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Extraction of common microalgae by liquefied dimethyl ether: influence of species and pretreatment on oil yields and composition

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Abstract

Liquefied dimethyl ether (DME) is regarded as a promising, green solvent for biomass lipid extractions. It is non-toxic, applicable to wet feedstocks, and allows easy product separation by pressure reduction. Yet, knowledge about its usability in combination with oleaginous microalgae is limited. In the current work, four common microalgae and cyanobacteria species were used to study DME extraction characteristics: *Arthrospira platensis, Nannochloropsis gaditana, Phaeodactylum tricornutum*, and *Scenedesmus almeriensis*. Dried samples were subjected to a batch DME extraction and compared to a standard chloroform/methanol procedure. To evaluate the influence of pretreatment, particle size distributions of two different milling sequences (knife- and cryo-milling) and the resulting effects on DME extraction and oil composition were addressed. Additionally, an algae washing procedure was tested. DME extractions resulted in oil yields of 0.5–2.7% of dry mass (equal to 5–19% of total lipids) without further pretreatment. Cryo-milling reduced median particle sizes by 25–87% and simultaneously increased lipid yields to 1.7–5.6% of dry mass (17–50% of total lipids). *Phaeodactylum tricornutum* showed the highest extraction efficiency with DME, combined with a favorable fatty acid profile. Although being most affected by the additional milling pretreatment, *Arthrospira platensis* performed worst in both scenarios. DME extracted oils were generally characterized by enhanced contents of C14:0, C16:0, and C16:1 fatty acids. However, relative abundances were strongly influenced by the properties of the tested algae species. The additional cryo-milling pretreatment affected fatty acid compositions by increasing the shares of potentially valuable polyunsaturated fatty acids.

Keywords DME · Pressurized liquid extraction · Aquatic biomass · Fatty acid profiles · Biorefinery · Bioeconomy

1 Introduction

Algae are considered important biomass source, which will play a crucial role to achieve an industrial transition to a climate-neutral production of chemicals, commodities, and fuels. Depending on species, microalgae can excel yields of land-bound energy plants by a factor of 5 [1] and produce up to 20 times more oil than typical agricultural plants [2]. The downside is very complex and energy-consuming

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¹ Department of Conversion Technologies of Biobased Resources, Institute of Agricultural Engineering, University of Hohenheim, Garbenstr. 9, 70599 Stuttgart, Germany photobioreactor systems, especially if the biomass has to meet specific quality standards. Besides, harvesting and dewatering are challenging and energy-consuming tasks [1, 3].

Recently, cascade-like biorefinery approaches are being designed in order to obtain multiple marketable products from one algae species. Most concepts rely on different extractions to obtain valuable substances suitable for nutrition, healthcare, or cosmetics [4], prior to a possible end-of-pipe conversion.

During solid-liquid extraction, a targeted compound is transferred from the solid phase (biomass) to an immiscible liquid phase (solvent). In order to be successful, any extraction solvent must be able to penetrate the matrix enclosing the lipid material, make physical contact with the lipid, and solvate the lipid [5]. The process is highly dependent on solvent properties, such as solubility of the targeted compound(s), selectivity, or viscosity [6]. Processing parameters can affect those properties and enhance extraction efficiencies. High flow rates, for example, reduce the boundary layer of concentrated solute at particle surfaces [6] and elevated temperatures may decrease surface tension and viscosity of the solvent [7]. Traditionally, organic solvents, such as chloroform, hexane, or dichloromethane, are applied to recover lipids and other valuable compounds from biomass [8, 9]. However, with an increasing need for sustainable and non-toxic extractions, alternative concepts like supercritical fluids, gasexpanded liquids, or switchable solvents arose [7, 10]. The most prominent example, supercritical CO₂ (scCO₂) extraction, has already been applied for several decades. Besides commercial oil recovery from sesame seeds [11] or decaffeination of coffee [12], it has been used to extract various species of microalgae [13–16]. However, feedstocks with high water contents can represent a severe barrier for scCO₂ extractions [17] and its low polarity often requires the addition of an entrainer [15, 16].

In the case of algae, mechanical dewatering is only viable to about 80% dry matter and thermal drying requires high amounts of energy [18]. Consequently, particularly the processing of wet feedstocks is a key technology to enable algaesuccess on a large scale [19, 20].

Dimethyl ether (DME) offers many major advantages if used as extraction solvent. At normal conditions, DME is a colorless gas (normal boiling point: -24.82 °C [21]) with an ethereal odor [22]. It is categorized as non-toxic [23] and only needs a relatively low amount of energy for evaporation and pressurization [24]. The consecutive product separation via pressure reduction is simple and the temperature during extraction is normally set to 20-30 °C, which is mild enough to preserve the most valuable compounds. Furthermore, water is co-extracted and does not inhibit the extraction to a high extent [25] or, in some cases, even enhances the solvent's performance [26]. After DME extraction, a dry biomass residue is left, which contains high amounts of proteins and carbohydrates.

In contrast to many common solvents [8, 18, 27, 28], literature suggests that an energy-intensive pretreatment (e.g., bead milling, microwaving, or steam explosion) can be avoided when DME is used on wet feedstocks [29–31]. If this finding is related to the solvent itself, biomass water content or feedstock properties are largely unknown.

Up to now, DME was applied to multiple species of algae (and cyanobacteria), such as *Haematococcus pluvialis* [30], *Oscillatoria agardhii* [32], or *Botryococcus braunii* [24]. In many cases, oil yields were reported to be comparable with standard extractions [24, 29, 32, 33]. Different blue-green microalgae showed DME recoveries between 97.0 and 99.7% compared to a common chloroform/methanol procedure [32]. For *B. braunii* and *H. pluvialis*, DME yielded 99.1% and 69.8% of hexane and acetone extractions, respectively [24, 30].

In comparison to scCO₂, DME extracted 446% of oil during the treatment of wet *A. platensis* [34]. Non-microalgae sources, such as tuna liver [35] or labyrinthula *Aurantiochytrium limacinum* [31], showed recoveries of roughly 100% and 205%.

To enable an evaluation of different feedstocks under similar processing conditions, the current study relies on a simple batch system. It was assumed that differences in morphological properties, such as cell wall characteristics, lead to specific interactions between solvents and algae. The tested species were *Arthrospira platensis* (AP, Cyanophyceae), *Nannochloropsis gaditana* (NG, Eustigmatophyceae), *Phaeodactylum tricornutum* (PT, Bacillariophyceae), and *Scenedesmus almeriensis* (SA, Chlorophyceae) [36, 37]. They differ in structure and composition and were used in a dried state. NG, SA, and PT were selected because they offer a combination of high lipid productivity and sufficient content of valuable fatty acids [38–43]. AP was included as it is cultivated all around the world, contains high-quality proteins, and is well known among consumers [44, 45].

The four species were extracted with DME and compared to a common Folch based [46, 47] chloroform/methanol Soxhlet procedure with regard to oil yields and oil composition. DME was applied to differently milled samples in order to study the effects of pretreatment. In addition, a simple algae washing procedure [48] was applied and evaluated, which should help to correct biomass ash contents and extraction yield calculations.

Overall, the present work is intended to promote the understanding of the influences of algae species and pretreatment on DME extraction performance and oil quality.

2 Materials and methods

2.1 Microalgae feedstock

Freeze-dried *Nannochloropsis gaditana* (NG) and *Scenedesmus almeriensis* (SA) were obtained from the cultivation facility in Las Palmerillas (University of Almería, Spain). *Phaeodactylum tricornutum* (PT) slurry was provided by Roquette Klötze GmbH, Germany, and freeze-dried prior to use. Spray-dried *Arthrospira platensis* (AP) chips originated from the IGV GmbH, Germany. Additional information about the suppliers' drying procedures can be found in Online Resource 2.

2.2 Milling

Prior to experiments and analysis, the relatively inhomogeneous biomass samples (pictures of original feedstocks available in Online Resource 1) were milled to uniformity with the help of a knife mill (Pulverisette 19, Fritsch). It was equipped with a stainless steel knife rotor with five "V" shaped blades, which rotates inside the milling chamber (2800 rpm) against three fixed blades until the material passed a 0.5 mm screen. To further reduce the particle size and break open functional cell structures, a part of the samples was subjected to an additional ball milling (CryoMill, Retsch). Here, 5-7 g of knife-milled sample and a ø20 mm stainless steel grinding ball were loaded into a 35 ml grinding jar, which is inserted in the mill to perform radial oscillations in a horizontal position. Automatic pre-cooling and 30 s intermediate cooling with liquid nitrogen were applied and the samples were milled in two 3 min cycles with at a frequency of 25 Hz.

Figure 1 shows the knife- and cryo-milled algae feedstocks, which were subjected to the extractions.

2.3 Particle size analysis

Particle size distributions (PSD) of knife-milled and cryomilled samples were measured by a laser diffraction particle size analyzer (PSA 1190, Anton Paar). Prior to the measurement, microalgae samples were dispersed in ethanol at slow stirring speed and treated with ultrasound at medium stirring speed for 10 min. Measurements were conducted afterward at



Fig. 1 Knife- and cryo-milled algae feedstocks. AP, Arthrospira platensis; NG, Nannochloropsis gaditana; PT, Phaeodactylum tricornutum; SA, Scenedesmus almeriensis

medium stirring speed without further sonication. For each sample, at least two consecutive measurements were done and a mean value was calculated. In addition, mean D10, D50, and D90 values were calculated, which represent equivalent spherical particle diameters of the 10th, 50th, and 90th percentiles of total volume.

2.4 Algae characterization

Dry weight of the different microalgae feedstocks with the respective milling pretreatments was determined by drying 0.5 g of sample at 60 °C for 48 h (based on the NREL method NREL/TP-5100-60943) [49]. The determination was done in quadruplicate.

To obtain the ash content, dry samples were subjected to 575 °C in a muffle furnace according to the NREL algae biomass method (NREL/TP-5100-60956) [50] with three replicates.

Since the ash content of the studied samples was very high in some cases, a washing procedure was tested, which should help to correct dry matter contents in the following calculations (as extraction yields are based on dry matter). In accordance with Zhu and Lee, algae were washed with an isotonic 0.5 M ammonium hydrogen carbonate solution (99% purity, Acros Organics) [48]. Therefore, 0.3 g of each species (knifemilled, dry) was mixed with 35 ml of washing solution in a 50 ml falcon tube, vortexed for 20 seconds, and centrifuged for 10 min at 10,000 rpm under room temperature. The supernatant was vacuum filtrated over a pre-weight, dry 0.2 µm filter and washed with an additional 20 ml of ammonium hydrogen carbonate solution. The tube, containing the washed algae pellet, and the filter were dried at 60 °C for 48 h and weighed to determine the washed dry matter. This treatment was conducted in triplicate. The ash content of washed samples was determined as stated above.

A CHNS Analyzer (HEKAtech Euro EA, Germany) was used to determine C, H, N, and S contents of washed and unwashed algae samples (dry and finely ground) in duplicate. The oxygen content was calculated by difference.

2.5 Lipid extraction procedures

Crude lipid extracts were obtained via modified Folch (2.5.1) and DME (2.5.2) extractions. The experiments were done in triplicate. For a better overview, Fig. 2 contains the applied methods and the most important parameters.

2.5.1 Soxhlet extraction (modified Folch method, adapted from [46])

In brief, 2 g of knife-milled sample was weighed into a cellulose crucible and subjected to a 20 h extraction with 200 ml of a chloroform/methanol mixture (2:1, v/v) in a



Fig. 2 Applied methods for algae lipid extraction

common Soxhlet apparatus (chloroform was purchased from Merck with 99.0% purity, methanol from VWR with \geq 99.9% purity). Cryo-milled samples were not further investigated here, as the fine particles clumped together and inhibited the extraction in most cases (data not shown). Heating was set to minimum with an Isopad heater (Tyco Thermal Controls) to ensure a number of at least 4 solvent cycles per hour. In the following, the extract was vacuum filtrated (0.2 µm pore size) and filled up to 220 ml with fresh chloroform/methanol (2:1, v/v), which was also used to wash the involved glassware and filter three times. After the addition of 55 ml 0.7% NaCl (aq) solution (99.9% purity, VWR) up to a final ratio of 8:4:3 chloroform/methanol/NaCl (aq), the mixture was shaken and thereafter allowed to settle in a separation funnel for at least 20 h. The lower, organic phase was transferred into a pre-weight flask, evaporated in the rotary evaporator (50 °C), and additionally dried at 50 °C for 24 h before the oil yield was determined gravimetrically.

2.5.2 Dimethyl ether extraction

For the DME extraction, 4 g of milled algae (knife- or cryomilled) was loaded into a batch extractor (80 ml, pressureresistant borosilicate glass flask with a screw lid) which was connected to an unlubricated stainless steel valve and mixed with 40 ± 3 g of liquefied DME (\geq 99.9% purity, Air Liquide). Figure 3 shows an extractor with dimensions and additional information. The loaded extraction vial was shaken for 90 min with high intensity (Köttermann 4020 shaker) at room temperature. Pressure occurred as a result of temperature and ranged between roughly 0.5–0.6 MPa [25]. In the following, about one-third of the liquid DME was directed over a 2 μ m sintered stainless steel filter (FITOK) into a second, empty, pre-weighed extractor vial. The transferred extract was weighed and eventually evaporated naturally.

The leftover oil was solved in 20 ml chloroform/methanol (2:1, v/v) and vacuum filtrated $(0.2 \mu m)$. Glass parts and filter were washed with an additional 10 ml of solvent mixture, which was added to the liquid. This combined extract was blended with 7.5 ml of 0.7% NaCl (aq) solution (to a final ratio of 8:4:3 chloroform/methanol/NaCl (aq)), shaken briefly, and allowed to settle in a separation funnel for at least 20 h.



Fig. 3 DME extraction vessel

The organic phase was transferred into a pre-weight flask. After rotary evaporation (50 °C) and drying the flask at 50 °C for 24 h, the oil yield was determined gravimetrically and calculated back onto the total amount of oil solved in the original flask and the total extraction yield according to formulas (1) and (2):

$$m_t = \frac{m_{\rm DME1}}{m_{\rm DME2}} \times m_{\rm Oil2} \tag{1}$$

where m_t is the mass of total oil extracted (g); m_{DME1} is the mass of DME in first extractor vial (g); m_{DME2} is the mass of DME in second extractor vial (g); m_{Oil2} is the mass of oil in second extractor vial (g).

$$M_t = \frac{m_t}{m_a} \times 100 \tag{2}$$

where M_t is the total oil yield (% of algae dry mass); m_t is the mass of total oil extracted (g); m_a is the mass of dry microalgae used (g).

2.6 Fatty acid profile analysis

To determine the fatty acid profiles, mixed oil samples of at least three identical extraction trials were prepared. They were solved in chloroform immediately after weighing and stored at 4 °C prior to analysis.

The following sample preparation consisted of evaporation of the chloroform, hydrolysis with 0.1 M hydrochloric acid (40 ml per g oil) for 1 h, filtration (filter paper MN 715 ¼, Macherey-Nagel), and neutralization with cold water. Extraction of the fatty acids was carried out on the dried retentate with petroleum ether for 6 h in a Soxhlet apparatus (30–50 °C). Then, 0.2–0.3 g of the pretreated, dried sample extract was dissolved in 5 ml of diisopropyl ether. Subsequent methylation was done by adding a 0.2 M methanolic solution of trimethylsulfoniumhydroxide (TMSH) to an aliquot of the sample extract dissolved in diisopropyl ether in a ratio of 1:2 (v/v). For analysis, this mixture was injected into a GC with FI detection (7890A, Agilent Technologies), equipped with a capillary column DB-23, 30 m \times 0.250 mm, film 0.25 μ m (Agilent Technologies).

The fatty acid standard mixture to be used for calibration consists of 37 individual components (C4:0 to C24:1). The amount of a methyl ester in % was calculated by determining the percentage corresponding to its peak area relative to the total sum of all FAME peak areas in the chromatogram. The calculated value was expressed as % relative to the total fatty acids.

2.7 Data evaluation

For each of the experiments (excluding GC-FID analysis), samples of a larger amount of homogenized feedstock were used. In the cases of elemental analysis and particle size distribution, the respective analyses were carried out in duplicate according to established best-practiced methods and a mean value was calculated. The other experiments' results were expressed as the average of at least three independent trials. In the case of the extraction experiments, standard deviations were additionally presented to allow an estimation of measurement uncertainties, mainly associated with the manual use of a separation funnel and the very low amount of product generated during each step.

3 Results

In the following sections, % represents g 100 g^{-1} dry weight unless stated differently. First, the influence of feedstock is reported.

3.1 Feedstock

Table 1 contains basic data about the microalgae from characterization experiments and literature.

Dry matter contents of all feedstocks ranged between 95.3 and 97.5% and only small differences could be found between cryo-milled and knife-milled samples.

According to literature, NG and PT show lipid yields between roughly 20 and 32% and SA between 11 and 22%. Even though AP was also reported to yield up to 16%, it is usually rather settled in the 4–9% region. Considering proteins, up to 70% can be found in AP, while PT ranks lowest with around 37%.

The results of the elemental analysis (not-washed feedstocks) displayed that the carbon contents of AP and SA were located at nearly 50%. In contrast, PT and NG showed remarkably lower C contents with 41.2% and 34.8%, respectively. Nitrogen contents of the tested species ranged between 5.1 and 10.0%. The amounts of oxygen, hydrogen, and sulfur were relatively similar between the algae.

A large difference among the species could be identified for the ash contents. Especially NG and PT showed high values for ash (27.0% and 16.7%), while SA was settled at 9.9% and AP at only 6.2%.

3.2 Pretreatments

3.2.1 Algae washing

Washing with ammonium hydrogen carbonate solution reduced the ash content and consequently increased the proportions of C, H, and N in all cases (Table 1). The most remarkable increase was observed in the C content of NG and accounted for nearly 10.4%. At the same time, in three of the four species (except for NG), the proportion of O was lower after washing. Figure 4

	Arthrospira platensis (AP)		Nannochloropsis gaditana (NG)		Phaeodactylum tricornutum (PT)		Scenedesmus almeriensis (SA)		
	Pretreated with knife mill (K) or cryo-mill (C)								
	Κ	С	Κ	С	К	С	Κ	С	
Dry weight (g 100 g^{-1} fresh weight)	95.30	95.95	97.44	97.49	95.96	96.74	96.54	96.41	
	Literature values								
	4.0–16.0 ⁱ		22.0–29.0 ⁱⁱ		20.0–32.2 ⁱⁱⁱ		11.2–21.8 ^{iv}		
Proteins (%)	46.0–70.0 ^v		36.0-49.0 ^{vi}		36.4–37.5 ^{vii}		41.8–51.7 ^{viii}		
	Not-washed (-) or washed** (w) feedstock								
	-	W	-	W	-	W	-	W	
C (%)	48.10	50.23	34.77	45.15	41.23	44.36	46.67	50.05	
Н (%)	6.78	7.31	5.11	6.72	6.05	6.70	6.69	7.23	
N (%)	9.95	11.43	5.07	7.01	6.75	8.12	7.05	8.28	
S (%)	0.55	0.52	0.82	1.09	0.87	0.96	0.94	0.75	
O (%)*	28.39	27.49	27.27	31.69	28.43	26.96	28.76	27.38	
Ash (%)	6.23	3.02	26.97	8.33	16.67	12.90	9.89	6.31	

 Table 1
 Properties of microalgae feedstocks (own experiments and literature data). Own values are expressed as a mean of two (elemental analysis) and three (dry weight determination) replicates

*Calculated by difference

**Washing conducted with knife-milled samples

^{i-viii} References: ⁱ [45, 51, 52], ⁱⁱ [53–56], ⁱⁱⁱ [43, 55–58], ^{iv} [36, 55, 59], ^v [45, 51, 52], ^{vi} [53, 55], ^{vii} [52, 55], ^{viii} [55, 59]

shows ash contents and ash-free dry mass (mass DAF) of all algae tested in % of initial dry mass.

Decreases of ash contents ranged between 3.8% (AP) and 22.2% (NG) of initial dry mass. PT and SA values were reduced by 7.5% and 4.7% of initial dry mass (Fig. 4). The lowest ash content after washing was measured in AP, while PT ranked highest (Table 1).

At the same time, ash-free dry mass was also removed during the washing. This led to further losses of 15.1%, 20.8%, 12.7%, and 21.3% of initial dry mass for AP, NG, SA, and PT. Alternative trials with two consecutive washing cycles led to greater removal of ash-free dry mass (Data not shown).

3.2.2 Algae milling

Table 2 and Fig. 5 contain the volume-based results of D10, D50, and D90 values and particle size distributions for knifemilled and cryo-milled algae samples, respectively.

Varying particle size distributions were found among the different feedstocks after both milling pretreatments.

PSD of knife-milled AP showed a peak at 400 μ m and a median particle size (D50) of 171.5 μ m, meaning that 50% of the particles had a size \leq 171.5 μ m. The other three species (knife-milled) consisted of smaller particles and peaked at 63 μ m (NG and PT) and 75 μ m (SA). Median particle sizes were also considerably lower with 39.0–54.3 μ m.

For each sample, a small particle fraction between 0.2 and $0.8 \ \mu m$ was found, which was increased after cryo-milling

(except for NG). It also decreased median particle sizes significantly to 12.2–33.7 μ m. PSD curves of AP and SA were changed drastically (new maxima at 50 μ m and 14 μ m, respectively), whereas PT was less impacted (new maximum at 50 μ m). The NG maximum was retained at 63 μ m, but the number of particles in this size-division was reduced.

3.3 Lipid extraction

In Fig. 6, crude oil yields are shown for the Soxhlet and DME extractions.

The Soxhlet (Folch) method with knife-milled feedstocks is expected to provide a good approximation of total lipid biomass contents and will be considered "total extractable oil" in the following.

Soxhlet oil yields ranged from 9.7% (NG) to 15.9% (SA) of dry mass. DME showed yields of 0.5% (AP) to 2.7% (SA) for knife-milled and 1.7% (AP) to 5.6% (SA) for cryo-milled feedstocks.

Of the tested species, AP revealed the lowest compatibility with the DME extraction procedure. Depending on pretreatment, extraction efficiencies between 5% (knife-milled) and 17% (cryo-milled) were realized by DME. NG yielded 8–18% and SA 17–35% for knife- and cryo-mill pretreatments, respectively. The highest efficiencies were achieved with PT, which ranged from 19% (knife-milled) to 50% (cryo-milled). Milling increased the yields in each case but showed different changes depending on the species. AP was most influenced, as DME yields increased by a factor of 3.5. PT, NG, and SA recoveries were

Fig. 4 Ash content and ash-free dry mass (mass DAF) of notwashed (-) and washed (w) algae as % of initial dry mass. Values are expressed as a mean of three replicates



Algae species and pretreatment

mass DAF

enhanced by factors of 2.0–2.6 by the additional milling. Each of the extracts with either method was colored in a dark greenishbrown and characterized by a high viscosity at room temperature.

3.4 Fatty acid compositions

Table 3 contains the fatty acid compositions of the extracted oils.

The four algae species displayed major differences in fatty acid compositions. Palmitic acid (C16:0) was found in every alga, although only for AP it was the main fatty acid with shares of 56.0-63.6%. Linoleic acid (C18:2n6) and γ -linolenic acid (GLA, C18:3n6) ranged between 11.7 and 14.9% and 9.3 and 12.6% for this species.

In NG and SA oils, α -linolenic acid (C18:3n3) was dominant with shares between 41.7–45.0% and 41.8–51.0%, respectively. Both species showed a relatively similar fatty acid profile with the remaining major fatty acids being C16:0, C16:1, C18:1n9, and C18:2n6.

For PT, C16:1 (31.5–37.5%) and C16:0 (23.0–28.6%) were dominant. In addition, eicosapentaenoic acid (EPA, C20:5n3) could be found in significant quantities (11.2–18.5%).

The total polyunsaturated fatty acids (PUFA) content of AP (about 23–27%) was roughly doubled by NG (56–59%). SA

ranked highest with 55–65% and PT lowest with only 14–22%. However, in contrast to the other three species, EPA was the dominant fatty acid and represented around 80% of the PUFA extracted from PT.

Ash

Concerning saturated fatty acids, AP oil contained by far the highest share (60–67%), followed by PT (36–42%), NG (30–34%), and SA (15–25%). Monounsaturated fatty acids were highest in PT (40–44%) and lowest in AP (7–13%).

The method of extraction and the respective solvents affected fatty acid compositions. With regard to the total numbers of saturated and monounsaturated fatty acids in knife-milled extractions, DME showed higher shares in AP and PT (approx. + 5%), was about even in NG, and around 9% lower in SA. Total PUFA shares behaved vice versa. In addition, DME led to an increase in relative contents of C14:0 and C16:0 for AP, NG, and PT and in relative content of C16:1 for AP, PT, and SA.

The effect of cryo-milling prior to DME extraction was apparent: total PUFA shares increased in each case, while the proportion of saturated and monounsaturated fatty acids was either nearly constant or decreased. This was by far most pronounced in PT, where the EPA share increased from 11.2 to 18.5% and total PUFA from 14.2 to 22.1% after cryo-milling.

Table 2Calculated mean valuesof at least 2 replicates for 10th(D10), 50th (D50), and 90th(D90) percentiles of total volumefor knife-milled (K) and cryo-milled (C) algae samples

	Arthrospira platensis (AP)		Nannochloropsis gaditana (NG)		Phaeodactylum tricornutum (PT)		Scenedesmus almeriensis (SA)	
	K	С	K	С	K	С	K	С
D10 (µm)	17.1	2.6	5.4	3.3	4.9	2.5	7.8	2.3
D50 (µm)	171.5	21.6	45.0	33.7	39.0	18.1	54.3	12.2
D90 (µm)	399.5	96.3	99.7	106.6	91.2	60.8	146.9	43.4





4 Discussion

4.1 Feedstock

Regarding the biomass water contents, cryo-milling only led to negligible changes for the tested samples (Table 1).

For both of the applied extraction methods, water can be tolerated to a certain extent. DME is miscible with water but may undergo a change of polarity, depending on the ratio of solvent to water [60, 61]. The applied chloroform/methanol system (2:1, v/v) is also applicable with small amounts of water but exhibits a phase separation above 3% water in the solvent mixture [62]. Also, in the case of wet biomass, a solvent shell can be formed, which may inhibit the extraction by the less polar chloroform [5]. Consequently, as the influences of biomass water contents on both extraction systems are expected to be high, only dried, stable feedstocks were tested. In this way, the effects of solvents and pretreatments on yields and composition of the oils could be studied and water-induced side effects and interferences were minimized.

Fig. 6 Crude oil yields (% dry mass) of algae extracted via chloroform/methanol (Soxhlet, knife-milled) and liquefied dimethyl ether (DME, knife- and cryo-milled). Values are expressed as a mean of three replicates; error bars represent standard deviation



 Table 3
 Fatty acid profiles (% of total fatty acids) of dimethyl ether (DME) and chloroform/methanol (Soxhlet) extracted oils. K, knife-milled algae feedstock; C, cryo-milled algae feedstock; SFA, saturated fatty

acids; MUFA, monounsaturated fatty acids; PUFA, polyunsaturated fatty acids; –, below 0.01% of total analyzed fatty acids

	Arthrospira platensis (AP)		Nannochloropsis gaditana (NG)			Phaeodactylum tricornutum (PT)			Scenedesmus almeriensis (SA)			
	Soxhlet	oxhlet DME		Soxhlet DME		Soxhlet	DME	DME		DME		
	К	K	С	К	K	С	Κ	K	С	K	K	С
C4:0	0.20	0.02	_	0.05	_	0.14	_	0.02	_	0.05	_	_
C6:0	_	_	_	_	_	0.03	-	-	-	-	-	_
C8:0	-	0.05	-	-	-	0.07	-	-	-	_	-	-
C10:0	-	0.40	-	-	-	0.06	-	-	-	-	-	-
C11:0	-	-	-	-	-	-	-	-	-	-	-	-
C12:0	-	0.20	0.10	-	-	0.12	-	0.04	0.02	_	-	-
C13:0	-	-	_	-	-	-	-	-	0.03	_	-	-
C14:0	0.40	0.54	0.51	0.77	1.06	0.88	8.24	10.80	8.67	0.50	0.50	0.30
C14:1	-			0.02	0.02	-	0.14	0.35	0.04	_	0.02	0.02
C15:0	0.19	0.18	0.19	0.47	0.55	0.53	0.46	0.58	0.48	0.13	0.12	0.10
C15:1	-	-	-	-	-	-	-	-	-	-	-	-
C16:0	56.02	61.19	63.60	21.14	21.99	19.29	23.03	28.58	25.43	21.59	13.79	13.31
C16:1	5.75	6.03	6.43	4.82	2.96	2.54	31.54	37.54	35.19	5.39	6.04	5.61
C17:0	0.76	0.59	0.67	0.86	0.63	0.31	6.45	-	-	0.49	0.50	0.41
C17:1	0.67	_	0.87	0.91	-	0.19	1.69	-	-	0.43	-	0.05
C18:0	2.26	1.92	1.80	2.66	3.30	2.48	1.18	0.60	0.71	1.34	0.88	0.87
C18:1n9	6.22	6.25	-	4.21	9.71	8.88	4.52	4.65	4.54	14.28	14.18	14.85
C18:2n6	14.00	11.72	14.86	13.30	12.31	12.63	1.21	1.06	1.57	12.28	13.08	12.81
C18:3n3	0.18	0.27	0.08	41.66	42.88	44.95	0.38	0.38	0.44	41.79	49.69	51.02
C18:3n6	12.62	9.75	9.29	-	0.07	0.09	0.55	0.45	0.59	0.40	0.28	0.28
C20:0	0.15	0.14	0.19	6.52	3.27	4.41	0.10	0.09	0.06	0.06	0.06	0.03
C20:1	0.18	0.17	0.15	0.29	0.28	0.15	0.57	0.09	-	0.24	0.16	0.22
C20:2	0.18	0.18	0.20	0.04	-	0.21	0.18	0.10	0.11	-	-	0.02
C20:3n3	0.24	0.17	0.23	-	-	-	-	0.06	0.05	-	-	-
C20:3n6	-	-	_	0.14	0.15	0.18	-	-	-	0.07	_	0.03
C20:4n6	_	_	_	_	_	_	0.63	0.47	0.71	-	_	_
C20:5n3	_	0.59	0.72	0.44	0.23	0.43	16.30	11.22	18.54	0.13	0.46	0.29
C21:0	_	_	_	_	0.16	0.20	0.37	-	-	-	_	_
C22:0	_	_	_	1.27	0.44	0.75	0.20	0.08	0.11	0.59	0.26	0.47
C22:1n9	_	_	_	_	_	-	-	0.44	0.04	-	_	_
C22:2	-	-	-	-	-	0.28	-	-	-	-	-	0.15
C22:6n3	-	-	-	-	-	-	-	0.46	0.13	-	-	-
C23:0	-	-	0.14	0.44	-	-	-	-	-	0.25	-	-
C24:0	-	-	-	-	-	0.25	1.07	1.29	0.93	-	-	-
C24:1	_	_	_	_	-	_	1.20	0.67	-	-	-	-
\sum SFA	59.98	65.23	67.20	34.18	31.40	29.52	41.10	42.08	36.44	25.00	16.11	15.49
\sum MUFA	12.82	12.45	7.45	10.25	12.97	11.76	39.66	43.74	39.81	20.34	20.40	20.75
\sum PUFA	27.22	22.68	25.38	55.58	55.64	58.77	19.25	14.20	22.14	54.67	63.51	64.60

However, with regard to a final application, systems that can tolerate high water contents are highly desirable [19, 20]. Experiments with varying water contents will be part of a consecutive study. With a view of previously reported lipid values, only the Soxhlet trials of AP and SA lie in the expected region (Table 1 and Fig. 6). The yield of NG revealed to be less than half of the predicted value. This finding, combined with the fatty acid profile found for NG (Table 3) led to the conclusion that the feedstock was most likely fundamentally different from a usual *Nannochloropsis gaditana* or heavily contaminated during cultivation (also see 4.4). Most other differences between reported and measured values can be owed to additional factors that influence microalgae compositions, such as subspecies, culture conditions (e.g., starvation, light regime, culture medium), and culture age [63, 64].

AP showed the highest nitrogen concentration, which correlates well with the expected, high protein content [44, 45]. The overall low contents of C, H, and N found in NG can be explained by the very high amount of ash. It accounted for over 25%, which results in a lower proportion of biomass constituents and calculated extraction efficiency. Ash contents of PT and SA were also relatively high, but the effect was less pronounced.

The marine species NG and PT are mostly cultivated in saline media. It is well known that the growth media has a high impact on cell composition of microalgae [65] and that salts can be absorbed on the cell surface and/or be solved in intercellular water [48]. To further address this issue, algae washing was conducted.

4.2 Pretreatments

4.2.1 Algae washing

The washing was mainly conducted in order to minimize the effects of adsorbed salts or non-cellular suspended solids (e.g., calcium carbonate) [66] on oil extraction yield calculations. For each of the tested species, washed samples showed lower ash contents than untreated ones, which suggests a successful removal. As a consequence, the relative contents of C, H, and N increased (Table 1). Only oxygen was rather unchanged, which can partly be explained by the nature of some common salts which are expected to have been washed out (e.g., Na₂SO₄, NaNO₃) [67]. The washing was very effective for the marine species—especially from NG and to a smaller extent from PT, substantial parts of ash were removed.

However, simultaneously, a large proportion of ash-free dry mass was lost during the treatment of each species (Fig. 4). After the washing, between 71 and 83% of the original biomass dry mass were left for AP, PT, and SA. In all three cases, the impact of mass DAF removal on total weight after washing was greater than the impact of ash removal. The most extreme result was obtained for NG, where only 57% of the original biomass retained.

It can be assumed that the algal cells of each of the tested species were already (partly) damaged during the drying or knife-milling and in this way also organic cell constituents were removed. Algae drying, for example, may already lower cells' elasticity, as shown by Miranda et al. in a study with *Scenedesmus obliquus* [68].

In contrast to the applied procedure, algae were washed directly after harvesting in previous works [48, 55, 69]. In this way, cells were presumably still intact. For the presented feed-stocks, algae washing was not suitable as a pre-processing step and was not included in further calculations.

4.2.2 Algae milling

In the present work, grinding in liquid nitrogen (cryo-milling) was applied to avoid excessive heat (and destruction or reduction of desired products [70]), as well as effectively disrupt algal cells in a relatively short time [71]. This ensured a high accuracy during the experiments but is not suitable for large-scale application.

Algae cell walls often form strong barriers to solvents, which have to make physical contact with the lipids in order to successfully solvate them [5]. Cell wall architecture and composition and consequently the resistance to physical or chemical stresses among different algae and cyanobacteria are highly variable [8, 68, 70]. In addition, culture conditions can have a large impact on the biomass quality, as they do not only influence lipid synthesis and salt content but can also directly impact cell wall thickness and stability [72].

Crucial differences can be noted in the cell wall architecture of the applied species. AP is a cyanobacterium with a sixlayered wall. Its inner cell membrane is covered by a peptidoglycan layer, which is located between two fibrillary layers. The outer membrane is covered with a sheath of acidic polysaccharides and tightly connected to the peptidoglycan layer [70].

NG and SA both obtain cellulose layers and algaenancontaining layers. The latter are insoluble, nonhydrolysable, highly resistant aliphatic polymers of distinct biochemical origin. They are believed to be primarily responsible for the cell's robustness, but their composition varies between algal species [70, 73].

In particular, the plasma membrane of NG is connected to a thick, porous, cellulose-based layer via struts. It is protected by an algaenan layer with extensions that protrude from the surface. The *Scenedesmus* cell wall consists of an inner cellulosic layer that surrounds the membranes of individual cells and a thin, algaenan-based trilaminar sheath [74]. It is often described as extremely robust [68, 74, 75] and highly resistant to bacterial degradation [75], chemical attacks [68], and lytic enzymes [74].

PT exhibits major differences in composition, surface ultrastructure, and mechanical properties depending on morphotype [76, 77]. Ovoid cell walls, for example, are strongly silicificated and about five times stiffer than other types [76]. Martínez-Sanz et al. showed that *Spirulina* cells could simply be disrupted by stirring, whereas *Nannochloropsis* gaditana and *Scenedesmus almeriensis* required intense sonication [78]. Such a need for cell rupture was reported repeatedly during extractions of different components from NG, PT, and SA, such as pigments [79–83], amino acid concentrates [59] or lipids [81, 84].

It is commonly assumed that peptidoglycan-based cell walls, like in the case of AP, are softer than cellulosecontaining ones [85]. This relationship was confirmed in the present measurements. AP median particle size was reduced by 87% (Fig. 5 and Table 2), followed by SA (reduction of 76%), PT (54%), and NG (25%). A similar trend was observed for the PSD spectra (Fig. 5). NG and PT curves appeared rather unchanged with a maximum of particles close to the original value and some smaller fractions becoming more prevalent. In contrast, the SA and AP curves were completely shifted towards smaller particle sizes. The appearance of a higher number of particles below the size of a single cell is an indicator of cell debris or other cell constituents that were released by the mechanical force.

Weakening or rupture of cell walls also explains the enhanced extractability with DME (see Fig. 6). Micrographs of the feedstocks (Online Resource 5) additionally documented a noticeable increase in particle surface roughness and cell disintegration with higher pretreatment intensity.

Overall, the feedstocks can be classified with regard to their resistance to cryo-milling in the order NG > PT > SA > AP (from highest to lowest resistance) for the current experiments.

4.3 Lipid extraction

Chloroform/methanol(/water) systems, such as the commonly applied Bligh and Dyer method (1:2:0.8 chloroform/methanol/water with an initial solvent-to-sample ratio of 3:1 (v/m) [86]), are often recognized as "gold standard" [62]. The original Folch method works with a different ratio of 2:1 chloroform/methanol (v/v) and a solvent-to-sample ratio of 20:1 (v/m) [47]. It showed to be superior to the Bligh & Dyer in the determination of total lipid contents for samples of marine tissues with lipid contents above 2% [87]. In the current work, an increased solvent-to-sample ratio of 100:1 was applied in order to prevent solubility-limitation. This method was expected to provide a reliable value for total lipid contents of the different microalgae samples. Nonetheless, it is considered unsuitable for large scale industrial application, because of health and environmental risks associated with the use of chloroform [88]. It should also be noted that a complete lipid removal is usually only achieved by applying a high number of consecutive extractions (due to a lowered solvent mixture's capacity with an increasing amount of nonpolar lipids and a decreasing oil concentration in the starting material [5]). However, extracting the remaining lipids often requires a disproportionately high number of additional extraction steps [5] and would not add much value here.

As GRAS (generally recognized as safe) alternative, an ethanol-based reference extraction was tested (adapted from [89]), but not further considered due to low total oil recoveries (Online Resources 3, 4).

Previous studies reported the salvation of additional organic compounds (e.g., proteins or polyphenols) in DME [26, 90]. In order to mitigate these effects on yield determination, crude oil samples were solved in chloroform/methanol and processed similarly to the Soxhlet method (mixed with salt solution and consecutively separated in two phases) for the removal of unsaponifiable lipids and non-lipid material [91, 92]. Nonetheless, extracts were dark green to brown in color and thus obviously contaminated with algal pigments to a certain extent. Those pigments were most likely carotenoids, as chlorophylls tend to remain in the polar phase [54]. It was also noted that especially during the Soxhlet extractions, after solvent evaporation, a dark, brown to black residue was found, which was hardly soluble. Residual proteins that accumulated in the phase boundary have previously been reported [93, 94] and could be a part of this "insoluble fraction." Regarding the overall mass balance, the influence of pigments and residual proteins is expected to be small.

Overall, DME extracted 5–19% of total lipids without further pretreatment and 17–50% with additional cryo-milling (Fig. 6). In both scenarios, PT was the best performing species, followed by SA, NG, and AP. This did not correlate with the total lipid contents of the species. AP, NG, and PT showed very similar total oil contents, but AP and NG yields (knifemilled) with DME were at only 5–8% of total oil in contrast to 19% for PT. It is consequently assumed that the specific properties of the tested algae species were a major influential factor for the differences in observed extraction performances with DME.

4.3.1 Effects of milling on DME extractions

Cryo-milling reduced particle sizes drastically (Fig. 5 and Table 2) and did thus not only damage algae cells, but also enlarge the specific surface area of the samples [6]. Direct contact between solvent and solute and an enlarged contact area (shorter diffusion paths) can enhance overall extraction performance [95, 96]. Yap et al. described monophasic algae extraction systems as binary in nature with a fast lipid recovery from ruptured cells, followed by a slow, linear, diffusion-driven extraction from intact cells [18]. For each of the batch trials, by using long contact times with constant shaking, the formation of equilibrium between solvent and solute was allowed and kinetic limitation should be avoided.

However, it was still found that extraction yields were greatly enhanced after additionally cryo-milling the algae feedstocks (Fig. 6). This indicates that cell walls of the dried algae formed a strong barrier to DME. In particular, yields between knife- and cryo-milled samples of the DME extractions increased by factors of 2.0 < 2.3 < 2.6 < 3.5 for SA < NG < PT < AP.

Interestingly, changes in extraction yields after cryomilling do not match with the relative or absolute changes in particle sizes. As an example, SA was the 2nd most-affected species by milling and additionally exhibited the smallest absolute particle size after milling. Nevertheless, it showed the lowest increase in oil yield with the cryo-milling pretreatment.

In experiments with *Saccharomyces cerevisiae*, Spiden et al. demonstrated that PSD measurements were insufficient to quantify cell rupture, as they could not detect major flaws in cell walls (which might allow a release of metabolites) [97]. Also, in the present case, neither size reduction nor absolute particle size or a combination of both parameters could be directly linked to the degree of cell rupture and thus be utilized to predict extraction performances.

4.3.2 Study implications and limitations

To the authors' knowledge, most of the applied species have not been tested with DME so far. Hoshino et al. [34] previously worked with wet AP in a fixed-bed system, but focused on a simultaneous dewatering and oil extraction. Consequently, solvent amounts and extraction system were different from the given conditions.

Opposing to the presented results, similar extraction efficiencies between DME and alternative solvents were reported previously, including chloroform/methanol mixtures [29, 31]. With regard to hexane or $scCO_2$, DME was even found to excel those in several works with various biomasses [34, 98, 99].

Also, the displayed need for cryo-milling to maximize extraction performance is somewhat in contrast to literature [29–31]. As a recent example, Kanda et al. investigated the DME extraction of microalgae species with a rigid, bio-mineralized cell wall. They concluded that pretreatment was not needed to obtain satisfactory extraction yields with DME [33].

As previous studies differ in multiple major processing parameters, such as algae species, mode of extraction (fixedbed vs batch), solvent-to-sample ratio, and feedstock water contents, a direct comparison is highly problematic. The presented DME batch experiments were conducted to reveal species-specific differences between various microalgae rather than obtaining a maximum oil yield. They were performed with relatively low amounts of solvent and long contact time in order to ensure equilibrium between solvent and solute. Further optimization of the extraction was not carried out.

Previous works showed that solvent-to-sample ratio is a key factor for maximizing oil recoveries. DME consumption

was found to be positively correlated to lipid yields in both, fixed-bed [29, 30, 33, 100] and batch [101] extractions. As an example, Boonnoun et al. reported an increase in oil yield by a factor of 6 after changing the amount of DME from 6 to 200 g when extracting *Haematococcus pluvialis* [30]. Although it was demonstrated that DME extractions can work with ratios of 2.0–8.7 g DME g⁻¹ dry mass [99, 102, 103], during most fixed-bed extractions, solvent-to-sample ratios of up to 890 g g⁻¹ dry mass were applied [34]. This equals to about 500– 9000 g of solvent per g of lipid [25]. One of the few batch studies, which showed good results with regard to the reference treatment (hexane), used a solvent-to-sample ratio of roughly 160 g g⁻¹ dry mass for maximum extraction of *Euglena gracilis* in comparison to 10 ± 1.5 g g⁻¹ here [101].

In the current setup, a limited solubility (due to low solvent-to-sample ratios) might explain the lower overall extraction efficiency with cryo-milled samples. The solubility of lipids is owed to a number of interacting factors, such as oil composition and lipids arrangement, which strongly vary with algae species. Nevertheless, this explanation cannot cover the pretreatment-induced differences between knife- and cryo-milled algae. In the latter case, a similar quantity of DME solved more than double the amount of oil. As an example, for PT the amount of DME per lipids extracted changed from 565 g g⁻¹ for knife-milled samples to 210 g g⁻¹ after cryo-milling.

In this context, the downsides of high amounts of solvent should also be touched briefly. Energy requirements for pumping and DME recovery increases with the amount of solvent applied. Moreover, if 5% of DME is lost during an extraction cycle as stated by Oshita et al. [104], it adds up to 10–450 g of DME lost per g oil extracted for the reported 200–9000 g of solvent per g of lipid (literature and own values). Especially for biodiesel-focused approaches (e.g., [33, 34, 105]) such ratios may raise questions about sustainability, as DME is regarded as alternative fuel itself [106].

As another major influential factor, the initial water contents of feedstocks must be taken into account. Algae with roughly 74–95% water were used in previous studies [24, 30, 32, 100, 101]. This is advantageous with regard to a final application [19, 20], but water is expected to have a promoting or at least influencing effect on the overall extraction. On the one hand, it is partly solved by DME and is suspected to change the solvent mixture's polarity [25]. On the other hand, due to the mutual solubility, cell wall penetration, and consecutive salvation of lipid compounds might be facilitated. In addition, water transporting mechanisms of living cells could also affect the extraction in comparison to dried feedstocks.

In experiments with the brown seaweed Undaria pinnatifida, Billakanti et al. reported a decrease of DME yields

by about 6% of total lipids when comparing wet to freezedried samples [60]. Although this is a rather small difference compared to the presented results, the possibility of water being a main reason for the low extraction efficiencies cannot be ruled out.

As discussed, the conducted experiments differ from previous studies in multiple key parameters. It is consequently not recommended to use the present results as a basis of comparison for DME extraction efficiency.

4.4 Fatty acid compositions

Each of the tested species showed a unique fatty acid profile (Table 3). Regarding the results of total lipid extraction, AP and SA results were well in accordance with literature [36, 107, 108].

For PT, a predominance of C16:1 and C16:0 and a lack in C18 fatty acids were found. Also, a high share of EPA was confirmed in this species, which may vary with culture age and conditions [42, 43, 109, 110].

The measured fatty acid profile of NG strongly differs from other works. A composition of mainly C16:0 and C16:1 with small amounts of C14:0 and C18:1n9 and around 1% of C18:3n3 was reported [63, 111]. The latter, however, was dominant in this study with a share of 42%. Also, in literature, EPA yielded between 2.5 and 14% (for N starved and replete cultures) [63] in contrast to 0.4% here. It is consequently suspected that subspecies and/or culture conditions of the extracted NG were fundamentally different or that the culture was heavily contaminated.

In general, saturated and monounsaturated fatty acids with 14–20 C–atom chains are preferred for biodiesel applications [38, 112–114]. AP and PT oils contained the highest amounts of C16:0 and C16:1 and also small shares of C18:1n9. With combined ratios of saturated and monounsaturated fatty acids of 73% and 81%, respectively, those oils could be suitable as biodiesel feedstock.

As another major point of interest, PUFA, such as linoleic acid (C18:2n6), α -linolenic acid (C18:3n3), eicosapentaenoic acid (EPA, C20:5n3), and docosahexaenoic acid (DHA, C22:6n3) were examined.

Total amounts of PUFA among the tested feedstocks were highest in NG (56%), followed by SA (55%), AP (27%), and PT (19%).

PUFA can be found in various organisms as parts of membranes or storage organelles or as precursors for many biologically active eicosanoids—mainly in the form of glycolipids, phospholipids, sphingolipids, and lipoproteins [42, 115]. For photosynthetic organisms in standard conditions, phospholipids and glycolipids are mostly present in the endomembrane system and the plastid, respectively [110].

PUFA are entirely derived from the diet and essential for human health. A ω -6 to ω -3 fatty acid ratio of \leq 5:1 is generally recommended by health officials, which is often not met by typical western diets [116]. PUFA-rich products can help to balance out nutrient ratios or even for the treatment of illnesses and mental disorders [116, 117].

Of the named PUFA, especially C18:3n3 was very abundant in NG and SA oils with about 42% of total oil. C18:2n6 made up between 12 and 14% for AP, NG, and SA. However, both fatty acids can also be found in many vegetable oils, as well as in nuts, eggs, dairy, and meat [118].

As "more interesting" fatty acids, GLA, DHA, and EPA were examined. GLA is incorporated into infant formula and used to treat several diseases, e.g., atopic eczema [119]. DHA and EPA are included in therapies in order to prevent dementia [42]. EPA has further shown effectiveness in the treatment or prevention of cardiovascular diseases, high blood pressure, different forms of cancer, and schizophrenia, among others [42, 120, 121].

DHA was not present in relevant amounts and only AP showed meaningful GLA quantities around 13%. EPA was very abundant in PT oil with around 16%.

Although total PUFA quantities were high for NG and SA, AP and PT seemed to be more suitable biorefinery candidates. They showed high levels of lipids usable for biodiesel conversion, as well as a substantial portion of valuable GLA and EPA, respectively. Furthermore, they have the potential to provide other high-value side products, such as phycocyanin (AP) [122] or fucoxanthin (PT) [81].

Concerning DME extraction applications, PT is recommended for further trials. SA also showed good overall yields, but a less favorable lipid profile and AP and NG demonstrated to be least compatible with DME.

4.4.1 Effects of extraction procedure on fatty acid compositions

It was previously hypothesized that DME preferably extracts saturated and monounsaturated fatty acids, such as C16:0 and C16:1 [25] suitable for biodiesel purposes.

With regard to the total relative amounts of saturated and monounsaturated fatty acids, a preferred extraction of DME could not be verified. However, in most cases, the DME extracted oil was characterized by higher shares of C14:0, C16:0, and C16:1 fatty acids. A preferential extraction of C14:0, C16:0, and C18:0 species with DME in comparison to Bligh and Dyer, hexane, and scCO₂ was also previously reported by Hoshino et al during the extraction of *Aurantiochytrium limacinum* [31].

At the same time, it also became apparent that the used algae species interacted differently with the applied solvents and thereby influenced oil compositions. As an example, during the treatment of AP, the relative C16:0 amount was roughly 10% higher when DME was applied instead of chloroform/methanol. For SA, on the contrary, the relative C16:0 content changed from about 14% with DME to about 22% during Soxhlet.

Solvents (and procedures) showed to have an effect on the oil composition in the current work. The properties of the applied algae species and the specific interaction with the solvents seemed to be another determining factor for relative fatty acid contents in the resulting oils.

4.4.2 Effects of milling on fatty acid compositions

Cryo-milling increased the share of total PUFA during DME extraction for each species. Overall, relative PUFA recoveries were increased between 1.1% (SA) and 7.9% (PT) by cryo-milling. The most remarkable increase was observed for the relative EPA content in PT and accounted for roughly 7%.

According to literature, EPA is typically found in all membrane lipids of this species [110]. Since most PUFA are integrated as structural parts of membranes and organelles [42], these results are a strong indicator for advanced cell disruption and facilitated access of the solvent to inner cell compounds and building blocks.

As the intensive cryo-milling pretreatment led to increases in relative PUFA yields and changed the properties of the produced oils, no evidence was found that pressurized DME could compensate for the pretreatment in the current setup.

5 Conclusion

The present work showed that liquefied dimethyl ether can be a valuable solvent for biorefineries. Although working with low amounts of solvent in an unoptimized one-stage batch extraction, up to 50% of total oil could be recovered. DME extracted oils were characterized by increased relative proportions of C14:0, C16:0, and C16:1 fatty acids.

Phaeodactylum tricornutum proved to be most suitable with DME concerning extraction yields and content of valuable fatty acids in the current setup. Its oil was characterized by a high content of EPA in combination with biodieselrelevant fatty acids. Yields of *Nannochloropsis gaditana* and *Arthrospira platensis* were significantly lower and *Scenedesmus almeriensis* showed good recoveries but a less favorable fatty acid profile.

The applied cryo-milling pretreatment reduced particle sizes and simultaneously enhanced the DME yields significantly by factors of 2–3.5 for each species. It also helped to increase the ratios of PUFA in DME extracted oils. Hence, for the tested extractions, an intense cell-disrupting pretreatment was needed in order to optimize DME performance. It is assumed that algae species and water content strongly influenced this outcome.

As each microalgae feedstock exhibited specific morphological traits that interacted differently with pretreatments and solvents, the authors are convinced that there is no general answer to the question which algae species or solvent is "the best." Future extraction systems will have to be tested and adjusted with regard to biomass properties and desired products. However, the easy oil separation and dewatering ability are expected to make DME a versatile contender for future lipid extraction applications.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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