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# Protein quantity, not protein quality, accelerates whole-body leucine kinetics and the acute-phase response during acute infection in marasmic Malawian children

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The present study compared leucine kinetics and acute-phase-protein concentrations in three groups of marasmic, acutely infected Malawian children fed one of three isoenergetic diets. These were: an enhanced-protein-quality diet (egg-white+tryptophan, providing  $1\cdot2$  g protein/kg per d; n 14); an increased-protein-content diet (egg-white+tryptophan, providing  $1\cdot8$  g protein/kg per d; n 14); a standard-protein diet ( $1\cdot2$  g milk protein/kg per d; n 25). The hypotheses tested were that children receiving a diet with more protein would have greater rates of non-oxidative leucine disposal and that children receiving an isonitrogenous diet with a higher protein quality would have lower rates of leucine oxidation. The children were studied after 24 h of therapy using standard [ $^{13}$ C]leucine stable-isotope tracer techniques. The children receiving the higher-protein-content diet had greater leucine kinetic rates than those receiving the standard-protein-content diet; non-oxidative leucine disposal was 170 (sp 52) v. 122 (sp 30)  $\mu$  mol leucine/kg per h (P<0.01). Leucine oxidation was less in the children receiving the enhanced-protein-quality diet than in those receiving the standard-protein-quality diet; 34 (sp 12) v. 45 (sp 13)  $\mu$ mol leucine/kg per h (P<0.05). The children receiving the high-protein-content diet increased their serum concentration for five of six acute-phase proteins 24 h after starting therapy, while those receiving the standard-protein-content diet did not. These data suggest that there was greater whole-body protein synthesis, and a more vigorous acute-phase response associated with the higher-protein-content diet. The clinical benefits associated with a higher protein intake in marasmic, acutely infected children need further study.

Malnutrition: Therapy: Protein: Acute-phase response

Severe malnutrition and acute systemic infection are important determinants of whole-body protein kinetics. When dietary amino acid intake is reduced, such as occurs in malnutrition, there is decreased protein turnover (Jackson et al. 1983; Garlick et al. 1991). Proteolysis is reduced in malnutrition as well, which conserves the limited quantities of amino acids (Soares et al. 1994). Standard dietary recommendations for the initial treatment of children with severe malnutrition prescribe a very low total protein intake, 0.9-1.2 g/kg per d (World Health Organization, 1999), which may also contribute to the reduced rates of protein turnover. Acute infection, however, increases protein turnover (Long et al. 1977), probably by activating the acute-phase response, the accelerated synthesis of a group of proteins useful to the host in successfully responding to the infection (Szalai et al. 1995; Gabay & Kushner, 1999).

Acute infection and malnutrition are often coincident and synergistic in children (Scrimshaw & SanGiovanni, 1997).

In marasmic children with acute systemic infection receiving standard nutritional therapy, whole-body protein kinetics does not accelerate, and the acute-phase response is blunted (Manary *et al.* 2004). This blunted acute-phase response may well contribute to the compromised immune response typical of severe malnutrition. We tested two potential therapeutic strategies for overcoming this blunted acute-phase response in marasmus and acute infection. The first strategy was to provide dietary protein of higher quality, but in a standard amount. Higher dietary protein quality meant that the amino acid composition of the dietary protein was more similar to that of acute-phase proteins than was the standard milk protein. The second strategy was to provide a larger quantity of dietary protein.

The present study used a standard [<sup>13</sup>C]leucine stable-isotope tracer dilution method (Matthews *et al.* 1980) to measure whole-body leucine kinetics in marasmic Malawian children with acute infection receiving either a diet with a standard total protein content, but a higher protein

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quality, a diet with a higher total protein content or the standard milk-protein diet. Specifically, the hypotheses tested were that children receiving an isonitrogenous diet with a higher protein quality would have a lower rate of leucine oxidation, but no more non-oxidative disposal of leucine, than those receiving standard therapy, while children receiving a diet with more protein would have a greater rate of non-oxidative leucine disposal.

## Methods

#### Subjects

Children aged 12-60 months with marasmus admitted to Queen Elizabeth Central Hospital in Blantyre, Malawi from August 2000 to July 2002 were eligible. Marasmus was defined as having weight-for-age < 60 % of the international standard (set by the Wellcome Trust Working Party; Anonymous, 1970) and evidence of wasting, manifest by weight-for-height < 80 % of the international standard. After mothers gave their informed consent for their child's participation, each child was admitted to a special metabolic ward, which provided more intensive nursing care, better parenteral antibiotics, more frequent feedings and more careful clinical monitoring than the hospital ward. The initial evaluation of these children included blood culture, urine culture obtained by sterile catheter, chest X-ray, thick blood smear for malaria parasites and an ELISA for HIV (Vironostika HIV; Organon Teknika, Durham, NC, USA). Every child received parenteral ceftriaxone for the first 48 h after admission. Acute infection was defined as sepsis, malaria or pneumonia. Sepsis was defined as clinical signs of sepsis with a positive blood or urine culture, malaria was defined as clinical signs of falciparum malaria with a positive smear for malaria parasites, and pneumonia was defined as cough and tachypnea with a focal infiltrate on chest X-ray. The infections were believed to be acute because, according to the mothers' reports, each child's clinical condition had worsened within the day before admission. Children were said to recover if the symptoms of their acute infection resolved, their clinical status improved during their hospitalisation and they began to gain weight. The study was approved by the Human Studies Committee of Washington University in St Louis, MO and the College of Medicine Research Committee of the University of Malawi.

## Diets

The standard diet provided 350 kJ (85 kcal)/kg per d and 1·2 g milk protein/kg per d. The diet with improved amino acid quality provided the same amount of energy and protein, but used egg-white with an added 1·5 % tryptophan as a protein source. The high-protein diet provided 350 kJ/kg per d and 1·8 g protein/kg per d; egg-white with 1·5 % tryptophan was used as the dietary protein source. On the basis of the amino acid composition of a typical mixture of acute-phase proteins the fraction of each essential amino acid provided by the diet is shown in Table 1 (Reeds *et al.* 1994). Note that in the case of the standard milk diet, tryptophan is the limiting amino acid and the diet provides a complete complement of amino acids sufficient to synthesise 71 %

of a typical acute-phase response. For the egg-white +tryptophan diet providing a standard amount of protein, histidine is the limiting amino acid and the diet provides a complete complement of amino acids sufficient to synthesise 82% of a typical acute-phase response. The high-protein egg-white+tryptophan diet provides at least 120% of every essential amino acid needed for a typical acute-phase response. All children received a mineral—multivitamin supplement added to the feeding. Children were prospectively assigned to receive one of these diets by systematic allocation. Feedings were administered in equal amounts per kg body weight every 2 h, and children unable to take feedings by mouth were fed through a nasogastric tube.

## Metabolic study

The protocol and calculations for the metabolic study have been previously described and validated from a subset of the children included in the present report (Manary et al. 2002, 2003, 2004). Briefly, at 19h after admission each child's CO<sub>2</sub> appearance rate was determined with a primed (2·5 μmol/kg), constant (5 μmol/kg per h) 70 min infusion of [<sup>13</sup>C]sodium bicarbonate (99 % <sup>13</sup>C; Cambridge Isotopes, Andover, MA, USA). Before starting the [13C]sodium bicarbonate infusion breath samples were collected using a silicone rubber face mask connected to a non-diffusing gas collection bag (Hans Rudolph, Kansas City, MO, USA) to determine the baseline <sup>13</sup>CO<sub>2</sub> isotope abundance. Breath samples were collected 60, 65 and 70 min after the initiation of the [13C]sodium bicarbonate infusion for the measurement of the CO<sub>2</sub> appearance rate. Immediately following the collection of these breath samples, a primed constant intravenous infusion of [1-13C]leucine (99 % 13C; Cambridge Isotopes, Andover, MA, USA) was begun. Each child received [1-13C]leucine (prime 15.3 \(\mu\)mol/kg, infusion 4 \(\mu\)mol/kg per h) for 5 h. During the 5 h infusion children received feedings in small portions every 30 min. The amount of each feed was such that it provided the same rate of dietary energy intake (350 kJ/kg per d) as the 2 h feedings. This was done so that the metabolic measurements would reflect protein kinetics in the fed state rather than the postabsorptive state. At 4.5 and 5 h after the start of the leucine infusion, which corresponded to 24 h after admission and the initiation of feedings, a 1 ml blood sample was drawn to measure plasma  $[^{13}C]\alpha$ -ketoisocaproic acid enrichment. Three breath samples were collected 4.5, 4.75 and 5 h after the start of the leucine infusion for the measurement of <sup>13</sup>CO<sub>2</sub> isotope abundance, representative of leucine oxidation.

Sera for the measurement of the concentrations of six acute-phase proteins, C-reactive protein, properdin factor B, C3, haptoglobin,  $\alpha_1$ -anti-trypsin, and  $\alpha_1$ -acid glycoprotein were drawn on admission, 24 h after admission and 48 h after admission.

## Sample analyses

In 10 ml samples of breath, <sup>13</sup>CO<sub>2</sub>: <sup>12</sup>CO<sub>2</sub> abundance was measured using an automated gas isotope ratio mass spectrometer (Finnigan MAT Delta+ XL; Bremen, Germany; Manary *et al.* 2003).

Table 1. Essential amino acids provided for the typical acute-phase response in the milk and egg-white+tryptophan diets

Amino acid	Milk diet providing 1·2 g protein/kg per d	Egg-white+tryptophan diet providing 1·2 g protein/kg per d	Egg-white+tryptophan diet providing 1⋅8 g protein/kg per d	Estimated amino acid requirements for a typical acute-phase response (mg/kg)
Tryptophan (%)	71	525	555	24
Threonine (%)	84	84	126	65
Isoleucine (%)	135	126	189	54
Leucine (%)	132	114	171	89
Lysine (%)	106	91	136	90
Methionine (%)	131	180	270	23
Cysteine (%)	80	223	334	14
Phenylalanine (%)	74	89	134	79
Tyrosine (%)	106	85	128	55
Valine (%)	120	115	172	67
Arginine (%)	81	121	181	54
Histidine (%)	99	82	123	33

<sup>%,</sup> The fraction of amino acid provided in the diet to meet the estimated amino acid requirement for a typical acute-phase response (Reeds et al. 1994); the amount is listed in the last column.

Plasma  $\alpha$ -ketoisocaproic acid, the intracellular deamination product of leucine, was used to estimate intracellular [ $^{13}$ C]leucine enrichment. Serum samples were analysed by GC electron impact-quadrupole MS (Agilent Technologies 6890N capillary GC system and 5973 Network Mass Selection Detector; Foster City, CA, USA) after the  $\alpha$ -ketoisocaproic acid was converted to its quinoxalinol trimethylsilyl derivative (Manary *et al.* 1997).

The concentrations of acute-phase proteins were measured by rate nephelometry (Beckmann Image Nephelometry System; Beckmann, High Wycombe, UK).

# Calculations and statistics

The rates of appearance (Ra) of leucine and  $CO_2$  were calculated from the following equation, derived from a simple mass balance (Matthews *et al.* 1980):

$$Ra = ((Ei/Ep) - 1) \times I$$
,

where Ei is the isotopic enrichment of the tracer infused (99%), Ep is the isotopic enrichment of the tracer in serum or breath, and I is the infusion rate of the tracer.

The rate of leucine oxidation was estimated from the following equation:

$$\begin{aligned} Ra_{Leuox} &= (Ra_{CO_2}) \times (^{13}CO_2 \text{ enrichment in breath}) \\ &\times (Ei/Ep-1)/81, \end{aligned}$$

where  $Ra_{CO2}$  is the Ra of  $CO_2$  in breath, Ei is the isotopic purity of the leucine infused (98–99%), and Ep is the plasma [ $^{13}C$ ] $\alpha$ -ketoisocaproic acid enrichment averaged over  $4\cdot5-5$  h. The factor  $0\cdot81$  accounts for the fraction of  $CO_2$  that is produced by leucine oxidation, but not released from the body bicarbonate pool into the breath as determined by Matthews *et al.* (1980).

Leucine derived from whole-body proteolysis was calculated by subtracting the rate of leucine intake provided in the diet and [<sup>13</sup>C]leucine tracer infusion from Ra<sub>Leu</sub>, and non-oxidative leucine disposal was calculated by subtracting the rate of leucine oxidation from Ra<sub>Leu</sub>. Leucine derived from whole-body proteolysis and non-oxidative

leucine disposal are indices of whole-body proteolysis and protein synthesis respectively.

The estimated sample size was thirteen children in each group, assuming that protein kinetics would vary similarly as children with kwashiorkor (Manary et al. 1997) and that differences of 10 % between groups would be detected with 95% specificity and 80% power. Data are expressed as mean values and standard deviations. Anthropometric Z scores were calculated using Epi Info 2000 (WHO/Centers for Disease Control, Atlanta, GA, USA). The protein kinetic data and acute-phase protein concentrations were tested for normality using the Kolmogorov-Smirnov test. After determining that the data were normally distributed, ANOVA was used to compare leucine kinetics between the groups of children and Student's paired t test was used to compare acute-phase proteins within the groups over time. Statistical differences of P < 0.05 were considered to be significant. The children that received the standard milk-protein diet have had their protein kinetic data reported previously (Manary et al. 2004).

## Results

Fifty-three children were studied, with similar demographic and nutritional characteristics (Table 2). All twenty-eight children receiving either the high- or low-protein egg-white+tryptophan diet recovered, while of the twenty-five children receiving the milk-protein diet, twenty-two recovered and three died.

The constant [<sup>13</sup>C]sodium bicarbonate infusion reached an isotopic steady-state in breath <sup>13</sup>CO<sub>2</sub> after 55 min of infusion and the plasma [<sup>13</sup>C]leucine enrichment equilibrated within 4 h as previously documented in a subset of these children (Manary *et al.* 2002, 2003).

The rates of non-oxidative leucine disposal and leucine derived from whole-body proteolysis were similar in the children receiving 1·2 g protein/kg per d as either milk protein or egg-white+tryptophan, but greater in the children receiving 1·8 g protein/kg per d as egg-white+tryptophan (Fig. 1). The leucine oxidation rate was higher when the diet of lower protein quality (milk) was consumed than

**Table 2.** Demographic, anthropometric and clinical characteristics of study children (Mean values and standard deviations)

	Milk diet providing 1·2 g protein/kg per d		Egg-white+tryptophan diet providing 1·2 g protein/kg per d		Egg-white+tryptophan diet providing 1⋅8 g protein/kg per d	
	Mean	SD	Mean	SD	Mean	SD
Males	8		7		8	
Females	17		7		6	
Age (months)	31	12	35	19	38	20
Weight-for-age Z score	<b>−4.1</b>	0.6	-4.3	0.5	-4.3	0.5
Height-for-age Z score	<b>−3.4</b>	1.0	-3.5	1.2	-3.6	1.2
Weight-for-height Z score	-2.9	0.6	<b>−3</b> ·1	0⋅8	-3.0	0.9
Clinical infections*						
Pneumonia	12		7		8	
Sepsis	8		5		5	
Malaria	10		3		3	

<sup>\*</sup> For clinical infections the number of children listed with each infection is given; some children had more than one infection.

it was when the diet of higher protein quality (egg-white +tryptophan) was consumed (Fig. 1).

In the children receiving the higher-protein-content diet, five of the six acute-phase protein concentrations (C-reactive protein, C3, properdin factor B, haptoglobin and  $\alpha_1$ -acid glycoprotein) increased from admission to 24 h (Table 3; P < 0.05). For the children receiving either of the lower-protein-content diets there were no significant differences in the concentration of any of the acute-phase proteins when those on admission were compared with 24 h measurements.

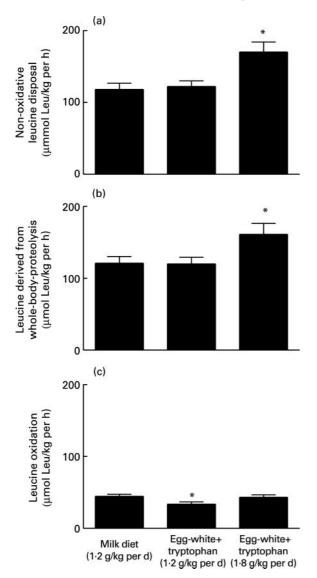
# Discussion

Marasmic, acutely infected children who received the highprotein diet had greater rates of non-oxidative leucine disposal, suggesting that there was greater whole-body protein synthesis, and they exhibited a more vigorous acute-phase response as measured by serum acute-phase protein concentrations. The marasmic, acutely infected children who received the isonitrogenous diet of presumed higher protein quality oxidised leucine at a slower rate, which supports the notion that they were wasting less amino acids, however their total non-oxidative leucine disposal rate was similar to the children receiving the standard milk-protein diet.

The present study assumes that whole-body leucine kinetics mirrors whole-body protein kinetics, an assumption that has been made in a variety of human metabolic investigations. Leucine solutions for human infusion are safe, easy to prepare and stable (Bodamer et al. 1997). Clinical methods for their administration are robust. Kinetic calculations using leucine enrichments are relatively straightforward, because leucine is an essential amino acid with quantifiable intracellular fates; incorporation into proteins or deamination and subsequent oxidative decarboxylation (Matthews et al. 1980). As a tracer, [13C]leucine is inexpensive, and the laboratory methods for its analyses are standardised and simple. The fraction of leucine in a typical mixture of acute-phase proteins is 10.5 %, so leucine turnover is substantial during acute infection (Reeds et al. 1994). For these reasons we chose [13C]leucine as the tracer for the present study, and we believe that leucine kinetics mirrors whole-body protein kinetics in malnutrition and acute infection.

The estimation of the rate of CO<sub>2</sub> production includes a factor to account for the CO2 which is produced, but not expired; this factor is based on experimental work done in adults and infants (Spear et al. 1995; Kien & McClead, 1996). This factor has been measured in diverse populations, including critically ill children, and varies between 0.70 and 0.85 (Hoerr et al. 1989; Bresson et al. 1990; Tissot et al. 1993) although most reports have identified values between 0.77 and 0.82 (including those from critically ill infants). We did not measure the amount of CO<sub>2</sub> produced that was then incorporated into other metabolic substrates in our population and this may have limited the accuracy of the present results. Another limitation of the study is that we measured simply the concentrations of acute-phase proteins, while others have measured their actual rate of synthesis (Jahoor et al. 1999). This would have required longer isotope infusions and much larger amounts of blood to be drawn from the subjects, which unfortunately was not possible in our context. No study subject had oedema at the time of their metabolic investigation, and thus these results may well not reflect protein metabolism in kwashiorkor and acute infection. Indeed whole-body protein kinetics in the two types of severe malnutrition, kwashiorkor and marasmus, has previously been shown to be different (Manary et al.

The estimations of amino acids needed for a typical acute phase were the only consideration in choosing the amino acid composition of the diet in the present study. In the time frame that was studied, the first 24 h of acute infection, it may be reasonable to consider amino acid requirements in this simple manner; synthesis of visceral proteins is suppressed and cell-mediated immune responses are just beginning to accelerate. However, whole-body protein synthesis is a dynamic and complex process which is incompletely understood; thus it is unlikely that amino acid requirements can be accurately estimated using such simplistic balance calculations. Several amino acids could potentially be limiting, and empirical data must be relied upon to determine how varying amino acid composition



**Fig. 1.** Mean rates of non-oxidative leucine disposal (a), leucine derived from whole-body proteolysis (b) and leucine oxidation (c) in marasmic Malawian children with acute infection fed one of three isoenergetic diets: a standard diet with milk protein providing 1·2 g protein/kg per d; an improved-protein-quality diet with egg-white with added tryptophan providing 1·2 g protein/kg per d; a higher-protein-content diet with egg-white with added tryptophan providing 1·8 g protein/kg per d. Values are means, with standard errors of the mean represented by vertical bars. \*Mean value was greater than those for the two other groups (*P*<0.05; ANOVA).

affects measurements of the immune response. Further studies on key amino acids hypothesised to be limiting protein synthesis during acute infection and malnutrition could be conducted by measuring non-oxidative leucine disposal in appropriate subjects assigned to receive diets that differ only in the amount of the single, key amino acid. The dietary threshold amount for a key amino acid could be determined by measuring the rate of non-oxidative leucine disposal and finding the point at which the dietary amount of the key amino acid was no longer associated with an increase in non-oxidative leucine disposal. Such studies may help define which amino acids are truly limiting in malnutrition and acute infection.

It is interesting that while the severity of malnutrition, measured by weight-for-height Z score, was similar between the children receiving either the standard- or higher-protein-content diets, the children receiving more dietary protein had greater rates of leucine derived from whole-body proteolysis. How might increased dietary protein intake stimulate proteolysis? Ubiquitin-proteosomemediated protein degradation increases with physiological stress, such as acute infection, and is stimulated by glucocorticoids and pro-inflammatory cytokines, such as TNFa and IL-1 (Hasselgren, 2000; Kee et al. 2003). Ubiquitinmediated proteolysis is not regulated by free amino concentrations, unlike lysosomal-mediated proteolysis (Kadowaki & Kanazawa, 2003). In animal models, prolonged starvation suppresses ubiquitin activity (Martin et al. 2002). Acutely infected, marasmic children have been shown to have elevated levels of pro-inflammatory cytokines (Manary et al. 2004). Perhaps the higher dietary protein intake in these marasmic, infected children ameliorates the factors that are suppressing ubiquitin-mediated proteolysis in malnutrition, and allows proteolysis to be stimulated by the stress of acute infection.

The pattern of changes in acute-phase protein concentration observed suggests that giving more dietary protein during malnutrition and acute infection results in a more vigorous acute-phase response. The serum acute-phase-protein concentrations among the marasmic children receiving 1.8 g protein/kg per d were similar to those in well-nourished Malawian children with acute infection (Manary *et al.* 2004).

The recommendations for dietary protein intake during the initial phase of severe childhood malnutrition are 0.9-1.2 g/kg per d, and they are accompanied by the caveat that if larger amounts of energy and protein are given 'the child may develop a serious metabolic imbal-(World Health Organization, 1999). Likewise, Waterlow (1992) recommends that 0.7 g protein/kg per d be given initially, and states that larger quantities of any nutrient run the risk of 'overloading the system at both the physiological and biochemical levels'. These recommendations were developed by the thoughtful experts with extensive experience treating severe malnutrition in the developing world (Golden, 2002). Comparisons of clinical outcomes or metabolic parameters of diets providing different quantities of protein have not been published. Reports from starvation victims in 1945 fed veryhigh-protein diets, diets with protein:energy ratios >30%, indicate that upon refeeding many of these individuals died (Burger et al. 1945; Lipscomb, 1945). Collins et al. (1998) compared therapeutic feeding in malnourished adults with either a diet with a protein:energy ratio of 16.4% or a diet with a protein: energy ratio of 8.5% and found that the lower-protein-content diet was associated with reduced mortality. The standard recommendations for children call for a protein: energy ratio of 2-4%, much lower than was used in these studies of adult malnutrition. While it seems that diets with very high protein intakes, >5 g protein/kg per d, during the initial phase of treatment are dangerous, our data suggest that very low protein intakes may not provide enough amino acids for appropriate metabolic responses to stress.

**Table 3.** Acute-phase protein concentrations (Mean values and standard deviations)

	Milk diet providing 1.2 g protein/kg per d		Egg-white+tryptophan diet providing 1⋅2 g protein/kg per d		Egg-white+tryptophan diet providing 1.8 g protein/kg per d	
Acute-phase protein	Mean	SD	Mean	SD	Mean	SD
C-reactive protein (mg/l)						
On admission	72	50	78	68	75	59
After 24 h	63	58	46	46	85*	52
After 48 h	39	37	25	27	35	33
C3(g/l)						
On admission	0.71	0.42	0.86	0.39	0.78	0.40
After 24 h	0.73	0.42	0.86	0.50	0.98*	0.69
After 48 h	0.73	0.40	0.73	0.40	1.01*	0.63
Properdin factor B (g/l)						
On admission	0.42	0.17	0.51	0.18	0.46	0.18
After 24 h	0.41	0⋅15	0.59	0.19	0.71*	0.43
After 48 h	0.39	0⋅15	0.51	0.12	0.68*	0.38
Haptoglobin (g/l)						
On admission	0.77	0.95	0.84	0.75	0.80	0.85
After 24 h	0.89	1.10	0.83	0.73	0⋅87*	1.03
After 48 h	0.78	0.96	0.73	0.62	0.74	0.84
$\alpha_1$ -Acid glycoprotein (g/l)						
On admission	2.6	0.4	2.9	0.7	2.7	0.6
After 24 h	2.6	0.4	3.0	0⋅8	3⋅1*	1.0
After 48 h	2.5	0⋅5	2.7	0.9	2.6	0⋅8
$\alpha_1$ -Antitrypsin (g/l)						
On admission	2.3	0.6	2.3	0.7	2.4	0⋅6
After 24 h	2.3	0.6	2.0	0⋅8	2⋅5	0.9
After 48 h	2.3	0.6	1.7	0⋅8	2.6	1.1

<sup>\*</sup>The concentrations of C-reactive protein, C3, properdin factor B, haptoglobin and α<sub>1</sub>-acid glycoprotein for the children receiving the high-protein diet increased from admission to 24 h or 48 h (*P*<0.05).

It is not known if the metabolic benefits associated with the higher-protein diet seen in the present study translate into genuine clinical benefits in the treatment of marasmic children, and a proper clinical trial of a higher-protein diet is needed to answer this question.

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