al.* (1994). Whole-body CP in pigs is estimated to have an energy density of 23.1–23.3 kJ/g, using the whole-body AA composition reported from different sources (Blaxter, 1989; Kyriazakis *et al.* 1993; Moughan, 1999). Even though there are clearly differences between the energy density of dietary as compared with body CP, a common value is generally used as a compromise: 23.3 kJ/g has been assumed for the energy density of both dietary CP and Pd in the calibration of the model for growing pigs. These same considerations can be applied to CFA, which will vary (to a lesser extent than CP)

with FA composition. In the calibration a common value of 39.3 kJ/g is used for the energy density of both dietary CFA and body Ld.

Energy densities obtained for ST are very consistent, which is important because ST makes up the major energy component in many pig diets, and ST is used for a critical step in model calibration. For SU intake the heats of combustion of pentoses and hexoses are less than 16 kJ/g: fructose, glucose, galactose, arabinose, xylose are all about 15.6 kJ/g, while values for lactose and sucrose are higher at 16.5 kJ/g (Critical Reviews in Chemistry, 1996). The high average value 16.8 kJ/g obtained for SU in the regression (Table 3) reflects the fact that pure sugars will be present in the nutrient fraction as oligosaccharides, which have energy densities greater than those of their constituent sugars in the same way that the heat of combustion of ST (17.35 kJ/g) is higher than that of its constituent sugar glucose (15.64 kJ/g). SU content in commercial pig diets is generally <100 g/kg diet, and the composition is variable and generally unknown, so the assumed average value of 16.8 kJ/g appears to be a reasonable estimate. Variation in the composition of the NSP fraction will significantly affect its energy density; the difference between gross energy and DE densities determined for NSP can probably be attributed to different compositions of digestible and non-digestible fractions. For instance, the very low digestibility of lignin implies that most lignin present in gross energy will not contribute to DE. The heat of combustion of lignin (26.3 kJ/g) is much higher than that of the other carbohydrates, which largely explains the higher values

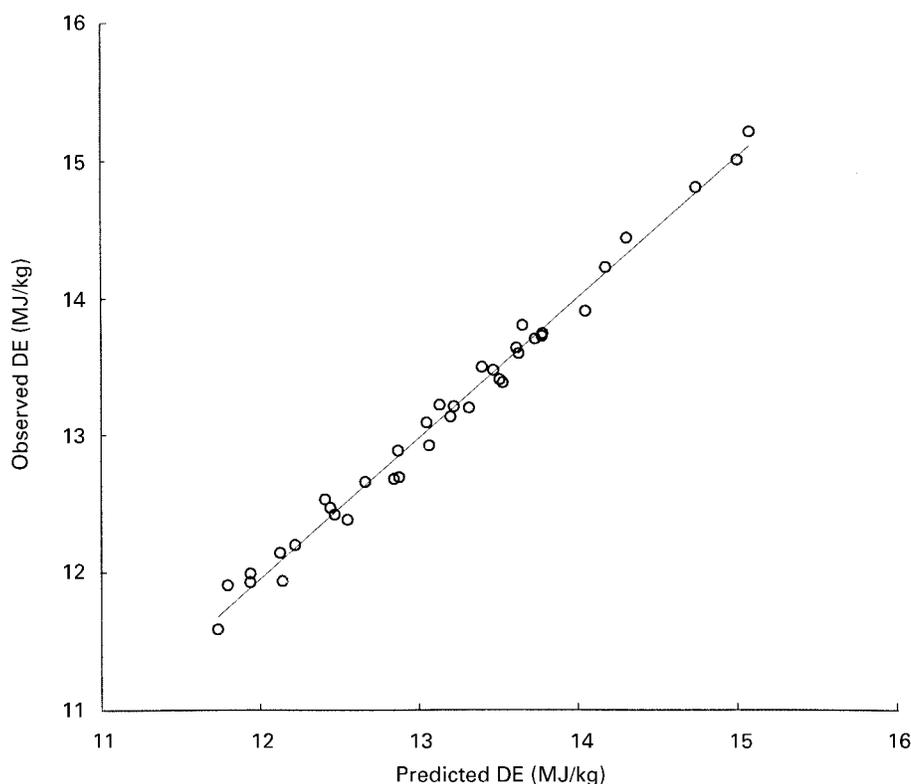


Fig. 2. Simulated diets from Noblet *et al.* (1989a). Predicted *v.* observed digestible energy (DE) values ($y = 1.03x - 0.35$, $R^2 0.99$).

obtained for NSP as compared with fNSP. Calculating the contribution of NSP to DE using a fixed average energy density is a reasonable approximation in the absence of further information on its composition.

To evaluate the prediction of DE content, diets were simulated using the stated ingredient compositions of the forty-one experimental diets of Noblet *et al.* (1989a), and ingredient nutrient composition data from Centraal Veevoeder Bureau (1998). The reported observed DE values are in good agreement (Fig. 2) with those predicted by calculation using the nutrient energy densities derived earlier, which may therefore be used with reasonable confidence in further comparison of model predictions to those of conventional energy-based models. It should be emphasized that the calibration procedure itself has been designed to rely only on the energy density values of CFA and ST, which are very well established and less variable than those of the other feed nutrients.

Simulation of metabolizable energy intake

To estimate simulated diet ME content for the nutritional process model requires an estimate of urinary and CH₄ energy losses. For pigs, simulated CH₄ energy losses are generally much smaller than urinary losses, and therefore, a reasonable enough estimate can be obtained from FS intake adapted from an equation of Kirchgeßner *et al.* (1991):

$$\text{CH}_4 \text{ energy loss (kJ/d)} = 160 + 0.041 \times \text{E(FS)} \\ (\text{kJ/d}), R^2 \text{ 0.59,}$$

where E(FS) is the energy intake in FS calculated using the energy densities estimated earlier.

For estimating urinary energy content, a relationship with urinary N (uN) content is often used, even though this is not

strong (see discussion in Birkett & de Lange, 2001a). A multiple regression with urinary C and N gives a better predictor for urinary energy (Hoffmann & Klein, 1980), however, since urinary C excretion is not predicted in the model, urinary energy content must be estimated from the simulated uN. For the calibration procedure a fixed value of 47.5 kJ/g uN (estimate from Hoffmann & Klein, 1980) is used and applied to uN derived from any source. The implications of variation in assumed urinary energy content are discussed in some detail below.

Simulated diet

The base diet used for simulations, with nutrient composition shown in Table 4, is formulated to represent a typical corn-soyabean meal pig diet. Simulated DE content is 14.25 kJ/g and lysine:CP ratio is 0.05. To avoid the non-linear response region, intake is determined so that nutrient supply is always sufficient to meet the requirements for the selected Pd level.

Calibration of marginal energetic response for pigs

Intake of faecal digestible nutrients (ATPd) and lipid deposition (ATPI)

The energetic efficiency of utilizing digestible dietary fat for Ld is consistently reported to be about 90% (Boyd & McCracken, 1979; Black, 1995; Whittemore, 1997), a value which implies a very low level of dietary fat catabolism. For supplemental intakes of ST and SU (GL, fructose, sucrose and lactose) the marginal energetic efficiency for Ld is reported to be 75% by Knox (1979), quoting Nehring *et al.* (1965), consistent with the efficiencies for ST and SU of between 74 and 76% given in both Armstrong (1969) and Black (1995). The first stage of model calibration is

Table 4. Nutrient composition of maize–soyabean meal base diet used for calibration simulations*

	Content (g/kg as fed)	Digestible content (g/kg as fed)	
		Apparent faecal	True ileal
DM	871	787	
Crude protein	175	149	146
Crude fat	32	22	
Starch	463	463	463
Sugars	32	32	27
NSP	127	82	
Lysine	8.5		7.3
Methionine	2.8		2.5
Methionine+cystine	5.8		5.0
Tryptophan	1.8		1.6
Histidine	4.8		4.3
Phenylalanine	8.6		7.6
Phenylalanine+tyrosine	14.9		13.1
Threonine	6.5		5.4
Leucine	15.9		14.0
Isoleucine	7.2		6.3
Valine	8.2		7.1
Arginine	11.0		10.1

* Simulated digestible energy content 14.25 kJ/g.

therefore based on achieving a simulated marginal energetic efficiency of 90 % for lipid retention from an increment of digestible CFA intake, and 75 % from digestible (non-fermented) ST and SU intake. This is accomplished by simultaneously adjusting the parameters ATPi and ATPd to find the unique solution for which the simulated marginal energetic efficiencies for Ld agree with these assumed values. The value of ATPx has a negligible influence on these simulated efficiencies and may be set to zero while adjusting ATPd and ATPi, because: (1) the digestibility of pure ST can reasonably be assumed to be 100 %; (2) due to the high energy density of CFA, moderate differences in digestibility have little effect on the simulated efficiency for CFA. The calibration of ATPd and ATPi is therefore essentially independent of ATPx.

After adjusting the two calibration parameters to obtain the assumed model response for incremental intakes of CFA and ST, ATPi is found to be 30 mol ATP/mol lipid retained and ATPd is 28 mol ATP/kg fDM. The minimum stoichiometric ATP requirement for synthesizing body lipid from FA and GY implies a (theoretical minimum) value for ATPi of 10 mol/d per mol lipid retained (Birkett & de Lange, 2001b). The calibrated value obtained for ATPd is therefore about three times the theoretical minimum for synthesis of Ld. Intake energy requirements as determined by the calibrated value for ATPd are quite large, and typically correspond to an ME cost of about 12–14 % total ME intake for the diet compositions of Noblet *et al.* (1989a).

Faecal excretion of non-digested materials (ATPx)

Müller & Kirchgeßner (1982) report an observed linear depressive effect on REL of 0.08 kJ in response to each kJ increment of 'non-digestible' straw added to a basal diet (energy digestibility for the straw supplement was 13 %). This data can be used to provide an independent estimate for the energy requirement of processing undigested xDM. Simulated straw increments were added to the DM intake of the base diet, and ATPx was adjusted until the simulated negative response in REL agreed with the experimental value reported. This procedure provides an estimated value for ATPx of 20 mol ATP/kg xDM. The energy requirements for processing xDM correspond to an ME cost of up to about 3 % total ME intake for the simulated treatments of Noblet *et al.* (1989a). On an energy basis the cost of processing xDM is typically about 10–12 % energy of the excreted xDM itself (kJ/kJ xDM), and this value varies with the energy density of xDM.

The independent estimates derived for ATPx and ATPd are consistent and imply that physical processing of nutrient intake accounts for about 70 % of the overall intake costs. This energy for intake processing must be provided whether or not the nutrients are eventually absorbed and subsequently metabolized. Carefully controlled experimental work is required to arrive at a definitive value for ATPx, but the present value will serve as a reasonable estimate, as confirmed by independent testing of direct model response to nutrient intake, using reported data from the forty-one diets of Noblet *et al.* (1989a), with widely varying non-digestible components (see Fig. 3).

Urinary excretion of non-metabolizable materials (ATPu)

Marginal efficiency is less easy to quantify for utilizing CP as an energy substrate, i.e. to generate the ATP and ACA to support Ld. For example, it is significantly affected by the AA composition of the protein, on account of the wide variation in biochemical efficiency of oxidation for different AA (Birkett & de Lange, 2001b). Reported experimental data generally relate to the measured response in terms of REL, rather than the marginal efficiency of ATP production *per se*. Kielanowski (1971) reports efficiencies of using CP intake for Ld in the range of 60–65 %, while Black (1995) and Whittemore (1997) both give a somewhat lower efficiency of 52–53 %. These values include the energetic effects of intake requirements, urea synthesis and urinary excretion, biochemical inefficiency of AA catabolism, and lipid retention. The actual ME cost related to urea excretion includes support costs for the excretion of both the N and the non-nitrogenous components of urine, various transport and metabolic processes that occur in the urea cycle, and other metabolic processes which are associated with urinary excretion (Birkett & de Lange, 2001b). Nevertheless, the ME cost for urinary excretion is closely related to uN, consistently reported to be about three times the theoretical value implied by the biochemical stoichiometry of urea synthesis. The parameter ATPu is therefore fixed in relation to the rate of uN excretion at 6 mol ATP/mol uN. Based on the previously determined values of ATPd, ATPi, and ATPx, this assumption provides a simulated marginal energetic efficiency of 56 % for REL derived from incremental dietary CP intake of the same AA composition and digestibility as that of the base diet, in agreement with the values reported earlier in the range of 50–65 %.

Fermentation

The marginal energetic efficiency for lipid retention from hindgut infusions of VFA is reported to be 65–71 % (Gädeken *et al.* 1989), 75–79 % (Roth *et al.* 1988), and 70 % (Müller & Kirchgeßner, 1991). The nutritional process model can be used to provide a simulated estimate for these experimental values by setting: (1) the fermentation efficiency factor temporarily to 100 %, so that an 'NSP supplement' then corresponds actually to a VFA supplement; (2) the ATPd parameter to zero, removing the intake costs and simulating a pure nutrient infusion into the hindgut. Faecal digestibility of VFA is close to 100 % (Kirchgeßner & Müller, 1991; Bakker, 1996) so simulations are not sensitive to ATPx. With these adjustments a simulated marginal energetic efficiency of 75 % is obtained, in good agreement with the observed experimental values cited earlier. (Actually a relatively small portion of the ATPd requirement will still apply to meet the energetic cost of absorption of VFA, so this simulated efficiency should be somewhat less.) Returning ATPd to the value determined in the calibration above lowers the simulated marginal efficiency to 63 %, a situation corresponding to dietary supplements of VFA which require ingestion, digestion, and absorption. Finally, returning the fermentation efficiency for deriving VFA from NSP to the assumed value of 83 %, a value of 53 % is obtained from the model for the marginal

efficiency of utilizing digestible NSP for REL. The faecal digestibility of NSP is very variable, typically not exceeding 60–65 % (Centraal Veevoeder Bureau, 1998), so a further (variable) reduction in simulated efficiency can be expected due to the effect of xDM and ATPx. The marginal energetic efficiency is lowered, for instance, to about 48 % for an NSP digestibility of 60 %. The implied range of simulated values for the marginal efficiency of utilizing ME from digestible NSP for REL is consistent with the range of the values reported in the literature: 43 % (Bakker, 1996), 59–63 % (Schiemann *et al.* 1989), 41–64 % (Kirchgeßner & Müller, 1991), 56 % (Jentsch *et al.* 1988), 59 % (Longland *et al.* 1989). These considerations indicate that no separate calibration parameter is required to represent the fermentation process adequately.

Protein retention (ATPp)

The biochemical stoichiometry of protein synthesis includes AA activation, initiation of peptide chain formation, elongation and termination. Armstrong (1969), Schulz (1978), van Es (1980), McDonald *et al.* (1995), and Blaxter (1989) all report energy requirements of 4–5 mol ATP for synthesizing 1 mol peptide bonds from constituent AA. Protein synthesis thus requires a minimum ME cost of some 4 kJ/g Pd, based on an estimated ME equivalent cost of 77 kJ/mol ATP and an average molar weight of 110 g/mol for body protein. Reeds *et al.* (1980) estimate that protein synthesis typically occurs at a rate between two and three times Pd, which suggests a minimum ME cost of between about 8 and 12 kJ/g Pd. Assuming no other requirement for energy would imply a maximum k_p of about 0.7, greater than any of the values reported in Table 2, which vary between about 0.44 and 0.60. Increments in various metabolic processes associated with an increase in Pd are likely responsible for observed k_p values significantly lower than the theoretical (maximum) k_p value of 0.7. Reeds *et al.* (1980) estimate a value for k_p of about 0.46, consistent with Webster *et al.* (1979), who estimate a heat increment associated with protein deposition of between 20 and 50 % total heat production, giving 52 kJ/g Pd using the mean of 35 %, or a k_p of 0.45. This value is also similar to the reported k_p of 0.44 in Pullar & Webster (1977), using rats fed a highly digestible, high protein (250 g CP/kg), synthetic diet, and a sophisticated experimental design that reduces the instability related to multi-collinearity in the regression analysis (Birkett & de Lange, 2001a).

Based on these considerations, and using a simulated high-protein diet (250 g CP/kg), the value of ATPp is adjusted until the simulated marginal energetic efficiency of using ME for Pd, i.e. the simulated k_p value, is 0.47. To avoid a non-linear response, care must be taken to ensure that protein retention at each level simulated is limited only by PdMax and not by AA intake. A value of 0.28 mol/g Pd is obtained for growing pigs when ATPp is adjusted until the simulated model marginal efficiency agrees with the assumed value. The minimum (theoretical) requirement for protein synthesis, 11 mol ATP/g Pd, and a typical synthesis rate of 2.2 times Pd (Reeds *et al.* 1980), would imply a value for ATPp of 0.10 mol/g Pd, therefore the

estimated value is about three times the theoretical one (as for ATPi).

There are, of course, some limitations to this calibration method for ATPp: the reported values for energetic efficiency k_p are quite variable (Birkett & de Lange (2001a); any assumption regarding k_p will ultimately influence the absolute model response (BE requirement); and it is assumed that the relative marginal efficiencies of using ME for REP are the same as those for supporting REL. Until more specific, highly controlled experimental data are available, the approach using simulated efficiency is probably the most reliable. Nevertheless, the estimate obtained for ATPp may be independently evaluated by comparing simulated and observed Ld and Pd response to nutrient intake using available datasets, without any need to consider partial efficiencies and the various problems associated with that concept (see p. 683).

Calibrating absolute energetic response

The last parameter (ATPb) representing BE is estimated based on comparing the absolute model-generated response, i.e. absolute RE in response to ME intake, to experimental observations (Birkett & de Lange, 2001b). Variation in BE can be anticipated, since it represents the residual energy expenditure. As a result, the practical application of the nutritional process model for predicting the growth of specific groups of animals under particular on-farm conditions will require ATPb to be adjusted so model predictions match production data for those specific conditions and animals. This provides a means to calibrate the model further for a particular (geno)type and production conditions, a procedure which is conceptually equivalent to adjusting maintenance energy when calibrating an energy-based model for particular on-farm conditions (de Lange & Schreurs, 1995). Therefore, the approach taken here is to determine a minimal BE (BE_{min}) as a function of the body protein content (P_{body}), with pigs kept under low levels of activity in stress-free and thermoneutral conditions (Birkett & de Lange, 2001b).

To illustrate the calibration of ATPb a minimum value was estimated using experimental data reported in Quiniou (1995), pertaining to pigs raised individually in metabolism crates. In this study N balances and energy expenditure measurements for three different genotypes (LP × PP castrates, LW castrates, LW × PP boars) are given at four different target live body weights (BW) (45, 65, 80 and 95 kg) for four different ME intake levels, expressed as 71, 80, 90 and 100 % voluntary ME intake, as determined by Quiniou (1995) for each genotype. Chemical body composition, in particular empty body protein (P_{body}) and lipid content, is reported at only the lowest and highest BW. Diets used for the different energy intake treatments are based on a common base diet, mixed at varying proportions with cornstarch. Simulated diets were formulated with the nutrient compositions stated in Quiniou (1995). Each of the treatments in experiment 3 of Quiniou (1995) was simulated using a dynamic growth model with an energy component based on the nutritional process model with marginal energetic response calibrated as described earlier.

Quiniou (1995) reports feed intake only at the target

BW, therefore feed intakes were represented with (model) calculated DE intake expressed as a percentage of reference National Research Council (1987) intake, adjusted for each simulation so that the (model) calculated ME intake at the target BW was identical to the reported ME intake for that BW. Furthermore, PdMax was adjusted to (artificially) force Pd into agreement with the reported value for that BW. Finally the value of BE was adjusted using ATPb until Ld was in agreement with the reported value when simulated BW was the target weight. This procedure ensures that all of the simulated ME intakes, Pd and Ld are the same as those reported for 45, 65, 80 and 95 kg BW. For each of these target weights, and for each energy intake level and genotype, the simulated residual BE requirement (mol ATP/d), and values of Pbody, BW and empty BW were obtained, a total of 48 points.

For each treatment, estimates for BE at the target BW were generated from simulations. These BE values were then correlated with variables that reflect the animal's body state, in particular Pbody, BW, and empty BW. Since Pbody is not reported at the two intermediate BW, it must be estimated with the model from predicted body composition at 65 and 80 kg BW, which requires growth simulation for four separate periods (20 kg BW was used as a starting weight) ending with each of the target weights.

Regression analysis was then used to relate BE to the body state variables. Each of Pbody, empty BW and BW is highly correlated with generated BE values, the best fit being obtained from a linear model with Pbody, with significant genotype specific (non-zero) intercept and slope. A reasonable estimate of BE for barrows can be obtained using the thirty-two combined model-generated points at BW of 45, 65, 80 and 95 kg for the two castrate genotypes. Linear and allometric models tested gave the following relationships:

$$\text{BE}_{\text{min}} \text{ (mol ATP/d)} = 40.4 + 4.23 \text{ Pbody (kg)},$$

$$R^2 \text{ 0.97, SEM 2.37,}$$

$$\text{BE}_{\text{min}} \text{ (mol ATP/d)} = 23.8 \text{ Pbody (kg)}^{0.54},$$

$$R^2 \text{ 0.96, SEM 2.53,}$$

which are indistinguishable in the experimental range 45–95 kg BW, although they diverge outside that range. Further evaluation comparing simulations and experimental data at BW below 45 kg and above 95 kg will be required to determine which BE_{min} regression is more appropriate, and to obtain more definitive values for the coefficients, although these will likely remain genotype specific.

Discussion

The practical use of the nutritional process model for quantitative simulation does not require conversion of nutrient intakes, retained protein and lipid, or ATP requirements, to their energy equivalents, even though energy flows were used for convenience in the calibration procedure. Nevertheless, to provide a basis for comparison of predictions to those of conventional energy-based models in the following discussion it is useful to be able to estimate ME equivalent costs for meeting simulated ATP requirements, i.e. the heat production associated with generating the ATP from dietary nutrient sources. The free energy released *in vivo* by breaking a high energy ATP bond varies somewhat according to thermodynamic conditions relating to temperature and the intercellular concentrations of ATP and ADP, but free energy is generally considered to be about 52 kJ/mol ATP (Burton, 1958; Armstrong, 1969). The nutrient composition of the ME used to generate ATP will affect the energetic efficiency, for example 74, 78 and 93 kJ ME/mol ATP being reported for ATP derived from GL, FA, and VFA respectively (Armstrong, 1969; van Es, 1980). For ME derived from the base diet of Table 4 an efficiency of 0.69 is predicted for generating ATP, equivalent to an ME cost of 75 kJ/mol ATP, and that value is assumed in the following discussion.

Marginal energetic response

Simulated marginal energetic efficiencies for deriving REL from ME are summarized in Table 5 for the main nutrient classes. These include values for the following scenarios: biochemical synthesis of lipid, pure nutrient infusions, dietary supplements assumed to be 100% digestible, and supplements with typical practical digestibilities. Simulated efficiencies obtained with the calibrated nutritional process model are in agreement with the range of reported values.

Table 5. Simulated marginal efficiencies of utilizing metabolizable energy (from different nutrient sources) for energy retained in lipid

	Calibration parameters				Simulated marginal efficiency ME to REL (%)				
	ATPd	ATP _x	ATP _u	ATPI	CFA	ST	CP	VFA	NSP
Biochemical synthesis*			✓		98	88	71†	78	
Pure nutrient infusions			✓	✓	95	85	69	75	
Diet sources (100% digestibility)	✓		✓	✓	90	75	57	63	53
Diet sources (typical digestibility)‡	✓	✓	✓	✓	89	75	56	63	48
Range of reported values (dietary digestible nutrients)	✓	✓	✓	✓	90	74–76	52–65	–	40–60

* Also including the urinary excretion cost associated with protein catabolism.

† Value reflects the actual (biological) energetic cost of N excretion with urine.

‡ Digestibilities assumed to be (%): CFA 80, fCP 85, NSP 60, VFA, ST and SU 100.

ME, metabolizable energy; REL, energy retained in lipid; ATPd, energy requirement for intake, digestion and absorption of nutrients from faecal digestible DM; ATPI, energy requirement for retention of body lipid; ATP_x, energy requirement for faecal excretion of non-digestible materials; ATP_u, energy requirement for urinary excretion of non-metabolizable materials; CFA, crude fat; ST, starch; CP, crude protein; VFA, volatile fatty acids; f, faecal.

The values reported in the second line of Table 5 indicate that the observed hierarchy of partial efficiencies (e.g. Noblet & Henry, 1991; Whittemore, 1997) for deriving REL in growing pigs from CFA, ST, CP or NSP is a consequence of combining biochemical efficiencies with the energy requirements to support the intake process (and also the cost of N excretion with urine). The actual efficiency values will vary with the energy requirement for lipid retention, but since the same value of ATP_I is applied to all nutrients the hierarchy will not be affected by it. Thus, it can be concluded that the hierarchical difference between calculated biochemical efficiencies and observed dietary efficiencies in pigs is primarily a consequence of the energy required to physically process the energy substrate itself, and the assumed relation that this is a function of nutrient mass, rather than energy density. This avoids the necessity for separate costs for L_d derived from diet FA and FA synthesized *de novo* (e.g. Emmans, 1994), which is seen to be a consequence of including the intake costs in the overall efficiency related to REL, thereby associating it with energy density rather than physical mass density of the nutrients processed.

Noblet *et al.* (1989a, 1994) report energy and N balances for pigs fed sixty-one different experimental diets of widely varied composition at two energy intake levels (high and low). These observations were made to study the quantitative effect of diet nutrient composition on the supply of 'useful' or net energy. The (high level) data

represent a valuable source of ME *v.* RE relationships for a wide variety of diet compositions and digestibilities (Birkett & de Lange, 2001a), which can be used as an independent test of the fully parameterized nutritional process model.

To evaluate the calibration procedure and the parameters determined for growing pigs, simulated diets were created to represent each of the forty-one diets reported in Noblet *et al.* (1989a), for which treatments detailed data were available. For each diet actual nutrient contents and digestibilities were used in the simulations. Missing diet information was derived from diet ingredient composition and ingredient characteristics according to Centraal Veevoeder Bureau (1998). Pd_{Max} was used to force the simulated Pd to match the observed value for each diet treatment, and the corresponding simulated L_d was recorded. Predicted L_d rates for the forty-one treatments were regressed against the observed L_d values reported in the dataset. The ATP_b parameter was adjusted once across all diets to determine the value which gave the best fit between predicted and observed L_d rates, by minimizing the standard error over the forty-one treatments. The relative hierarchy of L_d values across the forty-one treatments is not affected by this choice of ATP_b, since BW was very similar for all the animals in this study. Figure 3 shows observed *v.* predicted L_d values obtained using the nutritional process model calibrated with the parameters obtained above. The L_d values (*n* 41) have a reasonable standard error of prediction of 7 g/d.

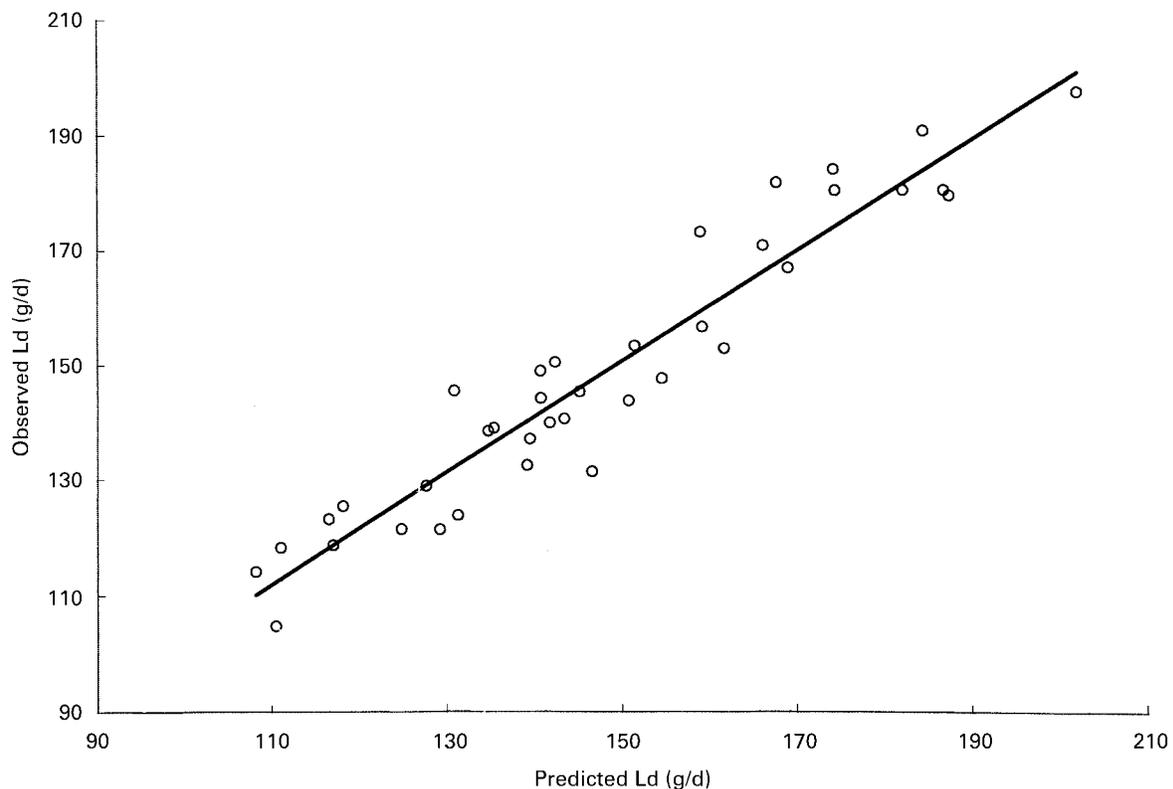


Fig. 3. Lipid retention (L_d) for growing pigs fed forty-one different experimental diets (Noblet *et al.* 1989a). Predicted *v.* observed L_d values ($y = 0.97x + 5.45$, R^2 0.91).

Protein and lipid retention

Combining the energy content of the Pd (23.3 kJ/g) with the energy requirements to support Pd, at an estimated equivalent ME cost of about 21 kJ/g Pd, gives a total ME cost for protein retention of about 44 kJ/g Pd, corresponding to a maximum theoretical biological efficiency for protein accretion of about 0.53. It can be estimated that about 50 % of the energy required to support Pd is not related directly to protein synthesis or turnover. In practice k_p will be reduced to between about 0.47 and 0.51, depending on the composition and digestibility of the ME as this affects the intake energy requirements and biochemical inefficiency of nutrient transformations. Energy requirements for Ld have an equivalent ME cost of about 2.5 kJ/g Ld, based on a typical molar weight of 900 g/mol Ld. Combining this value with the heat of combustion of the Ld (39.3 kJ/g) gives a total minimum ME cost for lipid retention of about 42 kJ/g Ld, very close to the value for Pd, and giving a theoretical upper limit of 0.94 for the marginal efficiency of utilizing ME for REL. In practice k_f will be reduced by other energy requirements to between about 0.62 to 0.71 for practical pig diets.

Simulated k_f and k_p values for ME derived from pig diets of different nutrient composition (Table 6) were calculated using multiple linear regression of model-generated ME against REL and REP data. Model response in terms of REL and REP is seen to be (essentially) multi-linear with regard to ME intake, therefore one can expect to obtain constant values for k_f and k_p . Corresponding values for $k_g = RE/(ME - MEm)$ are shown for incremental REL:REP ratios of 1.0, 2.0, and 3.0. The enhanced energetic efficiency for diets with increased CFA is evident in the increment in k_g of almost 3 percentage points associated with a moderate increase in CFA content from 3 to 13 %. On the other hand a depressive effect on simulated k_g is seen to be a consequence of increased diet content of either NSP, or CP of the same AA composition as that of the base diet. A more well-balanced CP intake, achieved by increasing the essential AA:non-essential AA ratio, enhances k_f , but reduces k_p , with little net effect on k_g . Simulated k_g is also enhanced by an increased lipid:protein ratio in the gain.

These dietary and animal effects on k_g are as anticipated and reflect biological reality. In some cases, the increase in k_g is associated with increases in both k_f and k_p , as these are affected by the efficiency of generating ATP from nutrient intake. In general, k_f is more variable than k_p , since it is also affected by alternative pathways for nutrients.

The complexities of interpreting experiments, important statistical issues, assumptions related to MEM, and the fact that REP is often considerably smaller than REL (Birkett & de Lange, 2001a), are no doubt responsible for considerable variation in experimental estimates of k_p . However, model-generated k_p values avoid these pitfalls to a large extent, and the comparatively small variation in the simulated k_p values of Table 6 may perhaps more accurately reflect biological reality. Alternatively, a better understanding of the factors which contribute to variation in the rate of protein synthesis as compared to retention, i.e. explicit modelling of protein turnover, might possibly generate more variation in simulated k_p values. There will always be some uncertainty in the simulated ME intake value related to the estimate of urinary energy content, an increase in assumed urinary energy density over the range from 33 to 60 kJ/g uN increasing k_f by 3–5 percentage points, and decreasing k_p by 1–3 percentage points. Further unpredictable effects may also result if different urinary energy contents apply at the two intake levels used to derive the incremental intake.

It is important to determine experimental data in the region where the animal's response to nutrient intake is linear. For example, the simulated response surface of ME v. REP and REL becomes noticeably curvilinear in the region where Pd is governed by AA limitation. For such situations k_f and k_p are not constant and will vary according to an animal's utilization of nutrient intake. Multiple linear regression of ME v. REP and REL using experimental data for which some Pd values are limited by AA intake is therefore not valid, and the non-linear response will affect the interpretation of k_f and k_p values. Experimental design and animal utilization of dietary nutrients must be given careful consideration in any examination of marginal energetic efficiencies, particularly when very low intakes are used.

Table 6. Simulated k_f , k_p and k_g values for various practical pig diets*

Diet	k_f		k_p		k_g (Ld:Pd ratio)†					
					1		2		3	
	%	+/-‡	%	+/-‡	%	+/-‡	%	+/-‡	%	+/-‡
Base	68.7		48.8		59.6		62.8		64.4	
130 g CFA/kg	71.6	+2.9	50.5	+1.7	62.0	+2.3	65.4	+2.5	67.0	+2.6
220 g NSP/kg 0.50	65.9	-2.8	47.1	-1.7	57.4	-2.3	60.4	-2.5	61.8	-2.5
250 g CP/kg (Lys:CP=0.05)	66.8	-1.9	47.6	-1.2	58.1	-1.6	61.2	-1.7	62.6	-1.7
130 g CP/kg (Lys:CP=0.06)§	70.0	+1.3	47.6	-1.2	59.6	-0.1	63.2	+0.4	65.0	+0.6

Ld, lipid retained; Pd, protein retained; CFA, crude fat; CP, crude protein; Lys, lysine.

* For details of composition of maize-soyabean base diet (175 g CP/kg), see Table 4. Other diets were formulated by adjusting the ST content to achieve the stated nutrient level.

† k_g values shown at three levels of Ld:Pd ratio (1,2 and 3 g/g).

‡ +/- denotes percentage increment with regard to values for base diet.

§ Non-essential amino acids replaced with starch.

|| Assumed 50% digestible.

Influence of digestibility

As discussed in Birkett & de Lange (2001a), the high intake level data of Noblet *et al.* (1989a) exhibit a diet quality bias, with a positive correlation between digestibility and feed intake. To illustrate the importance of ATPx, simulated Ld values were calculated for these forty-one high intake level dietary treatments, with ATPx set to zero and all other calibrated parameters unchanged. Under these conditions, i.e. ignoring the effects of xDM on efficiency, a slope of only 0.90 (R^2 0.90) is obtained in a linear regression of predicted against observed Ld values, as compared with a slope of 0.95 (R^2 0.91) obtained when the properly calibrated value for ATPx is used. Therefore, it can be concluded that the increased heat production associated with the lower quality, less digestible, diets cannot be completely predicted without considering the effects of non-digested material, thus confirming the essential significance of the ATPx parameter.

Energy requirements for processing non-digested material have an equivalent ME cost of about 1.5 kJ/kg xDM. The independently derived parameters ATPx and ATPd are consistent and their ratio (0.70) is biologically realistic and consistent with the definitions of the processes they drive. Emmans (1994) derives an ME cost for faecal organic matter of 3.8 kJ/g faecal organic matter, using data derived for cattle and poultry, extrapolating this to all species. In particular, the estimate is not based on a direct estimate for pigs (GC Emmans, personal communication). This ME cost of faecal organic matter is more than twice the equivalent ME cost estimated for processing xDM in the nutritional process model.

The effect of nutrient faecal digestibility on simulated marginal energetic efficiencies of utilizing different types of digestible nutrients for Ld are shown in Table 7 for various extreme diets in Noblet *et al.* (1989a). The high energy density of CFA constrains variation in k_f to a narrow range between 0.85 to 0.89, despite a very wide range of digestibilities from 38 to 75%. As a consequence of its low energy density, digestible NSP shows a large variation in k_f between 0.21 and 0.49, due to the effect of digestibility ranging from 19 to 67%. For CP the k_f of about 0.56 is largely unaffected by variation in CP digestibility over the typical narrow range of 70 to 94%. However, the assumed urinary energy density used in the calculation of ME intake has a major influence on the simulated k_f value for CP, which varies from 0.50 if a low 33 kJ/g uN is used, to 0.65 if

urinary energy is a high 60 kJ/g uN. This is a consequence of the relatively high energy requirement to support urinary excretion (ATPu), which is equivalent to an ME cost varying between 1 and 5% total ME intake in the diets of Noblet *et al.* (1989a). The fraction of a digestible nutrient intake which is fermented has a significant influence on simulated k_f values. For example k_f for SU varies from 0.64 to 0.74 as a consequence of varying the fraction of SU that is enzymatically digested as opposed to fermented over the range of 60 to 97%. These latter values were estimated from ingredient characterizations according to Centraal Veevoeder Bureau (1998), since ileal SU digestibilities were not determined by Noblet *et al.* (1989a).

Maintenance energy requirements

Tess (1981) gives a review of literature values of maintenance energy requirements and fasting heat production. Variation in reported MEM can be attributed to various experimental or statistical reasons (Birkett & de Lange, 2001a). In addition, metabolic factors can influence the relationship between MEM and BW, for example those which affect the efficiency of utilization of nutrients for generating ATP, such as diet composition, or which affect the lipid:protein ratio in the gain, such as feeding level, BW, AA intake, or pig genotype. Typical values for MEM in growing pigs are reported in the range 800–900 kJ/kg^{0.60} per d, for instance 825 kJ/kg^{0.60} per d (Agricultural Research Council, 1981; Quiniou, 1995), although values as large as 1000 kJ/kg^{0.60} per d have been suggested (Noblet *et al.* 1991). For the purpose of facilitating comparison between MEM values from various sources, which are expressed with metabolic BW exponents ranging from 0.54 to 0.75, values were converted to a common metabolic BW exponent of 0.60 (using a basis of 50 kg BW).

The effect on MEM due to variation in partitioning ME intake can be illustrated by calculating MEM from model simulations, at various BW and for pigs with different Pd values. A BE value of $1.3 \times \text{BE}_{\text{min}}$, typical for pigs kept under on-farm production conditions, was assumed. Model-generated values of MEM, calculated as:

$$\text{MEM} = \text{ME} - \text{REL}/k_f - \text{REP}/k_p,$$

are shown in Table 8, for BW of 20, 50, 80, and 110 kg, using ME derived from the base diet of Table 4, with efficiencies of $k_f = 0.69$ and $k_p = 0.49$. The simulated values are consistent with reported values (Agricultural Research Council, 1981; Quiniou, 1995). The two pig types shown in Table 8, low v. high Pd, have different simulated MEM at the same BW, on account of the difference in lipid:protein ratio in the gain, and its effect on the efficiency (k_g) of nutrient utilization for RE. To illustrate the dependence of MEM on diet composition, simulated MEM values were generated for diets with various changes to the nutrient composition of the base diet: (1) with 100 g ST/kg replaced by CFA, MEM was about 4% lower than those shown in Table 8; (2) with CP increased to 250 g/kg at the expense of ST, and no change in the AA composition of CP, MEM was increased by about 4%; (3) with 100 g ST/kg replaced by crude fibre (faecal digestibility 0.50), MEM was

Table 7. Effect of faecal digestibility on simulated partial efficiency k_f for various nutrients

	Faecal digestibility		Simulated k_f	
	Min	Max	Min	Max
CFA	0.38	0.75	0.85	0.89
NSP	0.19	0.67	0.21	0.49
CP	0.70	0.94	0.56	0.57

Min, minimum; Max, maximum; CFA, crude fat; CP, crude protein.

Table 8. Simulated metabolizable energy for maintenance requirement (ME_m) at four live body weights for two pig genotypes*†

	ME _m ‡							
	BW (kg/Pd 135 g/d)				BW (kg/Pd 220 g/d)			
	20	50	80	110	20	50	80	110
MJ/d	5.2	8.6	10.9	13.0	5.0	8.5	11.5	13.6
kJ/d per kg BW ^{0.75}	550	460	410	380	530	450	430	400
kJ/d per kg BW ^{0.60}	860	820	790	780	830	810	830	810
kJ/d per kg BW ^{0.54}	1000	1000	1000	1000	1000	1000	1100	1100

BW, body weight; Pd, protein retained; Ld, lipid retained.

* Basal free energy expenditure (BE) = 1.3 × BE_m, where BE_m = 23.8 × body protein (kg)^{0.54}.

† For details of composition of diet, see Table 4 (diet was fed at 90% *ad libitum* intake, according to the National Research Council (1987)).

‡ ME_m calculated as ME intake – REL/k_l – REP/k_p, with k_p = 0.69 and k_l = 0.49, where ME is metabolizable energy intake, REL is energy retained in lipid and REP is energy retained in protein.

increased by 3%. These simulated variations in ME_m are related only to changes in metabolic and biochemical processes and their associated energy requirements and efficiencies.

In experiments with pigs under fasting conditions long enough to achieve a stable heat production, van Es (1972) reports an estimated fasting heat production equivalent to 575 kJ/d/kg^{0.60}. In a series of experiments designed to measure fasting heat production in six different genotypes of pigs, van Milgen *et al.* (1998) report fasting heat production values of about 700 kJ/kg^{0.60} per d, when corrected to resting conditions. In comparison, BE requirements obtained for simulated experiments are generally in the range of 7.3 to 8.3 mol ATP/kg^{0.60} per d, and BE_m is equivalent to about 7.1 mol ATP/kg^{0.60} per d. Thus, model-generated BE is generally in the range of about 7–8 mol ATP/kg^{0.60} per d, equivalent to free energy requirement in the range from 360 to 420 kJ/kg^{0.60} per d. This range of free energy values is consistent with the definition of BE and the range of heat production estimates reported for fasting heat production (600–700 kJ/kg^{0.60} per d) and ME_m (800 to 900 kJ/kg^{0.60} per d).

Conclusions

A calibration procedure has been described that can be used to adjust the marginal and absolute energetic response of a nutritional process model to match the response of growing pigs to nutrient intake. The parameters obtained are intended to be reasonable estimates, however further studies will be required for a more definitive calibration. The values given illustrate that the calibration procedure is capable of parameterizing the model effectively and meaningfully, as shown by the evaluation of simulated response against independent datasets.

Models based on factorial analysis of ME intake emphasize animal response, but cannot account for variation in the contribution of the various nutrients to energy intake. Current net energy systems can account for variation in dietary nutrient composition, but not in the animal's utilization of that net energy intake. The calibrated nutritional process model described here can account for variation in both dietary nutrient composition and metabolic

utilization of nutrient intake. Linear programming optimization methods for least cost feed formulation are no longer possible with the nutrient process model, since the net energy supplied by feed ingredients is not constant for a given nutrient composition and will vary according to the use of nutrients for the various body functions. However, the integration of feed formulation with animal response provided by the new model provides a different, more sophisticated optimization capability.

This nutritional process model represents animal biology explicitly, thereby associating heat production with biological reality and the various processes which are responsible for it. The well-known hierarchy of biological marginal energetic efficiencies for Ld derived from different nutrient classes can be predicted by the model, as a consequence only of energy density, digestibility, and biochemical stoichiometry.

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