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***Desmodesmus* sp. pigment and FAME profiles under different illuminations and nitrogen status**

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Abstract

Acknowledging the limited knowledge available on microalgae *Desmodesmus sp.*, a specimen isolated from nearby freshwater lake was studied. Photobioreactor runs were led at two different illuminations (moderate - 150 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ - and high - 300 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$) and under three nitrogen status: sufficient, deficient and starved. Pigment and FAME profilings were carried out. *Desmodesmus sp.* has a Chlorophyceae common pigment profile with natural occurrence of lutein. Nitrogen deficiency induced both chlorophyll *a* and *b* contents decrease while not modifying the overall profile. Long term starvation induced pigment profile modification: disappearance of violaxanthin and stronger expression of carotenoids such as canthaxantin. FAME profiling showed that the three major fatty acids produced by this strain are oleic (7.0 %g/gDW), linoleic (5.0 %g/gDW) and γ -linolenic (6.9 %g/gDW) acids. The first two being the storage lipids of this strain. Those two fatty acids have valuable health properties and may pave the way toward nutraceuticals covalorization.

Keywords

Desmodesmus sp., Covalorisation, Pigment, FAME, Profiling

1. Introduction

Microalgae started to draw scientists and engineers attention about 50 years ago for many reasons. The most striking ones being: photosynthesis, robustness and fast growth. In more details, while growing, microalgae can derive their energy from sunlight and fix inorganic carbon. Furthermore, compared with higher plants they show superior photosynthetic efficiencies 4-5 % (compared with 1-2 %) (Walker 2009), no intrinsic sensibility to seasonality and faster growth. Another

edge of microalgae is that their culture can be deployed on non-agricultural land, without pesticides and using wastewater. Thanks to all this, it is nowadays commonly admitted that microalgae can be of help in facing the numerous challenges mankind is confronted with. Among them climate change, induced by
45 fossil carbon release, may be the most urging one. As part of the solution, microalgae can sequester atmospheric carbon dioxide, mitigate its emissions, and be a substrate of choice for third generation biofuel (Sati et al. 2019).

Still, after years of research, the scientific community acknowledged the fact that the sole production of biofuel is not viable economically (Rizwan et al. 2018).
50 Indeed, even in the most optimistic scenarios, whole biomass cost is about 0.82 €/kg dry weight (Acién Fernández et al. 2019). Thus biofuel production will only be possible as part of a co-valorization process of microalgal biomass. Several strategies can be envisioned: growing biomass on formulated wastewater (alleviating 0.20 €/kg dry weight (Acién Fernández et al. 2019)) or extracting
55 high added-value molecules before turning spent biomass into biofuel.

In addition to this paradigm shift, strain selection and optimization will also be key (Lim et al. 2017), as it would ease the lifting of the three bottlenecks of microalgae large scale cultivation: dense culture (as cultures only contain only 1 - 20 g/l) (Enamala et al. 2018), harvest (Fasaei et al. 2018), extraction (Rizwan et al. 2018).
60 Luckily only a fraction of the microalgae species have been discovered (Guiry 2012), and even fewer studied. That is why, with this in mind, researchers have led environmental surveys and strain characterization - the initiative of the National Alliance for Advanced Biofuels and Bioproducts being of note. In their work, Neofotis et al. pointed out that, even though widespread, *Desmodesmus* genus is
65 far from well-known and may hide potential for biofuels and bioactive molecules production (Neofotis et al. 2016). This can be confirmed by a simple analysis of Web of Science database. In the date of the 28th of January 2020, topic search for
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Chlorella genus yielded 13732 articles, *Scenedesmus* 4979 and *Desmodesmus* only 417. Thus, the present work aims at gaining general knowledge about

70 *Desmodesmus* genus when it comes to its growth rate, medium affinity, pigment and Fatty Acids Methyl Ester (FAME) profiles.

To do so, a *Desmodesmus sp.* strain was isolated from nearby lake. Indeed, obtaining strains from closeby ponds ensures that they are already adapted to local light and temperatures conditions. Hence it would ease local deployment in case

75 of promising discovery. After collection, this strain response to varying growth media was screened. Then, its pigment and FAME profiles were extracted. This was done under different growth conditions as microalgae are known to be able to adapt such profiles depending on environmental conditions (Ratha et al. 2012).

80 Nitrogen limitation or starvation are techniques frequently applied to microalgae to manipulate their secondary metabolites profiles. For example, Pancha et al. demonstrated that nitrogen limitation leads to a decrease of photosynthetic activity while triggering highest lipid and carotenoid production (Pancha et al. 2014).

Similar findings have been reported by many other authors, thus those can now be considered as routine culture protocols (Lari et al. 2016; Mandal et al. 2009;

85 González-Garcinuño et al. 2014; Ratha et al. 2012). Lighting strategies have also be shown to impact microalgal growth. In this regard, the pioneering work of Myers and Kok has to be acknowledged (Kok 1953; Phillips et al. 1954). More recently, excess light has been shown to modify both lipid and pigment profiles (Leonardi et al. 2019; Gris et al. 2014; Sarat Chandra et al. 2017; Sforza et al.

90 2014). Carotenoids and other xanthophylls pigments are expressed as a means to protect the cell from upset photosynthesis induced Reactive Oxygen Species (ROS) (Müller et al. 2001; Choudhury et al. 2001). Thus, after determining the most promising growth medium, *Desmodesmus sp.* pigment and FAME profiles

were determined under nitrogen sufficient, nitrogen deficient as well as moderate
95 and high light conditions.

2. Materials and methods

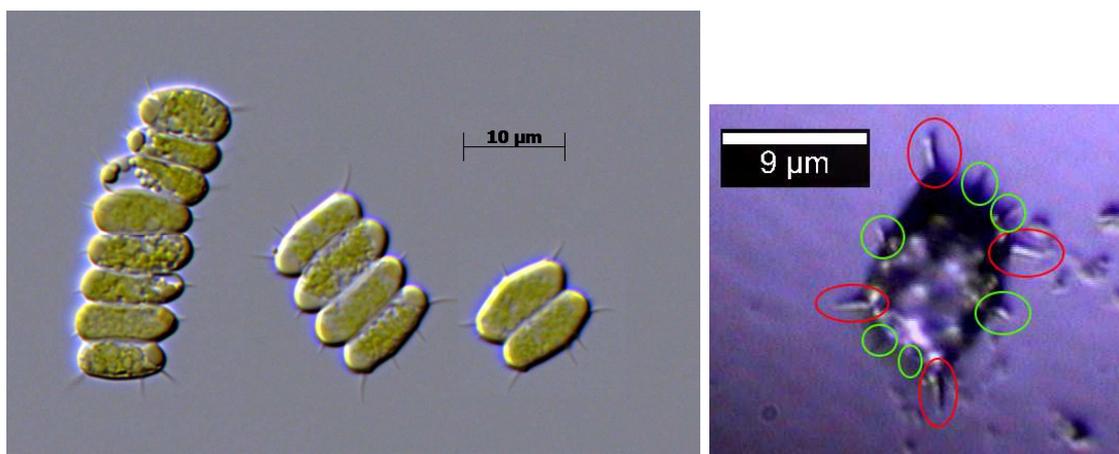
2.1 Strain

The strain was obtained by sampling in late Autumn 2018 from freshwater
Bazancourt lake (GPS: 49.365412, 4.162591), France. Microscope examination
100 confirmed the presence of *Chlorella sp.*, *Scenedesmus sp.*, *Desmodesmus sp.*,
Dunaliella sp., *Euglenozoa sp.*, several protozoa and bacteria. The obtained
consortium was cultured over BG-11 (Blue-Green medium with 11 elements)
medium in 250 ml flasks (50 ml culture medium). Then Petri dish isolation was
conducted. From the obtained strains, *Desmodesmus sp.* was latter purified and
105 kept axenic.

With the help of Hegewald pioneering work (Hegewald et al. 2005), the strain was
further characterized. Two key factors were used: population distribution and
microscopy similarities. This strain population exhibits 2, 4 and 8-cell colonies
(Fig. 1, right). Even though cells were extensively observed, no singled cell
110 colonies nor 16-cell colonies were found. The most common ones being the 4-cell
colonies, when no stress, nor synchronization is applied (Ševčíková et al. 2013).
All of the colonies featured distinguishable long spines, while short spines could
not be identified with certainty using benchtop optical microscope. Thus
additional observations were carried out with a confocal microscope (Alpha 300
115 R+ Witec) in order to obtain observations whose focal plan matches the spines
position. Those observations revealed the presence of short spines in addition to
the long ones (Fig. 1, left). The fact that the strain possesses both short and long
spines points towards *Desmodesmus* genus while the types of observed colonies

narrow it down to *pleiomorphus* species. Still, not being taxonomists, the authors
120 would remain cautious and referred to the strain as *Desmodesmus sp.*.

The generation process of this genus is similar to other colony living genus, such
as *Scenedesmus*. Generation takes place in an independent manner within each cell
composing the colony. When a cell enter the autosporeulation process it will form
either a 2, 4 or 8-cell colony, depending on the environmental conditions and not
125 on the number of cells present in its current colony. Then, the newly form colony
departs from its former host, leaving an empty spot. Finally, manipulation of the
light photoperiod allows to select which type of colonies are expressed by the
strain (Ševčíková et al. 2013).



130 *Fig. 1: Visual observations of the Desmodesmus sp. strain used for this study. Sampled
from a subculture Right: Microscope observations, magnification x100 (image
manipulation: contrast enhancement). Left: Confocal observations. Circled in red: long
spines. Circled in green: short spines*

2.2 Subculturing

135 The first action was to identify optimal growth medium for this strain. To do so,
Fraquil, WC (Wright's Cryptophyte), BBM (Bold's Basal Medium), B3N (BBM
with 3 times more nitrates) and BG-11 media were screened. Those media were
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selected as they are indicated for *Scenedesmus* genus growth in Andersen media compendium (Andersen et al. 2005). The second criterium of choice was the range of nitrogen contents those media offered (from 0.10 mM for Fraquil to 17.6 mM for BG-11)

In order not to bias the screening by former BG-11 subculturing, the strain was subcultured on each medium for three weeks. Three passagings consisted in a 100-fold dilution to 50 ml of culture medium in 250 ml flasks. Illumination was set as 75 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ and temperature at 20 °C. They were all conducted in triplicates. Fraquil medium showed sign of early exhaustion and was therefore excluded from the screening. Cell concentration was tracked after a fourth passaging, in triplicate again. The monitoring was realized twice daily, optical density at 750 nm (OD_{750} hereafter) was used as proxy of cell concentration (more details are available in Sec. 2.4).

Media screening are detailed in Section 3.1. Briefly, B3N was the most suited medium and was therefore used for subculturing during the remaining of this study. Subculturing took place every Monday morning, with unchanged passaging procedure: 1/100 sampling, 250 ml flasks, 50 ml culture medium, 75 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$, 20 °C, duplicate. Once per month, culture plated on glucose enriched Lysogeny Broth (LB, promoting bacteria proliferation) to check for potential mother culture contamination. None were found.

2.3 Culture protocol

The culture was fed in a 2.5-liter bubble column photobioreactor. Such a large vessel was used for two reasons. First of all, it is closer to industrial scale culture protocols than large capacity shake flasks. Second, it allows for large sampling volumes, meaning that, for every sample, enough biomass is collected to later process it and acquire both pigment and FAME profiles. External lighting was

ensured by cool white LED ribbons (OSRAM GW JDSRS1.EC) placed all around
165 the photobioreactor. LEDs were closely packed together across all the
photobioreactor height to ensure uniform illumination. Lighting capacity ranges
from 30 to 300 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ at the surface of the culture. In addition, as
it is known that light spectrum influences microalgae (Mohsenpour et al. 2012),
light spectra were acquired across this range (Ocean Optics Flame-T-UV-Vis
170 portable spectrophotometer). Light spectrum exhibits no variation over the
aforementioned range. Finally, culture temperature is controlled using a heater-
chiller directly injecting water into the thermal jacket.

The cultures were led with 2 liters of B3N medium. Aeration was set at 0.2 vvm,
with 2.5 % CO_2 -enriched air. Culture temperature was set at 20 °C. Two different
175 illuminations were tested 150 and 300 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$. The first one
represents moderate illumination, at least for the resembling *Scenedesmus* genus
(Liu et al. 2012; Přibyl et al. 2016) while the second should trigger light stress.
Inoculation was realized after sterilization by injecting 50 ml of B3N subcultured
cells passaged 10 days before. This delay was chosen to ensure that cells were in
180 the exponential phase at the time of inoculation. At the end of each run, the
remaining culture was removed and the photobioreactor cleaned and sterilized.

Finally, long term nitrate starvation effects were also explored. Sadly, despite the
fact that a liter size capacity photobioreactor was used in this study, this volume
was still too low to allow for one month or more follow-up (with at least 120 ml
185 withdrawn each day). Thus shake flasks subcultures, without continuous sampling,
were used for this last part of our investigation. To do so, subcultures were left
under incubation conditions (75 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$, 20 °C, air). They were
analyzed one month after passaging.

2.4 Cell concentration monitoring

190 Samples were drawn out of the culture vessel twice daily, 1 ml for shake flasks, 50
ml for the photobioreactor (discarding the 10 first milliliters). Cells growth was
monitored using optical density (Shimadzu UV-1800 spectrophotometer) as proxy
of cells dry weight. Samples were diluted to an optical density of 0.4 or less before
recording the value. Following Griffiths' advices (Griffiths et al. 2011), the optical
195 density was acquired at 750 nm, as it allows to almost only account for cell walls
and not pigments, hence measuring only for biomass. Once more, following his
recommendations, a calibration curve linking optical density and dry weight was
produced using late exponential phase cells (7 points, range: 0.02 to 0.5, $R^2 =$
0.997). With this curve it is possible to compute biomass density from optical
200 density readings. Finally, offline pH measurements and frequent microscope
observations were done to check the cells condition.

2.5 Nitrate concentration monitoring

As nitrogen is known to have a dramatic impact on cell conditions, nitrate
concentration was monitored until exhaustion. Samples were prepared by
205 centrifugation (11000 rpm, 4 °C, 10 min, Eppendorf 5804 R). Supernatant was
then filtered (0.22 µm polypropylene) before being diluted with milliQ water
(Integral-5, Merck, also analyzed for correction) in order to be in the range of 0.2
to 10 mg/l. Nitrate quantification was carried out on an ICS-5000+ Ion
Chromatography system (Thermo Fisher Scientific) coupled with a conductivity
210 detector. Separation was achieved on an AS11-HC column (2x250 mm, 4 µm)
protected by a guard column AG11-HC (2x50 mm, 4 µm). Column temperature
was maintained at 35 °C. The mobile phase of 30 mM of potassium hydroxide was
at a flow rate of 0.2 mL/min in isocratic mode and was delivered by an EGC 500
eluent generator. The eluent generated was then purified by a CR-TC to trap

215 impurities such as carbonate. The suppressor was an ADRS 600, 4 mm, operated
in recycle mode at 15 mA. The injection volume was 2 μL and total run analysis
lasted 30 minutes. Nitrate was identified by comparison of its retention time with
standard solutions. Quantification was achieved using the height of the peak in
external calibration (0.2 to 10 mg/l range). Nitrate standard was purchased from
220 Sigma-Aldrich with a TraceCert quality which is a standard at 1000 ± 4 mg/l.

2.6 FAME profile determination

Transesterification done using an in-situ transesterification following NREL
protocol (Van Wychen et al. 2013). Quantification of the FAME was carried out
on a GC-2010 Plus coupled to a GCMS-TQ8040 triple quadrupole mass
225 spectrometer (Shimadzu). Separation was achieved on a HP-88 capillary column,
100m x 0.25 mm x 0.20 μm (Agilent). Helium (99.9999 %) was used as the carrier
gas with a column flow rate of 2 mL/min, the linear velocity was kept constant
during the analysis. The injection volume and temperature were 1 μL and 240 $^{\circ}\text{C}$,
samples were injected in splitless mode. The oven temperature started at 50 $^{\circ}\text{C}$ for
230 1 minute and raised to 180 $^{\circ}\text{C}$ with a rate of 20 $^{\circ}\text{C}/\text{min}$, held for 5 minutes. It was
successively raised with a rate of 10 $^{\circ}\text{C}$ at 200, 210, 220 and 230 $^{\circ}\text{C}$ and
respectively for 5,5,1 and 6 minutes. Total run time was 34.5 minutes. Detection
was done by the triple quadrupole in Electron Ionisation (EI) with an energy of 70
eV. The temperature of the ion source was set at 200 $^{\circ}\text{C}$ and the interface at 210
235 $^{\circ}\text{C}$. The acquisition was done in MRM mode (Multiple Reaction Mode),
transitions and energy of collision for each FAME are supplied in the
supplemental material section. Compounds were identified by comparison with
their retention time with standards, for each compound 3 different transitions were
recorded, one was used for quantification using the area of the chromatographic
240 peak and the two others were used as qualifiers. The ratio of intensity between
quantifiers and qualifiers was also used for additional identification. Internal
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calibration was done using C13:0 as the internal standard. FAME standard was the Supelco 37 component FAME mix, CRM47885 from Sigma-Aldrich. All samples were also acquired in scan mode using the third quadrupole from 50 to 400 m/z in
245 order to visualize any unquantified compound.

2.7 Pigment profile determination

Samples were prepared by centrifugation of 50 ml of culture (4 °C, 11000 rpm, 10 min). Supernatant was used for nitrate quantification. The pellet was frozen, then freeze-dried (1 day primary drying, 2 days secondary drying, Christ alpha 1-2 LD
250 +). 1 mg of dried microalgae powder was homogenized in 5 ml pure methanol using MP Biomedicals FastPrep42 bead miller. Liquid was then filtered (0.22 µm) and stored in dark vials at 4 °C before quantification.

Quantification of pigments were carried out on an Ultima 3000 HPLC (Thermo Fisher Scientific) coupled with an UV Detector. Separation was achieved on an
255 Acclaim Polar Advantage II C18 column (4.6 x 150 mm, 3 µm, 120 Å) from Thermo Fisher Scientific. Column temperature was maintained at 30 °C. Pure methanol was the mobile phase, the flow rate was 0.5 mL/min and the elution set in isocratic mode. Injection volume was 5 µL and the total run analysis was 40 minutes. Compounds were identified by comparison of their retention time and of
260 their UV-Vis spectra with standard solutions. UV-Vis spectra were recorded from 200 nm to 700 nm, absorbance was recorded at 400, 450, 500 and 650 nm.

Pigments quantifications were done at 650 nm. It was achieved using the area of the peaks in external calibration, which concentrations ranged from 0.25 to 5 mg/l. Pigment standards and methanol were purchased from Sigma-Aldrich. Standards
265 had a purity greater than 97 %.

2.8 PAM measurement

In order to assess for the state of the photosynthetic apparatuses, chlorophyll fluorometry analyses were carried out during the exponential phase under nitrogen sufficient conditions. Experiments were led using pulse modulated
270 spectrophotometer (Dual-PAM-100, Walz). Two types of measurement were conducted. First Fv/Fm ratio was analysed, as it is an efficient proxy of photodamage. Even though fast and informative, Fv/Fm ratio can be seen as a too simplistic description of photosynthetic apparatuses. Indeed, for this measurement cells are dark adapted and exposed to only one flash. Hence this specific measure
275 does not yield information on actinic light management under different illuminations. Still, it would detect severe damage done to the photosynthetic apparatuses.

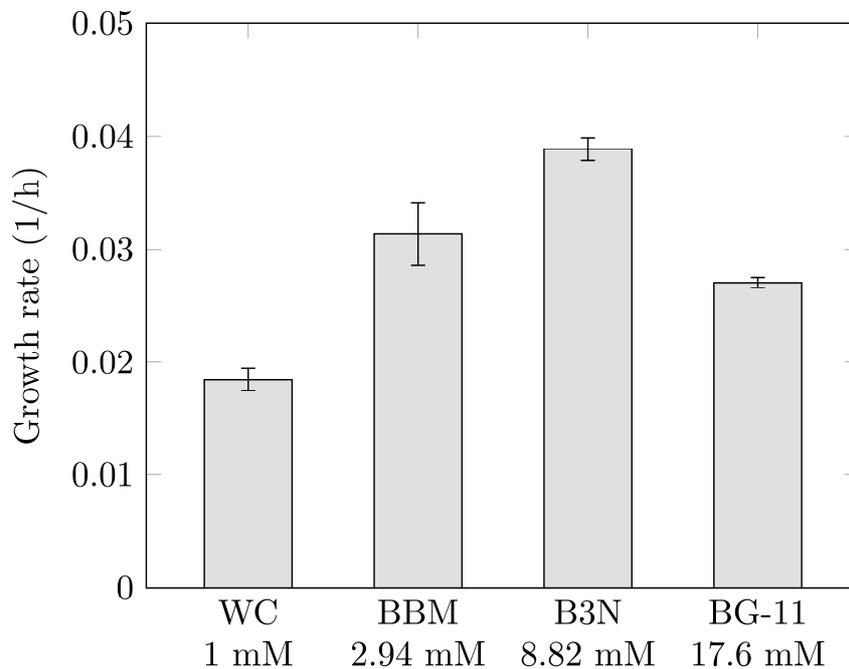
Thus, to avoid this shortfall, photosystem II yield (Y(II)) and NPQ yield (Y(NPQ)) were also recorded for 3 different actinic light intensities: 126
280 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$, to mimic the first photobioreactor conditions (150), 339, to mimic the second condition (300) and 1028 to investigate the cell reaction to excess light. Y(II) indicates the fraction of energy that is processed by the non-cyclic photosynthesis scheme, hence driven towards carbon fixing. Y(NPQ) quantifies the fraction of incident light energy that is safely regulated via
285 xanthophyll cycle.

Each measurement point was triplicated.

3. Results

3.1 Media screening

Media screening results are presented in Figure 2. As one can see, standard deviations are very low and results are statistically significantly different from medium to medium. The highest growth rate was obtained with B3N ($0.0389 \pm 0.001 \text{ h}^{-1}$) followed by BBM ($0.0313 \pm 0.003 \text{ h}^{-1}$), BG-11 ($0.0271 \pm 0.0004 \text{ h}^{-1}$) and WC ($0.0184 \pm 0.001 \text{ h}^{-1}$). This observation calls for a comment on strain affinity towards nitrogen. Indeed, WC, BBM, B3N and BG-11 show increasing nitrogen contents, and relatively similar composition for the last three of them. This means that the 3-fold increase in nitrogen content from BBM to B3N is favourable to this strain. On the other side, the same 3-fold increase from B3N to BG-11 exhibits detrimental effect. Thus, an optimal nitrogen concentration exists, probably not too far from B3N concentration.



300

Fig. 2: Growth rates resulting from the media screening. First abscissa line: media names, second abscissa line: nitrogen content. Error bars: standard deviation (n=3)

3.2 Cell growth, substrate monitoring and photosynthetic status

- 305 Shake flask and photobioreactor runs unfolded as expected. Repeatability was very good for moderate illumination runs (growth rates of 0.0460 and 0.0463 h⁻¹, exponential phase log fit R² > 0.97 for both runs) and reasonably for high illumination ones (growth rate of 0.0174 and 0.0153 h⁻¹, R² > 0.98 for both runs). Thus, only one run will presented and commented hereinafter.
- 310 Figure 3 presents the follow up of the first 150 μmolPhotonPAR/m²/s photobioreactor run. Culture started with a lag phase (0 to 72 h), followed by an exponential phase (72 to 129 h), a linear phase (129 to 241 h) and a stationary phase. Nitrate concentration exhibited a classical behaviour: almost plateau during the lag phase, with a value close to the one of the virgin medium, then a sharp
- 315 decrease during the exponential phase. One should note that nitrate exhaustion and exponential phase end took place at the same time. Finally, over the culture, pH rose from 6.3 to 7.6 at the end of the run.

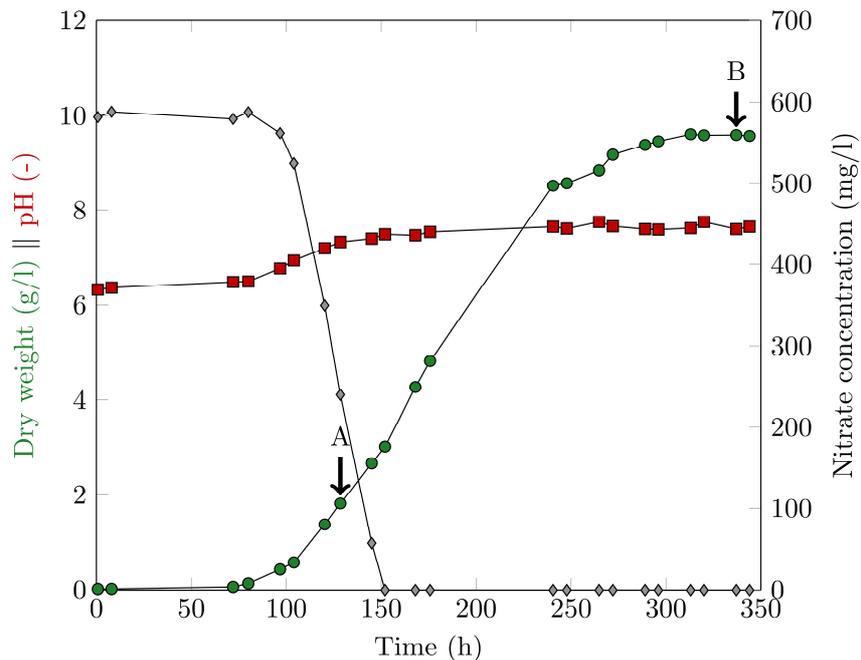
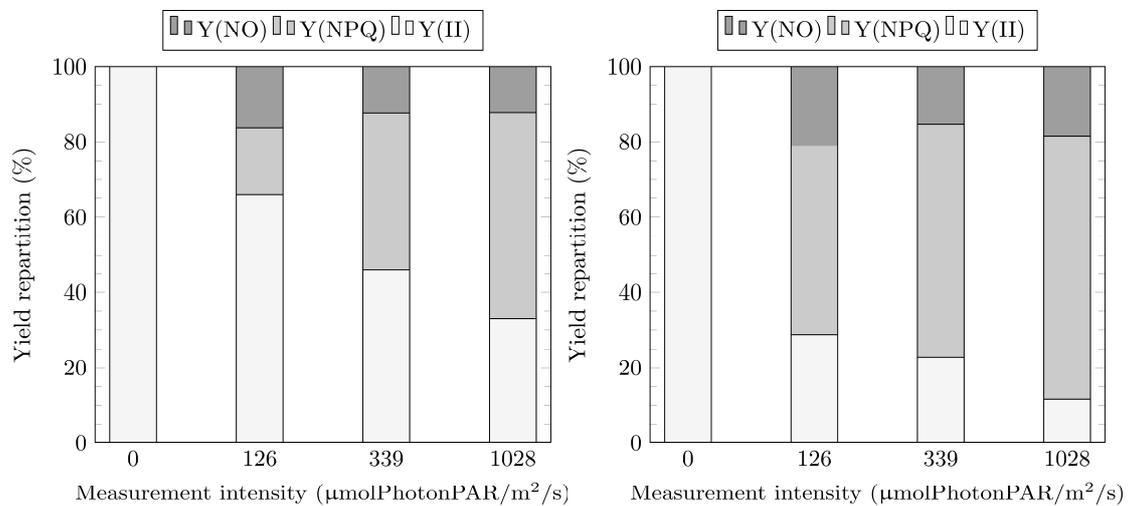


Fig. 3: 150 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ photobioreactor run monitoring. Green markers: biomass concentration. Red markers: pH. Grey markers: nitrate concentration. A mark: nitrogen sufficient sample, B mark: nitrogen deficient sample

Thanks to those curves, it was possible to determine the samples produced in nitrogen sufficient conditions and those produced under nitrogen depletion. Nitrogen sufficient samples were taken in exponential phase, while nitrogen was still available. Nitrogen limited samples were taken in late exponential (early stationary) phase. Once identified those samples were used for pigment and FAME profiling. Thus 5 profiles have been established: 150 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ nitrogen sufficient and deficient, 300 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ nitrogen sufficient and deficient and 75 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ nitrogen starved for 1 month.

Finally, chlorophyll fluorometry results are available in Figure 4. They are displayed as the energy flux repartition: Y(II) energy flux forwarded from PSII to PSI, Y(NPQ) safely regulated energy excess, Y(NO) the remaining energy *i.e.*

335 unregulated energy excess ($Y(NO) = 1 - Y(II) - Y(NPQ)$). They were obtained during the exponential phase under nitrogen sufficient conditions on the replicate runs. In addition, Fv/Fm ratio was 0.778 ± 0.038 for the culture under $150 \mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ and 0.570 ± 0.059 for the one under $300 \mu\text{molPhotonPAR}/\text{m}^2/\text{s}$.



340

Fig. 4: PAM measurements for both lighting conditions (left: $150 \mu\text{molPhotonPAR}/\text{m}^2/\text{s}$, right: $300 \mu\text{molPhotonPAR}/\text{m}^2/\text{s}$). Samples taken in exponential phase, nitrogen sufficient part. Standard deviation ($n=3$) below 0.06

3.3 Pigment profiles

345 First of all, pigment profiles were very repeatable from one run to the other. $150 \mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ and $300 \mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ photobioreactor yielded similar profiles, with different relative amounts of pigment species. Furthermore, the two growth conditions exhibited the same trend when going from nitrogen sufficient to nitrogen deficient phase. Over 1-month starvation, the culture turned

350 from green to orange/red color. From these profiles, four pigments can be identified with certainty in most of the conditions: chlorophyll *a*, chlorophyll *b*, lutein and violaxanthin. Other pigments, such as asthaxanthin and canthaxanthin have been detected in some samples, still their contents were too low to be quantified or even identified with certainty in all samples. Finally, the four main
 355 pigment contents have been quantified and are reported in Table 1

Table 1: Major pigments quantification results for the five growth conditions. BQL: Below Quantification Limit. LoQ: Limit of Quantification. LoD: Limit of Detection. Displayed as value \pm measurement uncertainty

Conditions		Pigment content (%g/gDW)			
		Chl <i>a</i>	Chl <i>b</i>	Lutein	Violaxanthine
150 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$	N sufficient	1.498 \pm 0.016	0.694 \pm 0.035	0.251 \pm 0.005	0.113 \pm 0.002
	N deficient	0.187 \pm 0.003	0.131 \pm 0.004	0.148 \pm 0.003	< LoQ
300 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$	N sufficient	1.427 \pm 0.014	0.638 \pm 0.032	0.244 \pm 0.005	0.097 \pm 0.002
	N deficient	0.838 \pm 0.009	0.432 \pm 0.018	0.185 \pm 0.004	0.076 \pm 0.001
1 month old subculture		0.076 \pm 0.000	0.031 \pm 0.001	0.039 \pm 0.000	< LoD

360 3.4 FAME profiles

Like for the pigments, FAME profiling is very repeatable. Still, while the detected species list does not evolve much, relative amounts of fatty acids vary over the tested conditions. Quantification of the seven major fatty acids is presented in Table 2. As one can see, going from nitrogen sufficient to nitrogen deficient
 365 condition triggered oleic acid accumulation. This trend is exacerbated by nitrogen starvation. Furthermore, nitrogen starvation also promotes storage of palmitic acid. In addition to nitrogen status manipulation, light stress induction also had an

impact on the strain FAME profile. Going from 150 to 300 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ reduced by a factor almost two the quantities of oleic and linoleic acids. Finally, total fatty acid content is also computed on the last row of the table. As one can see 1 month nitrogen starvation allowed the strain to accumulate a twofold quantity fatty acids compared with photobioreactor runs. This behavior, which mainly supported through palmitic and oleic acids production, is thought to be the origin of the appearance of the culture orange/red color.

375 **4. Discussion**

4.1 Light stress assessment

The first step was to ensure that 300 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ illumination could actually be considered as a light stress condition while 150 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ illumination should not be.

380 Table 2: Major fatty acids quantification results for the five growth conditions. LoD: Limit of Detection. Displayed as value \pm measurement uncertainty

Conditions	150 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$		300 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$		1 month old subculture
	N sufficient	N deficient	N sufficient	N deficient	
Palmitic acid (C16:0)	0.527 \pm 0.011	0.665 \pm 0.014	0.449 \pm 0.010	0.615 \pm 0.013	1.624 \pm 0.036
Heptadecanoic acid (C17:0)	0.049 \pm 0.001	0.054 \pm 0.001	0.035 \pm 0.000	0.078 \pm 0.001	0.148 \pm 0.003
Stearic acid (C18:0)	< LoD	0.194 \pm 0.004	< LoD	0.044 \pm 0.001	0.252 \pm 0.005
Lipid content (%g/gDW) Cis-9-oleic acid (C18:1n9c)	0.979 \pm 0.023	2.762 \pm 0.064	0.307 \pm 0.007	1.142 \pm 0.026	6.959 \pm 0.164
Linoleic acid (C18:2n6c)	2.070 \pm 0.048	2.312 \pm 0.054	0.915 \pm 0.021	0.742 \pm 0.017	5.027 \pm 0.119
Arachidic acid (C18:3n6)	0.105 \pm 0.002	0.099 \pm 0.002	0.063 \pm 0.001	0.095 \pm 0.002	0.158 \pm 0.003
γ -linolenic acid (C18:3n3)	6.461 \pm 0.152	4.322 \pm 0.101	6.819 \pm 0.160	6.663 \pm 0.156	6.865 \pm 0.162
Total	10.21	10.42	8.61	9.40	21.14

On one hand, 150 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ illumination runs yielded the same growth rate as B3N medium screening (around 0.0460 h^{-1} , under 75 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$). Light, as any substrate has a 3-phase impact on culture: 1. 385 affinity phase - growth rate linear increase, 2. plateau phase - stable growth rate, 3. inhibition - growth rate decrease (Ohad et al. 1990). The fact that subculturing and moderate light runs yielded the same growth rate is a token of fact that both were led in the plateau phase, hence no light limitation nor inhibition. This is confirmed by the value of the Fv/Fm ratio (0.778 ± 0.038) which indicates proper functioning 390 of the photosynthesis apparatus. Indeed, a value of 0.8 indicates a fully photosynthetically capable culture, while the culture is still considered 'healthy' down to 0.6 (Masojídek et al. 2003).

On the other hand, 300 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ illumination runs yielded a lower growth rate (0.0174 h^{-1}). In addition, photosynthetic apparatus analysis indicated a 395 hindered Fv/Fm ratio of 0.570 ± 0.059 . This could be further investigated using yields measurements (Fig. 4). Both cultures exhibit the same trend, a decreasing Y(II) with increasing light. Still lower values are reported for the high illumination culture. An important difference is highlighted under 1028 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$. In this case the moderately illuminated culture photosystems II forward about 33.1 400 % of incoming energy, while the highly illuminated cultures only 11.6 %.

Photosystems II energy forwarding at this illumination level is harmful for the whole photosynthetic apparatus on the long run (more than 20 minutes) (Baroli et al. 1998). In time, it will induce severe damages. Thus a low Y(II) under excessive illumination shows that the strain has adapted and diverts energy toward the NPQ 405 mechanisms. This is validated by the Y(NPQ) values which clearly indicate that the strain grown under high light is consistently more efficient at safely disposing excess energy. All this shows that the cells cultured under 300

$\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ illumination have deployed a strategy to cope with high light.

410 All in all this validates the occurrence of light stress under 300
 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ which is not the case under 150 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$. Still
this stress is not doing excessive harm to the photosynthetic apparatuses and
cannot be referred to as photodamage. With this in mind, the results will be further
discussed, first by commenting profiles obtained under nitrogen sufficient then
415 nitrogen limited conditions.

4.2 Pigment profile

The photobioreactor obtained pigment profiles are quite common among
Chlorophyta phylum. Indeed, they exhibit the two main pigments chlorophyll *a*
and *b* in large quantities. In addition, natural occurrence of lutein is observed.

420 Lutein is a primary photosynthetic carotenoid pigment widening light absorption
spectrum (Gong et al. 2016). This characteristic is also shared by resembling
Scenedesmus genus (Wiltshire et al. 2000). No major difference is observed in the
pigment profile between the two light conditions. Even though, 300
 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$ induces light stress, no carotenoid overexpression is
425 detected. This may suggest that pigment profile modification would have required
higher illumination.

On the contrary, nitrogen deficiency had an impact on cell pigment content. While
it did not modify the profile per se, nitrogen deficiency strongly decreased the total
amount of pigments per cell. Indeed, when confronted with nitrogen limitation
430 cells break down nitrogen containing materials, such as chlorophyll molecules, to
redirect nitrogen towards growth metabolism (Adams et al. 2014). Concomitantly,
lutein content also decreased. Lutein does not contain nitrogen, still this fall can be
explained by the nature of this pigment. As less chlorophylls molecules are

present, only a reduced number of antenna can have their absorption spectrum
435 widened.

Finally, long term nitrogen starvation altered the cell pigment profile. Chlorophyll
a and *b* as well as lutein contents reached very low levels. Violaxanthin is lacking
from this profile (confirmed by a second analysis). Furthermore potent antioxidant
carotenoids are detected (canthaxanthin and astaxanthin, β -caroten absence is
440 thought to come from its very apolar nature, hence it is barely extractible using
methanol). These results agree well with other authors findings on *Desmodesmus*
komarekii exposed to high light (250 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$) (Hanagata and
Dubinsky 1999). Still, in our case, the later carotenoids are thought to play a role
in mitigating ROS species targeting enhanced lipids synthesis products (Zhang et
445 al. 2013). Indeed, under 75 $\mu\text{molPhotonPAR}/\text{m}^2/\text{s}$, with low chlorophyll contents,
no violaxanthin - meaning xanthophyll cycle downregulation -, light induced ROS
production seems unlikely to happen. Furthermore, in our case, the production of
storage lipids is confirmed by FAME profiling.

4.3 FAME profile

450 First of all, the obtained FAME profiles are in agreement with those found in the
literature for similar microalgae (Wiltshire et al. 2000). In addition, amounts are
within well established range (3 to 60 %g/gDW (Morris et al. 2008; Gris et al.
2014; Vadiveloo et al. 2015)). Still, in our photobioreactor runs the total amount is
around 10 %g/gDW, while an increase could have been expected. This can be
455 explained by the fact that nitrogen deficient samples were chosen among those in
late exponential (or early stationary) phase. This procedure left only a limited
amount of time for the strain to accumulate lipids (Gifuni et al. 2019), yet profile
modification is observed, which was the objective. Furthermore, 1-month nitrogen
starvation allowed for longer lipid storage build up leading to a doubled strain

460 lipid content. In any configuration, the three major lipids are oleic acid, linoleic and γ -linolenic acid.

Going one step further, it is possible to determine which lipid is produced commonly or under light or nitrogen stress. As one can see in Table 2, linoleic and oleic acid productions are favoured by low light while γ -linolenic acid is enhanced
465 by high illumination. Higher content of given lipids under low light can be explained by the fact that they might be among the primary targets of light induced ROS (Zhang et al. 2013). Nitrogen deficiency induced appearance (with low amounts) of stearic acid, accumulation of palmitic and oleic acids while γ -linolenic acid concentration dwindled. Finally, long term (nitrogen starvation)
470 lipid content increase is mainly supported by oleic and linoleic acids. Hence pinpointing them as the storage lipids of this strain.

Oleic acid is a monounsaturated fatty acids with numerous health benefits both as feed and food. As feed, its inclusion in dairy cows diet has been shown to enhance milk production in the most productive cows (Souza et al. 2019). As food, it has
475 demonstrated positive effects in the treatment and prevention of cardiovascular and autoimmune diseases, metabolic disturbance and cancer (Sales-Campos et al. 2013). Linoleic acid is a polyunsaturated fatty acid which shows valuable health properties. Namely, from a metabolic perspective, it improves plasma lipid profile and promotes long-term glycaemic control and insulin resistance (Marangoni et al.
480 2020). From a population scale perspective, its consumption reduces incidence of cardiovascular diseases, metabolic syndrome and type 2 diabetes (Belury et al. 2018).

Conclusion

Freshwater microalgae *Desmodesmus sp.* has been isolated and profiled under
485 nitrogen sufficient and deficient conditions as well as under regular illumination
and light stress. Pigment profiles are common among Chlorophyta phylum, still
with natural presence of lutein. Lipid profiling showed that the three main lipids
produced by this strain are oleic, linoleic and γ -linolenic acids. While γ -linolenic
acid production is enhanced by high illumination, oleic and linoleic acids
490 production is favoured by nitrogen deficiency. Those two last fatty acids have
valuable health properties, hence pave the way towards potential
biofuel/nutraceuticals covalorization of this strain.

Declaration of author contributions

495 WL, PP and VP initiated and designed the study. WL, MP and VP led the
experimental work. CG developed and led the profiling procedure. WL and NG
led the photosynthetic apparatuses analysis. The authors critically interpreted the
results. VP drafted the manuscript, the other authors corrected it. All authors
approve the manuscript.

500 Conflict of interest

None to disclose.

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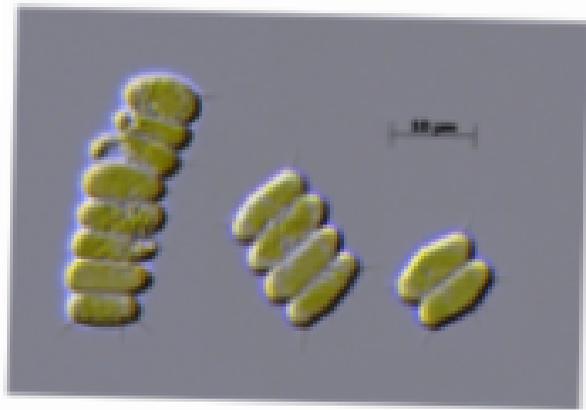
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Desmodesmus sp.

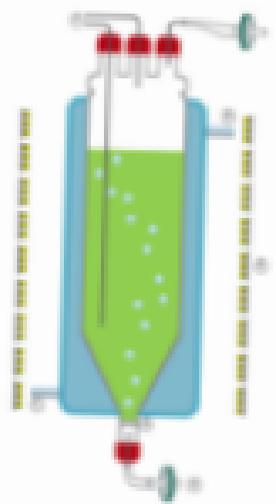


Media screening
Nitrogen affinity

Lipid & pigment
profiles

Identification of
valuable lipids

Potential biofuel
covalorization



CO₂

Light
stress

Nitrogen
stress