

Analytical Method Assessment for the Determination of DHA and Fatty Acids Present in Unextracted Aurantiochytrium limacinum **Biomass**

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Abstract

Research into long-chain polyunsaturated fatty acids (LC-PUFA), such as docosahexaenoic acid (DHA C22:6 n-3), has shown that their inclusion in the human diet is linked with many health benefits. This has led to an increased interest in the enrichment of certain foodstuffs with DHA by supplementing animal fed with DHA-rich ingredients which can lead to an increased uptake in the meat, milk and eggs animal by-products. The microalgae Aurantiochytrium limacinum has been found to be especially useful in this pursuit. It is subsequently desirable to availably have a simple and robust method for the routine analysis of DHA and other fatty acids in the algal biomass. The AOAC method 996.06 is often followed for the analysis of fatty acids in foods and demonstrating that its fitness for purpose in the analysis of DHA and additional fatty acids in Aurantiochytrium limacinum is therefore the objective of this paper. A validation of the method for the determination of DHA and three other fatty acids in Aurantiochytrium limacinum is presented. The method was found to be linear over the following ranges for each fatty acid methyl ester (FAME) analyte; 50 to 15,000 $\mu g/ml$ (C14:0), 300 to 95,000 (C16:0), 25 to 15,000 (C18:0) and 300 to 59,375 (C22:6). The accuracy, precision and LOD and LOQ of the method were confirmed and its robustness tested. The application of the method to assess the stability of Aurantiochytrium limacinum containing two alternative antioxidants was further examined. The investigation showed that DHA was stable over six months with the inclusion

of either Duralox[®] or ethoxyquin as an antioxidant and ethoxyquin could additionally stabilize DHA in *Aurantiochytrium limacinum* up to 24 months.

Keywords

DHA, GC-FID, Fatty Acids, Analytical Method, Fitness for Purpose, *Aurantiochytrium limacinum*

1. Introduction

In recent times, research into long-chain polyunsaturated fatty acids (LC-PUFA) has shown that they are associated with a range of health benefits including; improved brain development and cognitive function [1] as well as a reduction in the risk of depression [2], inflammatory processes [3] and cardiovascular disease [4]. LC-PUFA can be further classified as omega-6 (n-6) and omega-3 (n-3) fatty acids, where appropriate levels of both are required for optimum health [5]. The omega-3 fatty acid DHA is considered to be particularly important in the human diet as it is not easily derived in the body and must be supplemented through food sources [6]. Despite the awareness surrounding DHA, intake remains low in populations consuming a westernized diet [7]. Some estimates suggest that 80% of adults consume less than the recommended amount of DHA [8].

In order to address the deficiency of DHA in the human diet, enrichment of commonly consumed foods has gained popularity among researchers and food producers [9]. The DHA content of foods, such as animal by product, can be fortified by supplementing animal fed with ingredients rich in DHA [9] [10]. DHA enrichment of common food sources can therefore lead to a greater availability and accessibility of food choices rich in DHA and furthermore an increased consumption of DHA without any significant change to consumer diets [6].

Dietary supplementation of intensively farmed animals with fish oils, rich in both the omega-3 fatty acids DHA and eicosapentaenoic acid (EPA), has been used to increase the omega-3 content of animal meat, milk and egg products; [11] [12]. However, due to a global decline in fish stocks, as well as issues related to the stability to autoxidation of processed fish oils, a move to more sustainable alternatives of omega-3 rich feed supplements is desirable. Heterotrophically grown microalgae represent an alternative source containing high concentrations of protected DHA and can be used to supplement animal feed [13] [14]. The algal subject of this paper, *Aurantiochytrium limacinum* is a microalgae that has been shown to increase the DHA content of milk [15] [16], of pig muscle tissue [17] [18], broiler [19] and of laying hen eggs [20] when supplemented to the animal diets.

It is subsequently necessary to be able to quantify the fatty acid concentrations in DHA-rich feed ingredients. This can be achieved by using various methods such as Thin Layer Chromatography (TLC) interfaced with detection techniques such as Flame Ionization Detection (FID) and Matrix Assisted Laser Desorption Ionization Mass Spectrometry (MALDI-MS) [21] Likewise, Gas Chromatography (GC) and Liquid Chromatography (LC) methods hyphenated with various Mass Spectrometry (MS) detection systems which can yield a comprehensive fatty acid speciation of test subjects [22]. However, for routine and robust fatty acid analysis, GC-FID represents the method of choice due to its specific, sensitive and rapid analysis [23].

The fatty acid content in food & feed is commonly determined by extraction from their relevant sources using the method developed by Folch *et al.* [24] and by synthesising respective fatty acid methyl esters (FAME), which are then analysed by GC [25] [26]. The AOAC method 996.06, for example, is routinely followed for this purpose [27]. GC-FID has been reported in several research publications on the analysis of fatty acids in food and feed ingredients including the analysis of microalgae [28] [29]. Whilst these papers provide useful information on fatty acid content in a range of settings, there has not been up to now, a published demonstration of the fitness for purpose of the methodology for analysing DHA in an unextracted microalgae biomass in dry powder form. Moreover, this is the first disclosure of the DHA analysis of the specific strain of *Aurantiochytrium limacinum*. As algae has emerged as an ingredient of enormous potential for use in the enrichment of food, there is a need for a simple and robust method for the routine analysis of algae to monitor its FA composition and evaluate if they meet the desired specification.

The objective of this paper therefore is to establish the fitness for purpose vis-a-vis a validation of the method, AOAC 996.06, for the determination of DHA and other fatty acids present in the unextracted biomass of a heterotrophically grown and spray-dried Aurantiochytrium limacinum; specifically, tetradecanoic acid (C_{14:0}), hexadecanoic acid (C_{16:0}), octadecanoic acid (C_{18:0}). The parameters investigated as part of the validation study included; linearity and range, accuracy and precision, selectivity and the limit of detection (LOD) and limit of quantification (LOQ). A further objective of the paper was to apply the method to the stability assessment of DHA in respective batches of Aurantiochytrium limacinum containing two alternative antioxidants, ethoxyquin and Duralox[®]. Ethoxyquin has been widely used as an antioxidant in foods of animal origin and animal feed [30]. However, in 2017, the European Commission suspended its use as a technological additive in all species citing safety concerns of impurities and metabolites associated with the additive [31]. With a view to assessing the stability using a more natural alternative, the authors conducted a 6-month stability trial on batches of Aurantiochytrium limacinum containing Duralox[®] and compared the results to batches containing ethoxyquin. Duralox[®] is a commercial antioxidant containing rosemary extract and tocopherols.

2. Materials and Methods

2.1. Instrumentation and Chemicals & Reagents

The verification experiments were performed at Mérieux NutriSciences (Burnaby, Canada) in compliance with ISO 17025 requirements and ICH Q2B Guidelines. All experiments were performed on an Agilent 6890N gas chromatograph (Agilent, Ontario, Canada) equipped with a hydrogen FID and an Agilent 7683 autosampler (Agilent, Ontario, Canada). Solvents and reagents used during the extraction and methylation steps were sourced from Fisher Scientific (Ontario, Canada). These included diethyl ether, petroleum ether, water (HPLC grade), ethanol 95%, chloroform, toluene, hexane and methanol. Boron trifluoride, 7% was prepared by diluting boron fluoride 14% one to one with methanol. Hydrochloric acid, 8.3N was prepared by adding 250 ml of concentrated HCl (12 N) to 110 ml of deionized water. Pyrogallic acid was obtained from TCI America (Portland, USA).

The FAME reference solution (GLC-85) and the standards solutions comprising: methyl 4,7,10,13,16,19-docosahexaenoate; methyl octadecenoate, methyl hexadecanoate; and methyl tetradecenoate; internal standard for sample extraction, 1,2,3-triundecanoylglycerol (common name: triundecanoin, $C_{11:0}$); internal standards for calibration curve, methyl undecanoate and docosahexaenoic acid (DHA) were purchased from Nu-Chek-Prep Inc. (Minnesota, USA). The test microalgae sample (Unextracted *Aurantiochytrium limacinum* CCAP 4087/2) was provided by Alltech Inc. (Nicholasville, KY, USA).

2.2. Preparation of Stock Solutions, Internal Standards and Calibration Standards

The FAME standard stock solutions were prepared by dissolving 1.00 ± 0.05 g of commercial standards in hexane in a volumetric flask. A mixed FAME spiking solution with a final concentration of 10 mg/ml for C_{14:0} and C_{18:0} and 100 mg/ml for C_{16:0} and C_{22:6} were prepared from the FAME standard stock solution in a final volume of 10 ml of hexane. A mixed FAME spiking solution for the defatted matrix analysis with a final concentration of 2.5 mg/ml for C_{14:0}, 25 mg/ml for C_{16:0}, 1.25 mg/ml for C_{18:0} and 62.5 mg/ml for C_{22:6} was prepared by taking an aliquot of each FAME standard stock solution in a final volume of 10 ml of hexane. A methyl undecanoate internal standard solution was prepared in hexane with 1 g of methyl undecanoate, in a final volume of 10 ml using a volumetric flask. Linearity experiments were conducted over the following concentrations ranges of each analyte by adding given volumes of FAME standard solution to 100 µl of methyl undecanoate internal standard solution and making up to 2000 μ l with hexane. When not in use, solutions were stored at <-16°C. Linearity was established using 6-point calibration curves ranging from: 50 to 15,000 µg/ml (C $_{\rm 14:0}),\;300$ to 95,000 (C $_{\rm 16:0}),\;25$ to 15,000 (C $_{\rm 18:0})$ and 300 to 59,375 (C $_{\rm 22:6})$ and amended with C_{11:0} internal standard.

2.3. Preparation of Aurantiochytrium Biomass

In order to further explore the suitability of the method, a second test sample with an alternative matrix composition—defatted algae powder—was prepared and subjected to accuracy and precision experiments similar to the primary non-defatted algal test sample. To make the defatted matrix, algae powder was blended with water and then dried in a vacuum oven at 70°C. The sample was rinsed with acetone to remove excess water and then extracted using a soxhlet apparatus with petroleum ether. The soxhlet program was set to extract 45 minutes and rinse 45 minutes, followed by drying for 1 hour.

Extraction of Fat

Control algae powder was thoroughly mixed to ensure a homogeneous sample was obtained. A 0.15 \pm 0.01 g sample of the powder was weighed into a Mojonnier flask. For spiked algae samples, respective volumes of 100, 200 and 300 µl of mixed FAME stock solution were separately added into control algae samples to achieve the spike concentrations as outlined in Table 4 and Table 5. Pyrogallic acid (100 mg) was added to the flask which was then fortified with 75 µl of 40 mg/ml mixed FAME standard working solution. A 2 ml aliquot of triundecanoin internal standard solution, a few boiling granules and 2 ml of ethanol was added. The sample was mixed until completely wet. A 10 ml aliquot of 8.3 M hydrochloric acid was added. The flask was placed in a 70°C - 80°C water bath for one hour and the contents were mixed every ten minutes before being cooled to 20°C and vortex-mixed for 15 seconds. Excess ethanol was added to rinse down all particles adhering to the walls of the flask. A 25 ml portion of diethyl ether was added to the flask, which was stoppered and shaken gently. A 25 ml portion of petroleum ether was added, and the flask was again shaken and vortex-mixed for two minutes. The flask was centrifuged for five minutes at 600 rpm to yield a clear supernatant, which was decanted into a round bottomed flask and evaporated using a rotary evaporator and nitrogen to remove any residual solvent.

Methylation

The extracted fat residue was dissolved in 3 ml of chloroform. A 3 ml volume of diethyl ether was added, and the resulting solution was placed into a 10 ml glass tube. The round bottomed flask was washed with diethyl ether and the washings were added to the glass tube. The solution in the glass tube was evaporated to dryness under nitrogen at 40°C. A 2 ml volume of 7% BF3 reagent and 1ml of toluene was added to the glass tube which was vortex-mixed for 30 seconds. The tube was sealed tightly with a screw-cap top with a Teflon/silicone septum and heated in an oven for one hour at 100°C with gentle shaking every ten minutes. The tube was allowed to cool to 20°C and 5 ml of HPLC-grade water, 1 ml hexane and about 1 g sodium sulfate were added. The tube was capped and vortex-mixed for one minute and then centrifuged at 2000 rpm for 3 minutes. The resulting clear supernatant was dried again with sodium sulfate and injected into the GC-FID.

GC-FID Conditions

The GC system utilized a SP2560 100 cm long capillary column, 0.25 mm internal diameter; 0.20 μ m film thickness (Supelco, Pennsylvania, USA). The system used helium as a carrier gas with an initial flow of 1.1 ml/min and an average velocity of 19 cm/sec at a pressure of 35.74 psi. Air and hydrogen were used for the FID with pressures of 60 and 40 psi respectively. The initial oven temperature was 100°C which was held for four minutes and then ramped up at a rate of 3°C/min to 240°C and held for 19 minutes. The injector was employed in the split mode at a ratio of 200:1 and at a temperature of 225°C. The detector temperature was set to 285°C. System suitability was evaluated using the calibration GLC-85 reference FAME standards injected in five replicates.

2.4. Method Verification

Specificity

Blank algae powder samples were analyzed in six replicates and in three replicates on two successive days by the primary analyst and in three replicates on the second day by a secondary analyst in order to determine the endogenous levels of the analytes $C_{14:0}$, $C_{16:0}$, $C_{18:0}$ and $C_{22:6}$. In addition, an algae powder sample was spiked with a low concentration of analytes to determine if the method could distinguish between the analytes of interest and other interfering substances of the matrix.

Linearity

Standard curves of the respective analytes and a suitable range of concentrations were prepared to demonstrate linearity of the method. Linear regression, forced through the origin and with equal weighting, was applied to the peak area ratios plot for the construction of calibration curves and to provide information on the slope, coefficient of determination, and intercept.

Accuracy

In order to determine the accuracy of the method, blank algae powder was spiked with $C_{14:0}$, $C_{16:0}$, $C_{18:0}$ and $C_{22:6}$ in FAME form at three concentration levels as followed and % recoveries were calculated:

- C_{14:0} was spiked to final concentrations of approximately 3.2%, 3.9% and 4.6% (w/v),
- C_{16:0} was spiked to final concentrations of approximately 41.3%, 48.0% and 54.6% (w/v),
- C_{18:0} was spiked to final concentrations of approximately 1.7%, 2.4% and 3.0% (w/v), and
- C22:6 n3 was spiked to final concentrations of approximately 25.7%, 32.4% and 39.1% (w/v).

The accuracy study was repeated using a defatted blank matrix which contained the following concentrations of the fatty acid analytes; $C_{14:0}$ 1.4%, $C_{16:0}$ 15.4%, $C_{18:0}$ 0.48% and $C_{22:6}$ 7.7%. The blank algae powder was spiked with $C_{14:0}$, $C_{16:0}$, $C_{18:0}$ and $C_{22:6}$ in FAME form at three levels at the following concentrations:

- C_{14:0} was spiked to final concentrations of approximately 1.7%, 2.0% and 3.3% (w/v),
- C₁₆:0 was spiked to final concentrations of approximately 18.5%, 21.5% and 33.6% (w/v),
- C18:0 was spiked to final concentrations of approximately 0.6%, 0.8% and 1.4% (w/v), and
- C22:6 was spiked to final concentrations of approximately 10.1%, 12.3% and

21.6% (w/v).

The blank algae powder and defatted algae powder were extracted and analyzed in six replicates on Day 1 (first analyst only), and in three replicates on Day 2 by both the primary and secondary analysts. The average of the endogenous levels found in the blank was added to the amount spiked prior to the spike recoveries being calculated.

Precision

The spike data described from the accuracy experiments were analyzed for precision in terms of intra-day and inter-day variability. Intra-day precision was calculated for each analyte at each concentration. Inter-day precision was calculated to include data from all three analysis days, for each analyte.

Limit of Detection (LOD) and Limit of Quantification (LOQ)

To determine the method LOD and LOQ for $C_{14:0}$, $C_{16:0}$, $C_{18:0}$ and $C_{22:6}$, ten replicate determinations of algae powder were made and the mean and standard deviations (SD) for each analyte were calculated. The LOD was calculated as three times the SD and the LOQ was calculated as ten times the SD.

Stability

In order to assess the stability of DHA in *Aurantiochytrium limacinum*, alternative batches were prepared containing the antioxidants, ethoxyquin and Duralox[°] (Videka, Elven, France). The antioxidants were added post-heterotrophic fermentation under mixing in a chilled holding tank prior to spray drying. Ethoxyquin was added to three independent batches of *Aurantiochytrium limacinum* at a concentration of 200 mg/kg DM and the stability was determined over 24 months. Duralox was added to four batches of *Aurantiochytrium limacinum* at a concentration of 1000 mg/kg DM and the stability was assessed over six months.

3. Results

Specificity

The analysis of the algae powder confirmed the presence of endogenous levels of each investigated analyte and the results are included in **Table 1**. A chromatogram of the low concentration spike sample was compared to a chromatogram of the blank algae. The specificity of the method was demonstrated by the ability to distinguish between each FAME analyte from other compounds present in the matrix.

Linearity

The method was linear over the calibration ranges for all FAME analytes $C_{14:0}$, $C_{16:0}$ and $C_{18:0}$ and $C_{22:6}$. The coefficients of determination, R^2 , were consistently at 1.000 for each analyte's respective calibration curves. They were within the acceptable criteria of $R^2 \ge 0.990$ ($R \ge 0.995$) (Table 2).

Accuracy

To assess the accuracy, the mean of nine percent recovery determinations by the primary analyst and three by the secondary analyte were obtained. The spike recoveries for the four monitored analytes in the algae powder were calculated

Replicate No.	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}
1	2.59	34.7	1.06	19.1
2	2.62	35.1	1.06	19.3
3	2.63	35.1	1.06	19.2
4	2.62	35.1	1.06	19.3
5	2.68	35.9	1.08	19.8
6	2.64	35.5	1.07	19.6
7	2.54	34.4	1.07	18.9
8	2.61	35.3	1.07	19.3
9	2.60	35.0	1.07	19.3
Mean	2.61	35.1	1.07	19.3
SD	0.038	0.438	0.010	0.270
RSD (%)	1.47	1.25	0.900	1.40

 Table 1. Fatty acid % content in an unextracted Aurantiochytrium limacinum algae dried biomass.

Weight of FA in FAME form (g) = [(Response FAME*Weight of IS*Response Factor IS)]/Response IS*Response Factor FAME; Weight of FA in Fatty Acid Form (g) = Weight of FA in FAME form (g)*Conversion factor for FAME form to FA form; %FA = Weight of FA in Fatty Acid Form (g)/weight of sample (g)*100 [27]; RSD (%) = (SD/Mean)*100.

Table 2. Correlation coefficients and linearity equations for fatty acid analysis of unextracted biomass of *Aurantiochytrium limacinum* powder.

ID	Parameter	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}
Analysis #1	\mathbb{R}^2	1.000	1.000	1.000	1.000
	linear equation	y = 1.103x	y = 1.150x	y = 1.161x	y = 1.102x
Analysis #2	\mathbb{R}^2	1.000	1.000	1.000	1.000
	linear equation	y = 1.211x	y = 1.193x	y = 1.122x	y = 1.307x

and ranged from 91.6% to 107.3%, which was within the acceptable criteria of 90% - 110%. The SD of the recoveries was further calculated to determine the % relative standard deviation (%RSD). See **Table 3** for the complete summary of results.

The spike recoveries for the four monitored analytes in the defatted algae were also calculated and ranged from 80% to 119%. This was outside the desired criteria of 90% - 110%. See **Table 4** for the summary of the results. We have to note that the de-fatted algae powder was prepared only as a complementary evaluation to support the accuracy and precision of the method on the main matrix *i.e. Aurantiochytrium limacinum* algae powder. The impact of the defatting process is likely to have altered the overall stability of the matrix, with a consequent increase in the variability of the result compared to those obtained in the primary matrix.

Primary Analyst	<u>C_{14:0}</u>	<u>C_{16:0}</u>	<u>C_{18:0}</u>	<u>C_{22:6}</u>	<u>C_{14:0}</u>	<u>C_{16:0}</u>	<u>C_{18:0}</u>	<u>C_{22:6}</u>	<u>C_{14:0}</u>	<u>C_{16:0}</u>	<u>C_{18:0}</u>	<u>C_{22:6}</u>
Spiked Conc (%)	3.2	41.3	1.7	25.7	3.9	48.0	2.4	32.4	4.6	54.6	3.0	39.1
Replicate-1	100.8	99.9	97.5	100.9	103.7	102.8	100.1	106.0	103.6	101.9	100.1	107.2
Replicate-2	101.9	101.4	98.7	102.6	104.4	103.7	100.8	107.3	100.8	99.3	97.4	104.1
Replicate-3	102.0	101.3	99.1	102.7	100.7	99.9	97.7	103.6	101.9	100.4	98.2	104.7
Replicate-4	105.0	104.5	101.7	97.7	102.1	101.2	98.8	103.7	100.6	99.1	96.8	103.9
Replicate-5	102.2	101.8	99.4	103.2	101.2	100.4	98.9	103.8	101.6	100.2	97.8	103.9
Replicate-6	103.6	103.0	100.6	104.1	104.0	103.1	100.5	106.2	100.5	99.1	96.8	103.1
Replicate-7	101.1	101.4	99.2	103.3	102.5	102.0	99.8	103.8	101.3	101.3	99.0	104.3
Replicate-8	99.5	100.0	98.7	101.1	96.2	97.2	91.6	96.8	101.0	100.3	97.5	102.4
Replicate-9	100.4	100.7	98.5	101.7	101.1	100.4	99.9	104.9	103.1	103.5	101.4	106.8
Mean, primary analyst	101.8	101.6	99.3	101.9	101.8	101.2	98.7	104.0	101.6	100.6	98.3	104.5
Std Deviation	1.7	1.4	1.2	1.9	2.5	2.0	2.8	3.0	1.1	1.5	1.6	1.6
RSD (%)	1.6	1.4	1.2	1.9	2.4	2.0	2.9	2.9	1.1	1.4	1.6	1.5
Secondary Analyst	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}
Replicate-1	99.3	97.7	95.4	97.4	101.4	99.7	98.3	101.7	101.7	99.8	97.9	101.6
Replicate-2	100.5	99.3	97.3	99.2	101.5	100.1	98.1	101.5	101.6	100.0	97.6	102.2
Replicate-3	98.0	96.7	96.0	97.4	100.2	98.9	96.7	100.1	100.7	99.1	97.2	101.1
Mean, secondary analyst	99.2	97.9	96.2	98.0	101.0	99.6	97.7	101.1	101.3	99.6	97.6	101.6
RE (%) to Primary Analyst	-2.5	-3.6	-3.0	-3.9	-0.7	-1.6	-1.0	-2.8	-0.3	-1.0	-0.8	-2.7

Table 3. Accuracy of fatty acids recoveries (%) from the unextracted biomass of *Aurantiochytrium limacinum* algae powder spiked to three concentration levels.

RSD (% = (SD/Mean) * 100; RE (Relative error (%)) = ((mean, 2nd analyst - mean, 1st analyst)*(mean, 1st analyst)) * 100.

Precision

The precision of the method was established by calculating the coefficient of variation (CV, %) from the replicate determinations of the primary and secondary analysts. The precision ranged from 1.9% to 2.7% for the spiked algae powder and 9.8% to 11.5% for the defatted algae powder. See **Table 5** and **Table 6** for the complete summaries.

Limit of Detection (LOD) and Limit of Quantification (LOQ)

The results of LOD and LOQ determinations are included in **Table 7**. The LOD of the respective fatty acids were determined as 0.12% ($C_{14:0}$), 1.8% ($C_{16:0}$), 0.06% ($C_{18:0}$) and 1.2% ($C_{22:6}$) and the LOQs were as 0.4% ($C_{14:0}$), 6% ($C_{16:0}$), 0.2% ($C_{18:0}$) and 4% ($C_{22:6}$).

Stability of DHA in *Aurantiochytrium limacinum* Biomass with a Natural or Synthetic Antioxidant

The stability data over 24 months for *Aurantiochytrium limacinum* containing the synthetic antioxidant, ethoxyquin, demonstrates that ethoxyquin was very

Table 4. Accuracy of fa	tty acid recoverie.	s (%) in de-fat	ted unextracted	l biomass of	Aurantiochytrium	limacinum	algae	powder
spiked to three concentration	ation levels.							

Primary Analyst	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}
Approx. Spiked Conc (%)	1.7	18	0.6	10	2.0	22.0	0.8	12	3.3	34	1.4	22
Replicate-1	96.0	96.0	86.6	117.1	95.4	90.6	96.6	106.6	92.8	86.0	88.7	106.2
Replicate-2	87.2	88.2	88.5	113.1	89.7	82.6	86.9	102.8	96.8	91.3	91.9	112.8
Replicate-3	92.7	100.3	95.1	113.0	100.8	96.7	99.7	117.4	100.5	94.3	93.2	113.6
Replicate-4	97.5	95.6	97.8	119.2	99.1	93.9	88.9	108.1	100.1	95.0	94.5	114.4
Replicate-5	89.1	82.3	80.7	111.8	93.9	89.9	91.0	115.5	96.8	91.5	91.8	111.3
Replicate-6	99.1	94.7	91.9	112.8	99.6	96.0	91.5	114.5	92.7	86.1	89.1	106.6
Replicate-7	94.2	93.3	100.3	117.1	87.3	83.2	94.1	108.1	91.8	86.6	83.1	103.9
Replicate-8	82.9	82.9	97.0	100.7	85.6	81.0	86.5	104.9	92.1	89.3	90.0	109.6
Replicate-9	93.5	90.0	96.5	113.8	82.0	80.9	87.6	104.2	90.8	87.1	88.0	105.1
Mean, primary analyst	92.5	91.5	92.7	113.2	92.6	88.3	91.4	109.1	94.9	89.7	90.0	109.3
Std Deviation	5.2	6.1	6.3	5.3	6.8	6.5	4.6	5.3	3.7	3.5	3.4	3.9
RSD (%)	5.6	6.7	6.8	4.7	7.3	7.3	5.0	4.9	3.9	3.9	3.8	3.6
Secondary Analyst	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}
Spike-1	87.3	88.6	83.5	115.6	86.9	83.8	80.3	98.8	97.5	94.8	93.5	114.8
Spike-2	86.3	80.9	81.3	112.3	87.8	81.4	81.0	101.1	92.7	88.4	86.4	106.3
Spike-3	89.4	88.4	94.9	111.9	88.9	85.6	80.3	106.2	97.2	93.0	94.7	113.4
Mean, secondary analyst	87.7	86.0	86.6	113.3	87.9	83.6	80.5	102.0	95.8	92.1	91.5	111.5
RE (%) to Primary Analyst	-5.2	-6.0	-6.6	0.1	-5.1	-5.3	-11.9	-6.5	0.9	2.7	1.7	2.0

RSD (% = (SD/Mean) * 100; RE (Relative error (%)) = ((mean, 2nd analyst - mean, 1st analyst) * (mean, 1st analyst)) * 100.

 Table 5. Precision of fatty acid recoveries (%) from the unextracted biomass of Aurantiochytrium limacinum algae powder spiked to three concentration levels.

Intra-day Precision:	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}
Approx. Spiked Conc (%)	3.2	41.3	1.7	25.7	3.9	48.0	2.4	32.4	4.6	54.6	3.0	39.1
Day 1 \overline{x} (n = 6)	102.6	102.0	99.5	101.9	102.7	101.9	99.5	105.1	101.5	100.0	97.9	104.5
SD	1.5	1.6	1.5	2.3	1.5	1.6	1.2	1.6	1.2	1.1	1.2	1.4
% CV	1.5	1.6	1.5	2.3	1.5	1.5	1.2	1.5	1.2	1.1	1.2	1.4
Day 2 \overline{x} (n = 3)	100.3	100.7	98.8	102.1	99.9	99.9	97.1	101.8	101.8	101.7	99.3	104.5
SD	0.8	0.7	0.4	1.1	3.3	2.5	4.8	4.4	1.1	1.6	2.0	2.2
% CV	0.8	0.7	0.4	1.1	3.3	2.5	4.9	4.3	1.1	1.6	2.0	2.1
Day 3 \overline{x} (n = 3)	99.2	97.9	96.2	98.0	101.0	99.6	97.7	101.1	101.3	99.6	97.6	101.6
SD	1.3	1.3	0.9	1.0	0.8	0.6	0.9	0.9	0.5	0.5	0.3	0.5
% CV	1.3	1.3	1.0	1.0	0.8	0.6	0.9	0.8	0.5	0.5	0.3	0.5
Inter-day Precision:	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}
$\overline{\mathbf{x}}$ (n = 3)	101.2	100.6	98.5	100.9	101.6	100.8	98.4	103.3	101.5	100.3	98.2	103.8
SD	1.9	2.1	1.8	2.4	2.2	1.9	2.5	2.9	1.0	1.3	1.4	1.9
% CV	1.9	2.1	1.8	2.4	2.1	1.9	2.5	2.8	1.0	1.3	1.4	1.8

Intra-day Precision:	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}
Approx. Spiked Conc (%)	1.7	18	0.6	10	2.0	22.0	0.8	12	3.3	34	1.4	22
Day 1 \overline{x} (n = 6)	93.6	92.9	90.1	114.5	96.4	91.6	92.4	110.8	96.6	90.7	91.5	110.8
SD	4.8	6.5	6.2	2.9	4.2	5.2	4.8	5.8	3.4	3.9	2.3	3.6
% CV	5.1	7.0	6.9	2.6	4.4	5.7	5.2	5.2	3.5	4.3	2.5	3.2
Day 2 \overline{x} (n = 3)	90.2	88.7	97.9	110.5	85.0	81.7	89.4	105.7	91.6	87.7	87.0	106.2
SD	6.3	5.3	2.1	8.7	2.7	1.3	4.1	2.1	0.7	1.4	3.6	3.0
% CV	7.0	6.0	2.1	7.8	3.2	1.6	4.6	2.0	0.7	1.6	4.1	2.8
Day 3 \overline{x} (n = 3)	87.7	86.0	86.6	113.3	87.9	83.6	80.5	102.0	95.8	92.1	91.5	111.5
SD	1.6	4.4	7.3	2.0	1.0	2.1	0.4	3.8	2.7	3.3	4.5	4.6
% CV	1.8	5.1	8.4	1.8	1.1	2.5	0.5	3.7	2.8	3.6	4.9	4.1
Inter-day Precision:	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}	C _{14:0}	C _{16:0}	C _{18:0}	C _{22:6}
Average	91.3	90.1	91.2	113.2	91.4	87.1	88.7	107.4	95.2	90.3	90.4	109.8
SD	5.0	6.1	6.8	4.6	6.2	6.0	6.3	5.8	3.4	3.5	3.5	4.0
% CV	5.5	6.7	7.5	4.1	6.7	6.9	7.1	5.4	3.5	3.8	3.9	3.7

 Table 6. Precision of fatty acid recoveries (%) from de-fatted unextracted biomass of Aurantiochytrium limacinum algae powder spiked to three concentration levels.

Table 7. Limit of Detection (LOD) and Limit of Quantitation (LOQ) in unextracted biomass of *Aurantiochytrium limacinum* algae powder spiked to three concentration levels.

Replicate No.	% C _{14:0}	% C _{16:0}	% C _{18:0}	% C _{22:6}
Replicate-1	2.56	34.59	1.07	18.86
Replicate-2	2.65	35.96	1.10	19.78
Replicate-3	2.60	35.08	1.06	19.27
Replicate-4	2.60	35.15	1.06	19.34
Replicate-5	2.66	36.01	1.09	19.80
Replicate-6	2.60	35.01	1.06	19.06
Replicate-7	2.61	35.21	1.07	19.32
Replicate-8	2.60	35.11	1.07	19.28
Replicate-9	2.53	34.27	1.05	18.73
Replicate-10	2.53	34.28	1.05	18.67
Average	2.59	35.07	1.07	19.21
SD	0.04	0.60	0.02	0.4
LOD $(3 \times SD)$	0.12	2	0.06	1.2
$LOQ (10 \times SD)$	0.4	6	0.2	4

effective protecting the LC-PUFA, including DHA, from oxidation (Table 8). Results for all data points over the study were within +/- 3.5% of the initial determination. As ethoxyquin is understood to have high antioxidant efficiency and stability and protects lipids against peroxidation, the consistency of DHA over 24 months was not unexpected [32].

Table 9 shows the results for the stability assessment of DHA in *Aurantio-chytrium limacinum* containing Duralox^{*} and shows that DHA can be effectively

Time	Mean %	SD	Minimum	Maximum	Range
(Month)	(n = 3)				
0	16.56	0.51	16.08	17.1	1.02
1	17.56	0.70	17.14	18.36	1.22
2	17.36	0.51	16.89	17.91	1.02
3	17.16	0.59	16.81	17.84	1.03
6	17.30	0.42	17.05	17.79	0.74
9	17.16	1.06	16.36	18.36	2.00
12	16.93	0.43	16.57	17.4	0.83
15	17.06	0.55	16.64	17.68	1.04
18	16.16	0.44	15.75	16.62	0.87
21	16.01	0.40	15.7	16.46	0.76
24	17.16	0.69	16.73	17.96	1.23

Table 8. The stability of DHA (%) in in unextracted biomass of *Aurantiochytrium limacinum* powder over 24 months using the antioxidant, ethoxyquin.

Table 9. The stability of DHA in in unextracted biomass of *Aurantiochytrium limacinum* powder over six months using the antioxidant, Duralox[®].

Time	Mean %	SD	Minimum	Maximum	Range
(Month)	(n = 4)				
0	17.86	0.19	17.71	18.11	0.40
1	17.94	0.23	17.66	18.23	0.57
2	18.04	0.14	17.88	18.23	0.35
3	17.81	0.30	17.52	18.22	0.70
6	18.20	0.37	17.71	18.56	0.85

stabilized over six months with the inclusion of this antioxidant. Results for all data points over the study were within +/-2% of the initial determination.

4. Conclusion

The suitability of the AOAC method 996.06 for the determination of DHA (C22:6) and three additional fatty acids ($C_{14:0}$, $C_{16:0}$ and $C_{18:0}$) in a dried microalgae biomass was investigated. The method was found to be linear over the following ranges for each FAME analyte; 50 to 15,000 µg/ml ($C_{14:0}$), 300 to 95,000 ($C_{16:0}$), 25 to 15,000 ($C_{18:0}$) and 300 to 59,375 ($C_{22:6}$). The method was verified for use by examining the method parameters, accuracy and precision, specificity and by determining the LOD and LOQ for each analyte. The results of the accuracy and precision experiments were within acceptable limits for the primary matrix, *Aurantiochytrium limacinum* biomass. The robustness of the method was confirmed through limited variability of the analysis precision and accuracy when investigated by a second analyst. Therefore, the method is deemed fit for purpose for the analysis of the microalgae, *Aurantiochytrium limacinum*. The method could be further used by other researchers who are investigating *Aurantiochytrium limacinum* and who need to elucidate the fatty acid composition in various applications. The method could also be employed in examining other similar algal species. The present authors intend to use the method for routine analysis, quality control & product release. The results of the stability assessments show that Duralox[®] can be used as an alternative to ethoxyquin for stabilizing DHA over six months. Further analysis will be required to show the efficiency over a long time period.

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Declaration of Conflicting Interests

The funders had no role in the design of the study; in the collection, analyses, or the interpretation of data. GD, AY and CM work for Alltech, who sponsored the research, contributed to the writing of the manuscript and in the decision to publish the results.

Conflicts of Interest

The authors declare that there is no conflict of interest regarding the publication of this paper.

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