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Prediction of cellular ATP generation from foods in the adult human - application to developing specialist weight-loss foods

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Abstract

For the accurate prediction of the potential 'available energy' of a food at the cellular level (i.e. ATP generation from food) it is necessary to be able to predict both the quantity and location of uptake (upper-tract or colon) for each energy-yielding nutrient. The objective was to develop a valid model ('Combined Model') for predicting the (potential) ATP available to the body from absorbed nutrients across the total digestive tract. The model was intended for the adult human under conditions where energy intake ≤ energy expenditure and all absorbed nutrients are catabolised. The development of the model involved two parts: (i) the experimental development of a dual *in vivo* − *in vitro* digestibility assay ('dual digestibility assay') to predict human upper-tract nutrient digestibility, as modelled by the rat upper digestive tract, and colonic digestibility, as predicted by fermenting rat ileal digesta in an *in vitro* digestion system containing human faecal bacteria; and (ii) the development of a series of mathematical equations to predict the net ATP yielded during the post-absorptive catabolism of each absorbed nutrient at the cellular level.

A strong correlation (r=0.953, P=0.047) was found between total tract organic matter digestibility (OMD), as predicted with the newly developed dual *in vivo* – *in vitro* digestibility assay and with that determined in a metabolic study with humans for four mixed diets ranging considerably in nutrient content. There were no statistically significant (P>0.05) differences for mean OMD between the predicted and determined values for any of the diets.

The Combined Model (dual *in vivo* – *in vitro* digestibility assay + stoichiometric predictive equations) was applied to three meal replacement formulations and was successfully able to differentiate between the diets in terms of both energy digestibility and predicted ATP yields. When the energy content of each diet was compared to that of a baseline food (dextrin), some metabolisable energy (ME) models gave considerably different ratios compared to that predicted by the Combined Model. By way of example, for Diet C a ratio of 0.96 (Atwater and FDA models) was found

versus 0.75 (Combined Model). Thus, the model has practical application for predicting dietary available energy content, particularly in the research and development of specialised weight-loss foods, where it may be more accurate than some current ME models. Uniquely, the Combined Model is able to define a food in terms of ATP content (mol ATP / g food) using recent estimates of cellular P/O ratios and therefore, directly relates dietary energy intake to the quantity and form (ATP) of energy ultimately delivered at the cellular level.

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Abbreviations

ΔG Free Energy **AA** Amino Acid

AA_d Amino Acids Present In The Diet

AA_i Amino Acids Present In The Ileal Digesta

ADH Alcohol Dehydrogenase
ADP Adenosine Diphosphate

Ala Alanine

AMG Amyloglucosidase

AMP Adenosine Monophosphate

ANOVA Analysis Of Variance

AOAC Association Of Analytical Chemists

Arg Arginine
Asn Asparagine
Asp Aspartic Acid

ATP Adenosine Triphosphate

ATP_a Available ATP (ATP Yield)

ATP Yield From Amino Acids

ATP_d ATP Cost Of Digestion

ATP Yield From Fatty Acids

ATP_{net} Net ATP Yield

ATP Yield From Short Chain Fatty Acids

ATP_{ST+SU} ATP Yield From Starch And Sugars

ATP_t ATP Cost Of Absorption / Transportation

BMR Basal Metabolic Rate

BW Body Weight
CHO Carbohydrate
CoA Coenzyme A

CV Coefficient Of Variation

Cys Cysteine

dE Digestibility Of Energy

DE Digestible Energy

DF Dietary Fibre

dHE Increment Of Heat Energy

DIT Dietary Induced Thermogenesis

DM Dry Matter

DMD Dry Matter Digestibility

ER Endoplasmic Reticulum

EtOH Ethanol **FA** Fatty Acid

FA_d Fatty Acids Present In The DietFADH₂ Flavin Adenine Dinucleotide H₂

FA_i Fatty Acids Present In The Ileal Digesta

FAO Food And Agriculture Organization

FDA Food And Drug Administration

FFA Free Fatty Acid

FID Flame Ionisation Detector

GaE Gaseous Energy

GC Gas Chromatography

GE Gross Energy

GI Gastrointestinal

GL GlucoseGln Glutamine

Glu Glutamic Acid

Gly Glycine

GP Glycerol Phosphate

GTP Guanosine Triphosphate

GY Glycerol

HE Heat EnergyHF High FibreHis Histidine

IE Intake Energy

IleIsoleucineLeuLeucine

LF Low Fibre

LPL Lipoprotein Lipase

Lys Lysine

ME Metabolisable Energy

MEOS Microsomal Ethanol Oxidising System

Met Methionine
MF Mixed Fibre

NADH Nicotinamide Adenine Dinucleotide H

NADPH Nicotinamide Adenine Dinucleotide Phosphate H

N_d Nitrogen Present In The Diet

NDF Neutral Detergent Fibre

NE Net Energy

NEAT Non-Exercise Activity Thermogenesis

NEFA Non-Esterfied Fatty Acids

Nitrogen Present In The Ileal Digesta

NR Not Reported

NSP Non-Starch Polysaccharide

OFN Oxygen-Free Nitrogen

OM Organic Matter

OMD Organic Matter Digestibility

OM_{D+F} Organic Matter That Is Digested And Fermented By The Body

OM_i Organic Matter Present In The Ileal Digesta

 $\mathbf{OM}_{\mathrm{uf}}$ Unfermented Organic Matter At The End Of Incubation

P P-Value (Probability)

PE Pectin

PEG Polyethylene Glycol

Phe Phenylalanine

Pro Proline

PSP Phenolsulphonphthalein

PVTC Post-Valve T-Caecum

r Correlation Coefficient

RE Retained Energy

RMR Resting Metabolic Rate

RS Resistant Starch

SAPU Small Animal Production Unit

SCFA Short Chain Fatty Acid

SE Standard Error

SE Surface Energy

SEM Standard Error Of The Mean

Ser Serine

SI Le Système International D'unités

ST Starch

Starch Present In The Diet

Starch Present In The Ileal Digesta

SU Sugars

SU_d Sugars Present In The Diet

SU_i Sugars Present In The Ileal Digesta

TAG Triacylglycerol

TAG_d Triacylglycerol Present In The Diet

TAG_i Triacylglycerol Present In The Ileal Digesta

TCA Tricarboxylic Acid

TEE Total Energy Expenditure

Thr Threonine

Trp Tryptophan

Tyr Tyrosine

UC Unavailable Carbohydrate

UCP Uncoupling Protein

UE Urinary Energy

UV Ultraviolet

Val Valine

VFA Volatile Fatty Acid

VLDL Very Low-Density Lipoprotein

WB Wheat Bran

WHO World Health Organization

Preface

After ingestion, the energy-providing nutrients in food (carbohydrate, fats, protein, and for some individuals, ethanol) undergo a series of catabolic reactions in the human digestive tract, and then (primarily) in hepatocytes to release energy from their chemical bonds. This energy then becomes available to the body, primarily in the form of ATP (the universal currency of chemical energy in the body) and is subsequently converted into other forms of energy such as mechanical energy, thermic energy and so on. However, not all of the energy present in ingested food is ultimately converted to ATP due to the energy requirements involved with the digestion, absorption and intermediary metabolism of food, which vary with the type of food and the nutrients ingested. Some energy is also lost through the heat of fermentation of undigested dietary material entering the large intestine. Furthermore, nutrients vary in their degree of digestibility and absorption (i.e. uptake from the gut) and the efficiency by which they yield energy that is ultimately useful to the body (net ATP gains), with the energy made available to the body via short chain fatty acids from nutrients fermented in the hindgut being less than that obtained from direct nutrient uptake in the uppertract. For the accurate prediction of the potential 'available energy' (ATP) at the cellular level it is therefore important to be able to predict both the quantity and location of uptake (upper-tract or colon) for each nutrient. The use of metabolisable energy (ME) systems (e.g. Atwater system), as commonly used for food labelling purposes, may not be the most appropriate or accurate means of predicting the useful energy at the cellular level because amongst other weaknesses, ME systems do not account for the unique features of each diet, such as differences in digestibility or inter-nutrient interactions that may affect nutrient assimilation. A valid alternative means needs to be found to model and predict the available energy content of a food for the research and development of foods required to deliver a specific quantity of energy to the body at the cellular level, such as those specifically designed for weightloss. The need for such foods is growing in importance due to the rapid increase in overweight and obese persons in recent years.