





Whole-blood PUFA and associations with markers of nutritional and health status in acutely malnourished children in Cambodia

Sanne Sigh^{1,2,*} , Lotte Lauritzen¹ , Frank T Wieringa³, Arnaud Lailou⁴, Chhoun Chamnan², Nheb Angkeabos⁵, Duch Moniboth⁵, Jacques Berger³, Ken D Stark⁶ and Nanna Roos¹

¹Department of Nutrition, Exercise, and Sports, Faculty of Science, University of Copenhagen, Frederiksberg, Denmark; ²Department of Fisheries Post-Harvest Technologies and Quality Control, Fisheries Administration, #186 Preah Norodom Boulevard, 12101 Phnom Penh, Cambodia; ³Institut de recherche pour le développement (IRD)/ Université de Montpellier/SupAgro, Montpellier, France; ⁴Department of Child Survival and Development, UNICEF Cambodia, Phnom Penh, Cambodia; ⁵National Pediatric Hospital, Phnom Penh, Cambodia; ⁶Department of Kinesiology, University of Waterloo, Waterloo, Ontario, Canada

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Abstract

Objective: To measure fatty acid composition, particularly whole-blood PUFA content, in acutely malnourished children and identify associations with markers of nutritional and health status.

Design: PUFA were assessed in dried blood spots obtained from a cross-sectional study. Nutritional and health status were assessed by anthropometry, haemoglobinopathies, inflammation and blood counts.

Setting: Cambodia.

Participants: The study was conducted with 174 children aged 0.5–18 years with acute malnutrition.

Results: Among total fatty acids (FA), the relative percentage of total PUFA was 20% FA, with 14% of the children having very low PUFA (mead acid (MA): arachidonic acid (AA) >0.02, *n*-6 docosapentaenoic acid:DHA >0.2 and total *n*-6:*n*-3 PUFA >10.5). Wasting was not associated with any PUFA. Stunting and low height were consistently positively associated with total PUFA and positively with *n*-6 PUFA. Height was positively associated with *n*-3 long-chain PUFA (LCPUFA). The presence of haemoglobinopathies or inflammation was positively associated with MA:AA, but not total PUFA. Elevated blood platelet counts were positively correlated with linoleic acid and appeared to be influenced by anaemia ($P=0.010$) and inflammation ($P=0.002$). Monocyte counts were high during inflammation ($P=0.052$) and correlated positively with *n*-6 LCPUFA and *n*-3 LCPUFA.

Conclusions: Children with acute malnutrition or stunting had low PUFA, while elevated platelets and monocytes were associated with high PUFA. In acutely malnourished children, inflammation could lead to elevated blood cell counts resulting in increased whole-blood PUFA which does not reflect dietary intake or nutritional status.

Keywords

PUFA
Blood cell counts
Acutely malnourished children
Nutritional status
Inflammation

Acute malnutrition is a prevalent public health problem in many low- and middle-income countries. It contributes significantly to childhood morbidity and mortality and affects more than 70 million children annually^(1,2). Acute malnutrition is partly caused by insufficient intakes of energy and nutrients for maintaining essential body functions. The food consumption in low- and middle-income countries

is often a starch-based diet such as rice, legumes, fruits and vegetables that is low in PUFA^(3,4).

Dietary linoleic acid (18:2*n*-6; LA) and α -linolenic acid (18:3*n*-3; ALA) are essential fatty acids and can be metabolized to arachidonic acid (20:4*n*-6; AA) and DHA (22:6*n*-3), respectively. These long-chain PUFA (LCPUFA) are critical for healthy growth, immune function, brain

*Corresponding author: Email sannesigh@gmail.com



development and the skin–water barrier^(5–7). Acutely malnourished children may be particularly vulnerable to essential fatty acid deficiency owing to low food consumption, disturbed lipid metabolism and absorption, and/or diarrhoea caused by intestinal infections^(8–10).

In children with moderate acute malnutrition (MAM) and/or severe acute malnutrition (SAM), PUFA status has been shown to be altered in different blood pools such as sterols⁽¹¹⁾, triacylglycerols⁽¹²⁾, plasma^(13,14), erythrocytes⁽¹⁵⁾ and whole blood^(16–18). Whole-blood samples have logistical advantages for field research collection due to less sample processing immediately post collection, as well as the ability to collect dried blood spots. However, whole blood includes plasma and various types of blood cells that can differentially impact the overall fatty acid composition⁽¹⁹⁾. Acutely malnourished children living under poor conditions are vulnerable to infections, inflammation, anaemia, and other nutritional and health challenges^(10,20–24) that could affect blood cell counts, such as increased monocytes with infection^(25,26). Changes in cell counts can affect the overall fatty acid composition of whole blood, which could alter interpretations of whole-blood PUFA status. Another health challenge that is prevalent in South-East Asia is genetic Hb disorders (haemoglobinopathies), which may contribute to changes in whole-blood PUFA composition due to increased risk of anaemia and altered erythrocytes caused by these disorders^(27,28).

The purpose of the present study was to investigate the fatty acid composition, particularly the PUFA content, of whole blood and potential associations with markers of nutritional and health status in acutely malnourished Cambodian children.

Methods

Study design and setting

The present cross-sectional study uses baseline data from an 8-week interventional trial evaluating the effectiveness of a locally produced, fish-based, ready-to-use therapeutic food compared with a standard ready-to-use therapeutic food during outpatient treatment of acute malnutrition in Cambodian children (called the ‘SAM trial’)^(29,30). The SAM trial was conducted at the National Pediatric Hospital in Phnom Penh, Cambodia. The sample size in the trial was based on the superiority of one of two ready-to-use therapeutic foods on the main outcome, weight gain (g/kg per d). To detect a 10% difference with a standard deviation of 0.7 g/kg per d, a sample size of forty-nine in each group was required to achieve 80% power ($\alpha = 0.05$, two-sided). To allow for dropouts, a total of 120 children were enrolled⁽²⁹⁾. Initial recruitment was of 0.5–18-year-old children. After recruiting fifty-four children, the recruitment criterion for age was adjusted to enrol only 6–59-month-old children due to

limited funding and on request of the funding agency (UNICEF). The final sample was comprised of the fifty-four children aged up to 18 years and 120 children aged 6–59 months.

Eligibility and nutritional markers

The eligibility criteria for enrolment in the present study were children aged 0.5–18 years with non-complicated acute malnutrition, defined as weight-for-height Z-score (WHZ) or BMI-for-age Z-score (BAZ) of ≤ -2.8 . Children younger than 60 months were also re-enrolled based on mid-upper-arm circumference (MUAC) of ≤ 115 mm and/or presence of nutritional oedema⁽²⁹⁾. Children with complications affecting food intake or participating in other clinical trials were not eligible. Personal and socio-demographic information was obtained from caregivers using questionnaires. Age was obtained from birth growth monitoring cards, or by asking the caregiver if the card was not available. Current breast-feeding status was reported by interviewing caregivers using a structured questionnaire.

Nutritional markers were assessed by anthropometric measures (weight, length/height, MUAC, skinfolds) using standard equipment as described in Sigh *et al.*⁽²⁹⁾. Anthropometric Z-scores, namely WHZ or BAZ (wasting), weight-for-age Z-score (WAZ; underweight) and height-for-age Z-score (HAZ; stunting), were calculated from the WHO Growth Standards 2006 for children aged 6–59 months⁽³¹⁾ and the WHO Growth References 2007 for children aged 5–19 years⁽³²⁾. The mid-upper-arm fat percentage and muscle area were calculated according to Rolland-Cachera *et al.*⁽³³⁾ using MUAC and triceps skinfold measures, which were measured to nearest 0.1 mm using a Holtain skinfold calliper.

Blood sampling

Four millilitres of venous whole blood from the cubital vein were drawn using a sterile 23G/25G $\times \frac{3}{4}$ " scalp vein set (Vinahankook Medical Supplies Co., Hanoi, Vietnam) or a sterile 24G IV cannula needle with catheter and injection valve (Harsoria™, Gurgaon, India) by trained health staff. Two millilitres of whole blood were collected in 2 \times 2 ml EDTA-coated vacutainers and 2 ml of whole blood were collected in a 6 ml trace-element Sodium NH Heparin vacutainer (VACUETTE®; Greiner Bio-One International GmbH, Frickenhausen, Germany). One EDTA vacutainer was used to perform dried blood spots to measure PUFA and blood cell count analysis. Haemoglobinopathies were analysed using the second EDTA vacutainer and inflammation status was analysed using the sodium heparin vacutainer.

Dried blood spots and fatty acid analysis

Dried blood spots were performed using the laboratory facilities at the National Pediatric Hospital, following validated procedures⁽³⁴⁾. Dried blood spots were made

by applying 40 µl of EDTA whole blood on pre-treated chromatography paper (Grade 3MM, Whatman Ltd; Danish supplier: Frederiksen Scientific A/S, Ølgod, Denmark). The samples were dried at room temperature and stored in a polypropylene 'Ziplock' bag at 4°C for a maximum of 2 months prior to shipment to the University of Waterloo in Canada for fatty acid analysis^(34,35). The samples were analysed on a Varian 3900 gas chromatograph equipped with a DB-FFA 15 m × 0.10 mm capillary column coated with a 0.10 µm film of nitroterephthalic-acid-modified polyethylene glycol (J&W Scientific; Agilent Technologies, Mississauga, ON, Canada) using hydrogen as the carrier gas^(35–38). The eluted fatty acid methyl ester peaks were identified by comparing retention times with an external standard (GLC-246; Nu Chek Prep Inc., Elysian, MN, USA)⁽³⁹⁾. Fatty acid data are expressed as relative percentage of all fatty acids (% FA) and as fatty acid concentration (µg per 100 µg whole blood).

The inter- and intra-assay CV for the dried blood spot fatty acid analyses were similar to values previously reported for wet whole-blood analyses⁽⁴⁰⁾. For the fatty acids reported, all the inter-assay CV were below 7% and all the intra-assay CV were below 3%, with higher CV being associated with the fatty acids with relatively low abundance (i.e. docosapentaenoic acid (22:5n-6; DPA) and mead acid (20:3n-9; MA)).

Definition of essential fatty acid deficiency

Low LA and ALA are reflected in a ratio between their long-chain metabolites. The ratios MA:AA and n-6 DPA:DHA are commonly used biomarkers of essential fatty acid deficiencies^(41,42). For whole blood, deficiency levels for these markers have not been defined. However, it has been proposed that in whole blood, MA:AA > 0.02 indicates very low n-6 PUFA status and n-6 DPA:DHA > 0.2 and total n-6: n-3 PUFA > 10.5 indicate very low n-3 PUFA status⁽¹⁶⁾. We applied these cut-off values to define low PUFA.

Assessment of health markers

The markers assessed for health status were Hb concentration (for anaemia), blood cell counts (overall health status), Hb variations (for haemoglobinopathies), the presence of HIV (for HIV status), C-reactive protein (CRP; acute inflammation marker) and α₁-acid glycoprotein (AGP; progressed inflammation marker). Approximately 300 µl EDTA whole blood was used to assess blood cell counts, including Hb concentration, using a haematology analyser (Sysmex Kx-21, ID no. NP021 and NP023; Sysmex Corporation, Kobe, Japan) at the National Pediatric Hospital. Anaemia status was diagnosed based on Hb concentration using cut-off values recommended by WHO⁽⁴³⁾. The presence of haemoglobinopathies (only in the 6–59-month-old

children) was analysed using electrophoreses (Minicap; SEBIA, Lisses, France). Samples were analysed at the National Pediatric Hospital. Following a disruption of the capacity for analysing at the hospital, five samples were analysed at the Pasture Institute, Phnom Penh, using the same method and analysis protocol. Normal Hb was classified for 0–23-month-old children as HbA > 70%, HbF ≤ 30%, HbA2 < 2%; and for 24–59-month-old children as HbA > 95.5%, HbA2 = 2.0–3.5%, HbF < 5%. All other Hb variations were classified as haemoglobinopathies, except HbE/heterozygote, which was classified as HbE ~ 20–30%⁽⁴⁴⁾. Complete blood counts and haemoglobinopathies were analysed one sample at a time and no certified standard was available. Therefore, no inter- and intra-assay CV were available.

HIV status was self-reported via a questionnaire asking if the child had been tested and, if so, the result of the test. All HIV-positive children in the study received treatment at the National Pediatric Hospital. CRP and AGP were measured in a combined sandwich ELISA method in duplicate⁽⁴⁵⁾. The control samples' (n 40) inter-assay CV were 5.8% (CRP) and 8.1% (AGP). Inflammation was indicated by CRP > 5 mg/l and/or AGP > 1 g/l⁽⁴⁶⁾.

Statistical analysis

All data, except blood analyses, were entered directly into tablets and uploaded to an online software system (Kobotoolbox) or double entered into EpiData version 3.1 (The EpiData Association, Odense, Denmark). Statistical analyses were conducted using the software R studio (version 3.4.0 for Windows). Models were assessed by visual residuals and normal probability plots inspection. Non-normal distributed models were log₁₀ transformed. Descriptive statistics are presented as percentages and frequencies, means and standard deviations, or medians and interquartile ranges. All analyses, not pre-including age, are age-adjusted owing to large age variations in the study. To analyse correlations between PUFA and nutritional and health status markers, a simple or multiple linear regression was applied adjusted for age, sex and breast-feeding due to the well-known effects of breast-feeding on PUFA^(16,47–50). The analyses are reported as regression coefficients (β) with 95% confidence intervals and P values. Comparisons in cell counts between health markers were assessed using ANCOVA adjusted for age and sex with the Emmeans R package, which takes multiple comparisons into consideration. Statistical significance was defined as P < 0.05. P values were not adjusted for false discovery rates, as the correlation between PUFA and markers of nutrition status was consistent across variables, meaning the P values would be overcorrected if adjusted using the traditional Bonferroni method, resulting in a lower correlation factor and hence increased risk of type 2 errors.

**Table 1** Characteristics of the Cambodian children with acute malnutrition from the SAM trial conducted from September 2015 until January 2017 (mean and range; means and standard deviations; percentages and numbers; medians and interquartile ranges)

Characteristic	Children with acute malnutrition (<i>n</i> 174)	
	Mean, % or median	Min–max, SD, <i>n</i> or IQR
Age (years)†, mean and min–max	4.1	0.4–16.6
Sex, % and <i>n</i>		
Female	35	60
Currently breast-feeding, % and <i>n</i>	35	61
Nutritional status		
Wasting Z-score‡, mean and SD	–3.0	0.8
WAZ§, mean and SD	–3.3	0.9
Body weight (kg), mean and SD	11.2	7.8
HAZ, mean and SD	–2.4	1.3
Height (cm), mean and SD	88	26
MUAC (cm), mean and SD	12.8	24
SAM , % and <i>n</i>	58	101
MAM , % and <i>n</i>	36	63
Mid-upper-arm fat (%), mean and SD	31	9
Mid-upper-arm muscle area (cm ²), mean and SD	9.6	4.5
Health status		
Oedema, % and <i>n</i>	0.0	0
Hb (g/dl), mean and SD	11.1	1.5
Anaemia¶, % and <i>n</i>	52.3	91
Haemoglobinopathy††, % and <i>n</i>		
Hb normal	65.1	69
Hb disorders	34.9	37
HbE/heterozygote	20.8	22
Infection, % and <i>n</i>		
HIV positive	20	35
Inflammation markers		
Plasma CRP (mg/l), median and IQR	0.6	0.2–2.3
Plasma CRP (>5 mg/l), % and <i>n</i>	16	27
Plasma AGP (g/l), median and IQR	0.7	0.5–1.3
Plasma AGP (>1 g/l), % and <i>n</i>	35	61
CRP and AGP‡‡, % and <i>n</i>	12	20
Blood counts§§, mean and SD		
Erythrocytes (×10 ¹² /l)	4.6	0.7
Platelets (×10 ⁹ /l)	406	161
Leucocytes (×10 ⁹ /l)	11.7	4.5
Neutrophils (×10 ⁹ /l)	4.4	2.2
Lymphocytes (×10 ⁹ /l)	6.1	3.1
Monocytes (×10 ⁹ /l)	0.9	0.5
Eosinophils (×10 ⁹ /l)	0.3	0.4
Total cells (×10 ¹² /l)	4.8	0.7

IQR, interquartile range; WAZ, weight-for-age Z-score; HAZ, height-for-age Z-score; MUAC, mid-upper-arm circumference; SAM, severe acute malnutrition; MAM, moderate acute malnutrition; CRP, C-reactive protein; AGP, α_1 -acid glycoprotein.

†*n* 173, missing exact age and sex from one child.

‡Measured using weight-for-height Z-score (WHZ) or BMI-for-age Z-score (BAZ) as an indication of wasting.

§*n* 146, missing specific birthdate for twenty-eight children.

||Among children aged ≤ 59 months: SAM defined as either WHZ < –3 or MUAC < 115 mm; MAM defined as either WHZ between –2 and –3 or MUAC = 115–125 mm. Among children aged ≥ 60 months: SAM defined as either BAZ < –3 or MUAC < 129 mm (children aged 5–9 years), MUAC < 160 mm (children aged 10–14 years), MUAC < 190 mm (children aged 15–18 years); MAM defined as either BAZ between –2 and –3 or MUAC = 129–154 mm (children aged 5–9 years), MUAC = 160–185 mm (children aged 10–14 years), MUAC = 190–215 mm (children aged 15–18 years).

¶Anaemia cut-off is age- and sex-dependent.

††Haemoglobinopathies were only assessed in the 6–59-month-old children, *n* 106 out of the total 174; thus the percentage is calculated based on the *n* of the actual sampling.

‡‡Defined as CRP > 5 mg/l and AGP > 1 g/l.

§§*n* 163, data on blood cell counts were missing for ten children.

|||Erythrocytes, leucocytes and platelets combined.

Results

Participants

Out of 609 children screened, 174 (28.6%) were eligible for enrolment (Table 1). Their mean age was 4.1 years and their mean wasting Z-score was –3.0. The anaemia

prevalence was 52%, ~35% of the children had haemoglobinopathies and 20% (*n* 35) were HIV positive. Median levels of inflammation markers were 0.6 mg/l (CRP) and 0.7 g/l (AGP), reflecting an inflammation prevalence of 16 and 35% according to CRP and AGP, respectively.

Table 2 Whole-blood fatty acid composition of the malnourished Cambodian children from the SAM trial conducted from September 2015 until January 2017 (means and standard deviations)

Whole-blood fatty acid (% FA)	Children with acute malnutrition (n 174)		
	Mean	SD	
SFA			
Lauric acid	12:0	0.5	0.5
Myristic acid	14:0	1.7	0.9
Palmitic acid	16:0	27.4	2.1
Margaric acid	17:0	0.4	0.1
Steric acid	18:0	11.9	2.2
Arachidic acid	20:0	0.3	0.1
Behenic acid	22:0	0.8	0.2
Tricosylic acid	23:0	0.1	0.1
Lignoceric acid	24:0	1.4	0.4
Total SFA		45.5	3.4
MUFA			
Palmitoleic acid	16:1n-7	1.7	0.7
Vaccenic acid	18:1n-7	1.7	0.4
Oleic acid	18:1n-9	23.4	3.6
Gondoic acid	20:1n-9	0.3	0.1
Erucic acid	22:1n-9	2.4	1.0
Nervonic acid	24:1n-9	1.5	0.4
Total MUFA		31.1	3.7
Mead acid (MA)	20:3n-9	0.08	0.06
n-6 PUFA			
Linoleic acid (LA)	18:2n-6	13.2	2.7
γ -Linolenic acid	18:3n-6	0.1	0.1
Eicosadienoic acid	20:2n-6	0.2	0.0
Dihomo- γ -linolenic acid	20:3n-6	0.7	0.2
Arachidonic acid (AA)	20:4n-6	3.6	1.3
Docosadienoic acid	22:2n-6	0.1	0.0
Adrenic acid	22:4n-6	0.4	0.1
Docosapentaenoic acid (n-6 DPA)	22:5n-6	0.2	0.1
Total n-6 PUFA		18.6	3.5
n-3 PUFA			
α -Linolenic acid (ALA)	18:3n-3	0.2	0.1
EPA	20:5n-3	0.1	0.1
Docosapentaenoic acid (n-3 DPA)	22:5n-3	0.2	0.1
DHA	22:6n-3	0.8	0.3
Total n-3 PUFA		1.5	0.5
Total PUFA		20.1	3.7
Total fatty acid concentration†		349	131
Deficiency biomarkers			
MA:AA		0.02	0.02
n-6 DPA:DHA		0.4	0.2
n-6:n-3 PUFA		13.5	3.9

% FA, percentage of total fatty acids.

†Total fatty acid concentration is expressed as $\mu\text{g}/100\mu\text{l}$ whole blood.

Whole-blood PUFA and very low PUFA status

The relative contribution in whole blood was 13.2% FA for LA and was 0.2% FA for ALA (Table 2). The mean relative contribution for MA was 0.1% FA and this was reflected in a high MA:AA ratio, indicating very low n-6 PUFA status in the children according to the predefined cut-off (MA:AA > 0.02). Based on the predefined cut-offs (n-6 DPA:DHA > 0.2 and total n-6:n-3 PUFA > 10.5), 68% of the children appeared to have very low n-3 PUFA status. We estimated that 14% of the children may actually be both n-6 and n-3 deficient (MA:AA > 0.02, n-6 DPA:DHA > 0.2 and total n-6:n-3 PUFA > 10.5) based on these cut-offs.

Associations between whole-blood PUFA and markers of nutritional status

HAZ was significantly and positively correlated with total PUFA ($\beta = 0.57\%$ FA; 95% CI 0.14, 1.00% FA), LA ($\beta = 0.33\%$ FA; 95% CI 0.02, 0.64% FA), AA ($\beta = 0.16\%$ FA; 95% CI 0.01, 0.31% FA) and total n-6 LCPUFA ($\beta = 0.20\%$ FA; 95% CI 0.02, 0.38% FA; Table 3). The estimates indicated that an increase in HAZ of 1 was correlated with a ~0.6% FA-point increase in total PUFA. A similar association was found for absolute age-adjusted height. Height was also significantly positively associated with whole-blood DHA and n-3 LCPUFA (Table 4). A 6 cm increase in height was correlated with a ~0.7% FA-point increase in DHA (Table 4). A 50 cm increase in height was associated with a -0.01% FA-point decrease in MA:AA and hence with a decreased risk of n-6 essential fatty acid deficiencies.

MUAC was not significantly correlated with any of the PUFA measures (see online supplementary material, Supplemental Tables S1 and S2). However, separating MUAC into fat percentage and muscle area showed that a low mid-upper-arm fat percentage was associated with high AA, n-6 LCPUFA and n-3 LCPUFA (Tables 3 and 4), and a high mid-upper-arm muscle area was positively associated with AA and n-6 LCPUFA (Table 3).

Wasting Z-score and WAZ were not associated with any PUFA, but high body weight was associated with high EPA (20:5n-3; see online supplementary material, Supplemental Tables S1 and S2).

Associations between markers of health challenges and whole-blood PUFA

Hb concentration was not correlated with overall PUFA, n-6 PUFA, n-3 PUFA or either of the markers of low PUFA, MA:AA or n-6 DPA:DHA (Tables 3 and 4). Also, no associations were observed for anaemia (see online supplementary material, Supplemental Tables S1 and S2). However, the presence of haemoglobinopathies was associated with high MA:AA and a non-statistically significant tendency for low total PUFA ($\beta = -1.34\%$ FA-points; 95% CI -2.72, 0.04% FA-points; Table 3). An age-stratified analysis showed a 1.39% FA-point (95% CI -2.78, -0.00% FA-point, n 103) lower total PUFA in 6–59-month-old children with haemoglobinopathies than in the children without haemoglobinopathies.

HIV infection and high AGP concentration were associated with high MA:AA (Table 3). No associations were observed between CRP and any of the PUFA (Tables 3 and 4).

Unexpectedly, total whole-blood cell count was not associated with any of the PUFA measures. Furthermore, no correlations were observed for total count of leucocytes or erythrocytes. However, high platelet count was associated with high total PUFA and LA



Table 3 Associations of nutritional and health status markers with total PUFA and specific *n*-6 fatty acids in whole blood among malnourished Cambodian children from the SAM trial conducted from September 2015 until January 2017 (regression coefficients (β) and 95 % confidence intervals adjusted for age, sex and breast-feeding)

	<i>n</i> -6 PUFA									
	Total PUFA†		LA†		AA†		<i>n</i> -6 LCPUFA†		MA:AA†,‡	
	β	95 % CI	β	95 % CI	β	95 % CI	β	95 % CI	β	95 % CI
Nutritional status										
Wasting Z-score§, (n 168)	-0.06	-0.70, 0.83	-0.13	-0.43, 0.68	0.02	-0.29, 0.24	-0.03	-0.36, 0.29	-0.24	-4.82, 4.34
HAZ (n 168)	0.57	0.14, 1.00*	0.33	0.02, 0.64*	0.16	0.01, 0.31*	0.20	0.02, 0.38*	-1.85	-4.37, 0.66
Height (m) (n 168)	15.21	6.23, 4.19*	7.86	1.26, 14.46*	4.92	1.78, 8.05*	6.10	2.23, 9.97*	-16	-109, -4*
Mid-upper-arm fat (%) (n 168)	-0.05	-0.11, 0.03	-0.00	-0.28, 0.00	-0.03	-0.05, -0.00*	-0.03	-0.06, -0.00*	-0.05	-0.44, 0.35
Mid-upper-arm muscle area (cm ² , ×10) (n 168)	1.83	-0.83, 4.49	0.21	-1.73, 2.14	1.22	0.31, 2.13*	1.49	0.36, 2.61*	-7.43	-22.75, 7.89
Health status										
Hb (g/dl) (n 165)	0.24	-0.17, 0.64	0.14	-0.17, 0.43	0.07	-0.07, 0.21	0.09	-0.09, 0.26	-2.04	-4.36, 0.27
Haemoglobinopathy (yes v. no) (n 105)	-1.34	-2.72, 0.04	-1.01	-2.05, 0.03	-0.36	-0.90, 0.17	-0.33	-0.97, 0.31	8.53	0.12, 16.94*
Infection										
HIV infection, (yes v. no) (n 66)	-0.27	-3.52, 2.98	-1.07	-3.37, 1.23	0.35	-0.66, 1.35	0.75	-0.56, 2.06	24.82	8.11, 41.54*
Inflammation markers										
CRP (mg/l, ×10 ⁻⁴) (n 159)	-19	-88, 49	-17	-66, 32	1.12	-23.33, 25.58	0.13	-0.23, 1	133	-232, 499
AGP (g/l, ×10 ²) (n 159)	-598	-1299, 90	-432	-920, 57	-105	-351, 141	-99	-403, 206	6640	3102, 10177*
Blood counts										
Erythrocytes (×10 ¹⁵ /l) (n 165)	0.56	-2.30, 3.43	13.16	-48.37, 74.69	0.55	-29.04, 30.14	0.30	-36.23, 36.83	-4.49	-9.36, 3.77
Platelets (×10 ¹² /l) (n 165)	945	299, 1590*	0.38	0.12, 0.64*	0.10	-0.03, 0.22	0.12	-0.03, 0.28	-924	-2050, 202
Leucocytes (×10 ¹² /l) (n 165)	5.29	-1.23, 21.83	-3.20	-7.44, 13.84	0.89	-4.23, 6.00	1.09	-5.23, 7.41	-23.20	-51.99, 4.35
Monocytes (×10 ¹² /l) (n 165)	165	46, 284*	84	-3, 171	51	10, 93*	63	12, 114*	-179	-522, 164
Neutrophils (×10 ¹² /l) (n 165)	-64	-975, 847	-1.04	-20.40, 18.33	-1.06	-10.37, 8.25	-1.23	-12.72, 10.26	-1238	-2789, 313
Total cells¶ (×10 ¹² /l) (n 165)	68	-23, 359	0.15	-0.46, 0.75	0.01	-0.28, 0.30	0.01	-0.35, 0.37	-484	-978, 10

LA, linoleic acid; AA, arachidonic acid; LCPUFA, long-chain PUFA; MA, mead acid; HAZ, height-for-age Z-score; CRP, C-reactive protein; AGP, α_1 -acid glycoprotein.

* $P < 0.05$.

†For blood counts only, results are given as $\times 10^{-4}$ for total PUFA, $\times 10^3$ for LA and AA, $\times 10^2$ for *n*-6 LCPUFA and $\times 10^{-2}$ for MA:AA to present meaningful results.

‡For nutritional status and health status markers, results (except blood counts) are given as $\times 10^3$ to present meaningful results.

§Wasting Z-score is weight-for-height Z-score for children aged <5 years and BMI-for-age Z-score BAZ for children aged ≥ 5 years.

||Adjusted only for breast-feeding.

¶Erythrocytes, leucocytes and platelets combined.

Table 4 Associations of nutritional and health status markers with specific *n*-3 fatty acids in whole blood among malnourished Cambodian children from the SAM trial conducted from September 2015 until January 2017 (regression coefficients (β) and 95 % confidence intervals adjusted for age, sex and breast-feeding)

	<i>n</i> -3 PUFA							
	EPA†,‡		DHA‡		<i>n</i> -3 LCPUFA†,‡		<i>n</i> -6 DPA:DHA†,‡	
	β	95 % CI	β	95 % CI	β	95 % CI	β	95 % CI
Nutritional status								
Wasting Z-score§, (n 168)	5.7	-5.8, 17.2	7.4	-66.5, 1.3	-48.3	-152, 430	-8.4	-47.3, 30.6
HAZ (n 168)	5.2	-1.2, 11.7	25.3	-16.5, 7.1	30.5	-29.0, 89.1	-7.7	-29.8, 14.4
Height (m) (n 168)	175	-378, 313	1111	226, 1995*	1283	24, 2541*	-494	-962, -27*
Mid-upper-arm fat (%) (n 168)	-0.5	-1.5, 0.6	-4.8	-11.5, 1.9	-9.54	-18.9, -0.17*	3.2	-0.3, 6.7
Mid-upper-arm muscle area (cm ² , ×10) (n 168)	29	-11, 69	186	-72, 444	164	-202, 530	-95	-231, 41
Health status								
Hb (g/dl) (n 165)	-4.6	-10.7, 1.4	4.7	-34.7, 44.1	2.1	-3.5, 7.67	7.0	-13.8, 27.8
Haemoglobinopathy (yes v. no) (n 105)	1.4	-21.1, 23.8	-48	-201, 105	-0.7	-18.1, 16.7	6.4	-55.4, 68.2
Infection								
HIV infection (yes v. no) (n 66)	4.9	-37.6, 47.5	-5	-256, 245	-29.1	-496, 440	18.5	-158, 195
Inflammation markers								
CRP (mg/l, ×10 ⁻⁴) (n 159)	-157	-1, 897	-1367	-8136, 5401	-4676	-14 234, 4881	-39	-3653, 3574
AGP (g/l, ×10 ²) (n 159)	914	-11 556, 9727	-34 780	-102 858, 33 299	-78 210	-174 094, 17 682	8272	-28 150, 44 694
Blood counts								
Erythrocytes (×10 ¹⁵ /l) (n 165)	-7.2	-26.2, 11.8	2.6	-5.7, 10.8	4.3	-7.4, 16.0	-1.6	-7.2, 4.0
Platelets (×10 ¹² /l) (n 165)	3645	-712, 8003	0.02	-0.02, 0.05	0.02	-0.03, 0.07	-625	-1914, 664
Leucocytes (×10 ¹² /l) (n 165)	56.7	-53.1, 16.6	0.6	-0.9, 2.0	0.4	-1.7, 2.4	-12.5	-44.8, 19.8
Monocytes (×10 ¹² /l) (n 165)	818	-509, 2144	19	8, 31*	18	1, 34*	-273	-663, 116
Neutrophils (×10 ¹² /l) (n 165)	863	5183, 6910	0.5	-2.1, 3.1	1.2	-2.5, 4.8	-984	-2757, 789
Total cells¶ (×10 ¹² /l) (n 165)	-629	-2561, 1304	0.03	-0.05, 0.11	0.04	-0.07, 0.16	-183	-752, 385

LCPUFA, long-chain PUFA; DPA, docosapentanoic acid; CRP, C-reactive protein; AGP, α_1 -acid glycoprotein.

* $P < 0.05$.

†For blood counts only, results are total given as $\times 10^{-2}$ for EPA, $\times 10^2$ for *n*-3 LCPUFA and $\times 10^{-3}$ for *n*-6 DPA:DHA to present meaningful results.

‡For nutritional status and health status markers, results (except blood counts) are given as $\times 10^3$ to present meaningful results.

§Wasting Z-score is weight-for-height Z-score for children aged <5 years and BMI-for-age Z-score BAZ for children aged ≥ 5 years.

||Adjusted only for breast-feeding.

¶Erythrocytes, leucocytes and platelets combined.

Table 5 Comparison of blood counts in the presence or anaemia, HIV, inflammation or haemoglobinopathies among malnourished Cambodian children from the SAM trial conducted from September 2015 until January 2017 (means, standard deviations and *P* values adjusted for age and sex)

	Anaemia†					HIV‡				
	Anaemic (n91)		Non-anaemic (n77)		<i>P</i> value	HIV positive (n35)		HIV negative (n31)		<i>P</i> value
	Mean	SD	Mean	SD		Mean	SD	Mean	SD	
Erythrocytes (×10 ¹² /l)	4.6	0.1	4.7	0.1	0.229	3.7	0.2	4.9	0.2	<0.001
Platelets (×10 ⁹ /l)	434	17	371	18	0.010	349	34	343	4.3	0.915
Leucocytes (×10 ⁹ /l)	12.1	0.5	11.4	0.5	0.306	8.3	0.8	12.2	0.8	0.005
Neutrophils (×10 ⁹ /l)	0.6	0.0	0.6	0.0	0.950	2.8	0.5	4.9	0.5	0.006
Lymphocytes (×10 ⁹ /l)	6.4	0.3	5.8	0.3	0.237	4.7	0.5	6.0	0.5	0.125
Monocytes (×10 ⁹ /l)	0.9	0.1	0.8	0.1	0.176	0.7	0.1	0.8	0.1	0.250
Eosinophils (×10 ⁹ /l)	0.3	0.0	0.3	0.0	0.740	0.2	0.1	0.4	0.1	0.167
	Inflammation§					Haemoglobinopathies				
	Inflammation (n65)		No inflammation (n96)		<i>P</i> value	Hb disorder (n36)		No Hb disorder (n8)		<i>P</i> value
	Mean	SD	Mean	SD		Mean	SD	Mean	SD	
Erythrocytes (×10 ¹² /l)	4.5	0.1	4.7	0.1	0.069	4.8	0.1	4.8	0.1	0.795
Platelets (×10 ⁹ /l)	462	20	386	16	0.002	481	27	436	20	0.179
Leucocytes (×10 ⁹ /l)	12.6	0.5	11.5	0.4	0.073	1.2	0.7	12.9	0.5	0.140
Neutrophils (×10 ⁹ /l)	5.2	0.3	5.1	0.2	0.001	0.7	0.0	0.6	0.0	0.179
Lymphocytes (×10 ⁹ /l)	6.1	0.4	6.2	0.3	0.836	7.4	0.4	7.2	0.4	0.637
Monocytes (×10 ⁹ /l)	0.9	0.1	0.8	0.5	0.052	1.0	0.1	1.0	0.1	0.635
Eosinophils (×10 ⁹ /l)	0.3	0.1	0.3	0.1	0.583	0.5	0.1	0.3	0.1	0.021

†Analyses were not adjusted for age and sex, as this is included in the diagnostic criteria.

‡Only including children who had been tested for HIV.

§Inflammation was defined as C-reactive protein >5 mg/l or α₁-acid glycoprotein >1 g/l.

(Table 3), and high monocyte count was associated with high total PUFA, AA, *n*-6 LCPUFA, DHA and *n*-3 LCPUFA (Tables 3 and 4).

Comparing blood cell counts in children with and without health challenges

There was no significant difference observed in blood cell counts between children with SAM and MAM (see online supplementary material, Supplemental Table S3). Children with anaemia appeared to have a higher mean platelet count than those without anaemia (×10⁹/l: 434 *v.* 371, *P* = 0.010; Table 5). Furthermore, an association analysis adjusted for age and sex showed a non-significant tendency towards a negative correlation between Hb concentration and platelet count (×10⁹/l: -16.8; 95 % CI -33.8, 0.2), but not for monocyte count (×10⁹/l: -0.00; 95 % CI -0.05, 0.05). Presence of inflammation *v.* no inflammation resulted in a significantly higher mean platelet count (×10⁹/l: 462 *v.* 386, *P* = 0.002) and a strong tendency for a higher mean monocyte count (×10⁹/l: 0.9 *v.* 0.8, *P* = 0.052; Table 5). Children with inflammation defined as CRP > 5 mg/l (72 counts; 95 % CI 9, 135 counts) or AGP > 1 g/l (74 counts; 95 % CI 25, 123 counts) had higher mean platelets (×10⁹/l) compared with children with no signs of infection. Monocyte counts were elevated in children with CRP > 5 mg/l (estimated difference, ×10⁹/l: 0.20 counts; 95 % CI 0.01, 0.39 counts)

compared with children with normal CRP, but not in children with AGP above the cut-off of >1 g/l (estimated difference, ×10⁹/l: 0.09 counts; 95 % CI -0.06, 0.25 counts). Hence, the association was dependent on the stage of inflammation.

Children with HIV had lower counts of erythrocytes, leucocytes and neutrophils (Table 5). The presence of haemoglobinopathies was significantly associated with a higher mean eosinophil count (×10⁹/l: 0.5 *v.* 0.3, *P* = 0.021) and a tendency towards a lower mean platelet count (×10⁹/l: 481 *v.* 436, *P* = 0.179).

Discussion

We found low relative contributions of total whole-blood PUFA in children admitted to outpatient treatment of acute malnutrition. Stunting and low height were associated with low total PUFA, LA, AA and *n*-6 LCPUFA. Low mid-upper-arm muscle area and high mid-upper-arm fat percentage were also associated with low AA and *n*-6 LCPUFA, but no associations were observed for wasting or any other measures of weight except for EPA and total weight. No associations were observed between Hb, CRP or AGP, and total PUFA, sub-classes of PUFA or any of the individual PUFA. The presence of HIV infection, progressed inflammation (defined by AGP) and haemoglobin-



opathies were associated with high MA:AA, and hence a risk of very low *n*-6 PUFA status. The presence of haemoglobinopathies was also associated with low total PUFA, especially in the young children. High platelet and monocyte counts appeared to be linked with inflammation and anaemia. High platelet and monocyte counts were also associated with high total PUFA and LA and with *n*-6 and *n*-3 LCPUFA, respectively.

Whole-blood PUFA levels

The acutely malnourished children in the current study had considerably lower total PUFA, *n*-6 and *n*-3 PUFA, and AA and DHA compared with a population of rural Cambodian infants⁽⁴⁹⁾ living in an environment with limited dietary resources⁽⁵¹⁾ and vulnerable to stunting and wasting⁽⁵²⁾, respectively. The whole-blood ALA assessed in the present study children diagnosed with acute malnutrition was similar to what was reported in children with MAM from Burkina Faso (0.2% FA)⁽¹⁶⁾, while a study in Ugandan children with a more severe stage of SAM found even lower ALA (0.15% FA) as well as lower LA (10.5 *v.* 13.2% FA in the present study)⁽¹⁸⁾. We found relatively low MA (0.08% FA) compared with well-nourished Ugandan children (0.10% FA), which was unexpected, since the Ugandan children with SAM had higher whole-blood MA (0.12% FA)⁽¹⁸⁾. Our results are in line with the MAM study in Burkina Faso with regard to the essential fatty acid deficiency markers of MA:AA, *n*-6 DPA:DHA and *n*-6:*n*-3 PUFA⁽¹⁶⁾.

Low PUFA in acutely malnourished children could be due to infection and metabolic disturbances associated with malnutrition⁽¹⁵⁾. These disturbances could directly impair lipid digestion and uptake⁽¹⁵⁾ and also indirectly be affected by other nutritional imbalances such as Fe deficiency, which, in combination with acute malnutrition, could impact on PUFA metabolism through reduced Δ -6-desaturase enzyme activity⁽⁵³⁾ and consequently reduced capacity for the conversion of LA and ALA to their long-chain metabolites.

Whole-blood PUFA is highly influenced by diet, rather than by malnutrition *per se*⁽¹⁰⁾. Cambodian children generally consume a diet based on rice, with limited amounts of vegetables, legumes and fruits, and little or no animal-source foods^(3,4). This diet is at risk of being low in PUFA. Also, the cooking oils typically used are low in *n*-3 fatty acids⁽⁴⁾. Fish and fish products are the main animal-source food in the typical Cambodian diet⁽⁵⁴⁾ and, overall, fish is a highly important and valuable PUFA source in diets. The fish in the Cambodian diet are primarily freshwater fish available from the rich aquatic environment of the Mekong river basin. Tropical freshwater fish are generally lower in DHA and EPA compared with cold-water marine fish species⁽⁵⁵⁻⁵⁷⁾. Furthermore, despite fish being consumed very frequently in Cambodia (even daily in the peak fishing season) the

portion size can be very small, especially for children, resulting in low overall fish intake⁽⁵¹⁾ and a diet low in *n*-3 PUFA. The low PUFA status of the children in our study could therefore also be the result of a low dietary intake, further enforced by the suffering from acute malnutrition.

Nutritional markers associated with whole-blood PUFA

The associations between stunting, height and low *n*-6 PUFA support the idea that sufficient dietary PUFA intake may be important in preventing stunting. Height gain was found to be correlated with higher DHA in young children in treatment for SAM in Uganda⁽¹⁷⁾. A study of 2-6-year-old children from Tanzania found associations between risk of stunting and low whole-blood LA and total *n*-6 PUFA⁽⁵⁸⁾, while mixed findings between stunting and PUFA were reported in Ghanaian children^(59,60). The indication of a biological link between linear growth and PUFA status could be mediated by eicosanoids such as AA-derived prostaglandin E₂, which can affect proteolysis in skeletal muscle and thereby have an impact on linear growth⁽⁶¹⁾. Another pathway could go through prostaglandin E₂ stimulation of insulin-like growth factor-1 expression, which again mediates bone Ca accretion^(10,58,62). A few studies have found increased insulin-like growth factor-1 after fish-oil supplementation in children^(63,64).

We observed an association between high PUFA and muscle mass. Low mid-upper-arm fat percentage and high mid-upper-arm muscle area were associated with high AA and *n*-6 LCPUFA, meaning that high fat percentage or a reduced muscle mass was associated with less PUFA in the blood. Acute malnutrition causes muscle atrophy, which could be linked to increased stress metabolism and hypercortisolaemia⁽⁶⁵⁾. Although beyond the conclusions of the present study, we speculate that hypercortisolaemia may promote muscle proteolysis and increase energy expenditure and fat oxidation⁽⁶⁶⁾, which could result in a combined effect of lower PUFA and muscle mass. Cortisol and muscle proteolysis as well as low PUFA could also be caused by the combination of low dietary intake and malnutrition.

Compromised health is associated with very low PUFA status

Hb concentration and anaemia were not associated with any PUFA or the cut-off used to define very low PUFA status. This is inconsistent with studies of children with either MAM or SAM that show a correlation between Hb concentration and AA and DHA and low PUFA status (*n*-6 DPA: DHA and MA:AA)^(16,18).

Other studies observed an association between acute inflammation (CRP > 5 mg/l) and PUFA (LA, EPA and *n*-3 DPA) in children with SAM⁽¹⁸⁾ and between progressed inflammation (AGP \geq 1 g/l) and PUFA (AA, *n*-6 DPA, ALA



and DHA) in children with MAM⁽¹⁶⁾. The present study's finding that the inflammation markers were not associated with PUFA could probably be because the prevalence of inflammation was low compared with the two other studies^(16,18). Studies in various populations have shown reduced CRP following fish-oil supplementation^(67–69), which could support the suggestion that high PUFA may have a protective effect against inflammation.

Haemoglobinopathies were associated with low total PUFA in an age-stratified analysis in the young children. We speculate that haemoglobinopathies could cause a higher erythrocyte turnover, and thereby a higher phospholipid turnover. In the presence of haemoglobinopathies, a higher repair of blood cells, required due to oxidant stress and lipid damage of the erythrocyte membrane, is also speculated. Furthermore, because of lipid oxidation, the phospholipid repair system is insufficient in maintaining a proper molecular fatty acid composition in erythrocytes.

All blood cells are exposed to the same pool of fatty acid substrate from plasma, and the difference in PUFA measured in the whole-blood samples could be caused by the activity of enzymes using the fatty acid pool for phospholipid repair. Oxidant stress in erythrocytes may challenge the incorporation of PUFA, which affects the fatty acid composition of the membrane⁽⁷⁰⁾. This may lead to low PUFA in cells and thereby to lower PUFA in children with haemoglobinopathies than in children without haemoglobinopathies.

Health markers' influence on blood counts may indirectly implicate incorrect assessment of whole-blood PUFA status

Unexpectedly, the inflammation markers were not associated with any PUFA but were found to be associated with counts of platelets and monocytes, also formed as a response to infection. A sub-analysis showed an association between the markers of inflammation and platelets and monocytes, which confirms the expected biological response of CRP and AGP at the onset of infection and inflammation. The pattern of associations between the different markers used to define inflammation indicated differences in the biological response to acute and progressed inflammation.

Different blood fractions and cell types contribute different PUFA levels, so changes in the number of a specific cell could shift levels of PUFA in whole blood. This is likely to explain the increased PUFA we observed with increased monocyte and platelet counts, as monocytes consist of ~20% AA and AA increases with increased monocyte count⁽⁷¹⁾. Also, a decrease in erythrocytes, which is likely to be associated with anaemia, could result in reduced measurement of PUFA in whole-blood samples. These contra-acting mechanisms could explain the lack of associations of health markers for anaemia, inflammation and haemoglobinopathies with PUFA in the present study.

The combined effects of anaemia with elevated platelets and monocytes caused by inflammation may have masked changes in PUFA.

The fatty acid composition of immune cells (e.g. lymphocytes and monocytes) is also expected to alter in accordance with dietary intake, which impacts the capacity of the immune cells to produce eicosanoids, which are involved in immune system regulation⁽⁷²⁾. Thus, there is a two-way association between the immune system and PUFA. Inflammation is suggested to cause elevated cell counts in the present study, which may lead to misinterpretation of PUFA in whole blood. Inflammation markers were observed to influence platelets and monocytes differently, and platelets and monocytes were not associated with the same fatty acids. The complexity of these associations would make it difficult to adjust for inflammation markers and cell counts when assessing PUFA in whole blood in populations with severe health challenges such as acute malnutrition.

Limitations

The present study shares the limitation of other observational studies in that it identifies associations but does not identify causality. Another limitation is that we did not include data on total food consumption covering all dietary sources of PUFA such as quantity of fish and breast milk consumed. Also, the predefined cut-offs used to estimate essential fatty acid deficiencies were derived from infants and were applied to older children in the present study. The stage of deficiencies in children above infancy could be different. Finally, all the health markers used in the present study are to some degree related through cell counts, and it could be argued that all health markers should be adjusted for each other (anaemia, inflammation, HIV infection and haemoglobinopathies). Our study used secondary data and unfortunately it did not have insufficient power to adjust all health challenges for each of the others.

Conclusion

Overall, children with acute malnutrition in Cambodia had low total whole-blood PUFA and indicated to have a high prevalence of a very low PUFA. HAZ (stunting) and absolute height were both positively associated with PUFA, whereas no associations were found for wasting and other weight measures. Elevated platelets and monocytes, likely due to inflammation, were positively associated with PUFA, thus indicating that blood cell counts influence whole-blood PUFA levels. Therefore, in acutely malnourished Cambodian children, inflammation that elevates blood cell counts could result in increased blood levels of PUFA that do not reflect dietary intake and nutritional status. These findings indicate the need for further studies on the relationship between stunting and PUFA in acutely

malnourished children through randomized controlled trials with dietary PUFA supplementation. Furthermore, future field studies using whole blood and dried blood spots to assess PUFA status should consider co-morbidities associated with severe health challenges, such as malnourishment, that can change the cellular composition of blood.

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Supplementary material

To view supplementary material for this article, please visit <https://doi.org/10.1017/S1368980019003744>

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