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Biochemical analysis of cyanobacterial strains isolated from different Tunisian inland waters: therapeutical and nutritional potential

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ABSTRACT

The last decade period has been characterized by a remarkable increase in the need for food, bioenergy, and pharmaceutical compounds. Meanwhile, microalgae have been in the center of interest as a rich and sustainable source of highly valuable natural products and bioactive compounds. In this perspective, the present study aimed at investigating the biochemical composition of 10 cyanobacterial strains collected from several Tunisian inland waters, seeking to identify the possible use of cyanobacteria in nutraceuticals and pharmaceuticals industry, and for animal and human consumption. Ten local strains of cyanobacteria were isolated from several Tunisian inland waters and cultured in controlled laboratory conditions. Fatty Acid methyl esters were analyzed in each strain by Gas chromatography-Mass spectrometry (GC-MS) as well as total proteins, carbohydrates and different phycobiliproteins (phycocyanin, allophycocyanin and phycoerythrin). Results showed that Planktothrix agardhii, Leptolyngbya sp1., and Leptolyngbya sp2. contained 14 different fatty acids, among which oleic acid was the most abundant. These might be important in nutraceutical and agronomic sectors due to the potent antioxidant and larvicidal activities of oleic acid. Cylindrospermopsis raciborskii presents extraordinarily high levels of phycocyanin, making it an exceptional natural reservoir of effective and non-toxic drugs that are of particular interest against cancer and neurodegenerative diseases. Extracts from cyanobacterial strains represent a natural and available source of proteins, of highly valuable fatty acids, and especially of phycocyanin which is known by its extremely health beneficial properties. The nutritional and functional complements provided by these extracts can be used in the prevention of various diseases and also as effective and environmentally friendly insecticides. © 2019 International Formulae Group. All rights reserved.

Keywords: Cyanobacteria; biochemical composition; fatty acids; phycobiliproteins; valorization.

INTRODUCTION

Photoautotrophic microscopic organisms include a large variety of species of widespread occurrence and with diverse morphological, physiological, and biochemical properties (Kata et al., 2018). Among these organisms, cyanobacteria

occupy all types of habitats as they are able to overcome a plethora of environmental constraints, including cold, drought, salinity, photo-oxidation, anaerobiosis, osmotic pressure, and exposure to UV radiations (Samarakoon and Jeon, 2012; Moscoso et al., 2013). Furthermore, recognized as important

primary producers, cyanobacteria are major actors in the aquatic food chain, playing significant roles in carbon, nitrogen, and oxygen biochemical cycles (Moscoso et al., 2013).

Considering the increasing need for food, feeding, bioenergy, and pharmaceutical compounds, particular attention has been paid during the last decade to microalgae as a sustainable natural source of biologically active compounds that do not compete with the usual food resources. In particular, cyanobacteria have gained interest for producing third generation biofuels (both biomass and H₂ production) as well as for being valued in various fields, especially in agro-food, aquaculture, biotechnology, and pharmacy (Trabelsi et al., 2017). For example, cyanobacterial protein has received worldwide attention both as a food supplement and as an alternative source of food. Some species of Anabaena, Nostoc, and Arthrospira are consumed as food due to their high protein and fiber content (Plaza et al., 2009). The cyanobacterium Arthrospira platensis exhibits extremely high protein content (up to 85.5% of dry weight) (M'baye et al., 2011).

Furthermore, phycobiliproteins are among the most valuable accessory pigments that can be extracted from cyanobacteria, given their extremely health beneficial effects. These water-soluble, fluorescent proteins are largely considered as potent functional ingredients with very interesting applications in various fields, especially in biotechnology, medicine, and pharmacy (Jalal et al., 2013).

Therefore, the present research aimed at sampling, identifying, and characterizing 10 cyanobacterial strains collected from several Tunisian inland waters. Their biochemical composition (fatty acids, proteins, carbohydrates) as well their phycobiliprotein constituents (phycocyanin, allophycocyanin, and phycoerythrin) were determined with the purpose of identifying the cyanobacterial strains' possible use in nutraceuticals, pharmaceuticals, animal feed, and even as human food.

MATERIALS AND METHODS Cyanobacteria strains isolation an

Cyanobacteria strains: isolation and culture

Cyanobacteria strains were isolated from seven different Tunisian inland waters (river, lagoon, dam reservoir, spring water) located in northern, central and southern Tunisia (Table 1, Figure 1). The strains (Chroococcus sp.: Chroo-CH, A. platensis: Arthro-KB, Leptolyngbya sp1.: Osci-NB-01, Leptolyngbya sp2.: Lepto-CH, Limnothrix sp.: Osci-BM-01, Planktothrix agardhii: Plank-SS-01, Spirulina sp.: Spir-ML, Lyngbya sp.: Lyng-ML, Anabaenopsis circularis: Pseud-01, Cylindrospermopsis raciborskii: Cyl-NB-05) were isolated in the laboratory by capillary isolation on liquid medium and observed inverted using an microscope (Leica microsystems, Wetzlar, Germany). For each strain, isolation was performed on a single individual (a single cell, a colony or a filament) in 300 µl of appropriate culture medium and the growing was carried out by scaling-up culture volumes gradually from the initial isolation to the final batch. All strains were cultivated in 2 L volume laboratory flasks using monospecific batch culture system under sterile conditions. Each strain was cultured in either BG11 (Rippka et al., 1979) or CONWAY (Blancheton, 1985) culture medium depending on the origin of each strain (Table 1). Culture media were sterilized by autoclaving at 120 °C for 20 minutes before use. The cultures were incubated at 25 \pm 1 °C in a thermostatically controlled room and illuminated with cool white fluorescent tubes at an intensity of 2000 lux (approximately 10 μmol m⁻² s⁻¹) in a 16:8 h light: dark regime. Each culture was stirred continuously with air at a low and constant flow rate. The period of cultivation was from 8 to 26 days and the cells were harvested by centrifugation (3000 rpm for 10 minutes) and lyophilized using a Christ Alpha 2-4 LD Plus freeze-92 dryer (Harz, Germany). The freezedrying operation was carried out under vacuum at -70 °C and for 2 days long. For all biochemical analysis (fatty acids, proteins, carbohydrates, and phycobiliproteins), each culture was sampled in the exponential growth phase only.

Molecular identification

Total genomic DNA was extracted from fresh cells harvested by centrifugation during the exponential growth phase of each strain using the DNeasy® Plant Mini Kit (QIAGEN GmbH, Hilden, Germany) and following the protocol for Gram-negative bacteria as recommended by manufacturer. The DNA was subsequently kept frozen at -20 °C until use. DNA extracts were checked by electrophoresis in a 1% agarose gel. The DNA was quantified with a Thermo Fisher Scientific spectrometer (Massachusetts - USA) and subsequently diluted to 200 ng/µL solution.

Nuclear-encoded 16S rRNA gene segments were amplified with PCR primers 27 Fw and 1494 Rev under conditions described by (Fathalli et al., 2011). All PCR amplifications were performed in 20 µL aliquots containing 10 pmol of each forward and reverse primers (Invitrogen, Carlsbad, California), 1x Reaction Buffer (Invitrogen, Carlsbad, California), 250 µM of each deoxynucleosides triphosphate (OMEGA biotec TQAC136), 2.5 mM MgCl₂ (Invitrogen, Carlsbad, California), 0,5 U of Taq DNA polymerase (Invitrogen, Carlsbad, California) and 5 to 10 ng of genomic DNA template. All amplicons were analyzed electrophoresis in 1% agarose gel (Invitrogen, Carlsbad, California) run in 1× TBE buffer, ethidium stained with bromide photographed under UV trans-illumination.

A total of 100 μL of PCR of each amplified product was purified using a PureLink® PCR Purification Kit (Invitrogen, Carlsbad, California) following the protocol supplied by the manufacturer before being sent for direct sequencing. Nucleotide sequences were obtained and submitted to the (Basic Local Alignment Search Tool) BLAST database

(http://www.ncbi.nlm.nih.gov/BLAST) for identification.

All sequences were subsequently submitted to the GenBank database (accession numbers are given in Table 2).

Fatty acids analysis

For each strain, after lipid extraction mixture of dichloromethane: methanol: 0.7% aqueous NaCl (1:2:0.8, v/v/v) according to a method adapted from (Bligh and Dyer, 1959). Samples were then derivatized with 1 mL of 7% boron trifluoride in methanol (BF₃-MeOH) and 1 mL of toluene (100 °C for 60 minutes) according to a method adapted from (Morrison and Smith, 1964) to prepare Fatty Acid methyl esters (FAMEs). After heating, 2 mL of distilled water were added and samples were centrifuged at 3000 rpm during 4 minutes. The organic layer was separated and the aqueous layer was extracted three times with 1 mL of toluene. The obtained Fatty Acid Methyl Esters (FAMEs) were then analyzed by GC-MS (Gas chromatography-mass ThermoQuest spectrometry) on a GC/Thermo ITQ 900, ion trap spectrometer run with the Xcalibur 2.1 software package. The Electron Ionization method was used at 70 eV. FAMEs were separated on a Thermo TG-5MS column (30 m long, 0.25 mm i.d., 0.25 µm film thickness). The temperature program was as follows: initial, 160 °C; ramp 1.5 °C/min to 270 °C, with injector temperature maintained at 280 °C. The carrier gas used was helium at a constant flow of 1 mL/min, and solutions were injected in split mode with a ratio of 10. GC-MS analytical standards were purchased from Sigma-Aldrich (Menhaden:Capelin 1:1 fish oil; FAME Mix no. 47885-U). The GC-MS results were given in relative area (percentage of 1 peak = 1 FAME). Areas are calculated from raw data, and the ionization cross-section of each class of molecules was not taken into account. A blank sample (100% solvent) was injected between each analysis to avoid carryover. FAMEs were identified using the mass spectral database (NIST, 2008) and diagnostic ions in mass spectra, with additional FAME identification based on the retention index (equivalent chain length: ECL). Two sets of analytical standards were injected to confirm the FAME identification. According to the international nomenclature, the position of the first double bond is given by the (n-x) notation, by counting the number of carbon atoms from the methyl end (Revel et al., 2016). Each sample was injected in triplicate.

Total protein and carbohydrate estimation

Total protein was assessed by the Lowry method using bovine serum albumin (BSA) as standard instead of casein because of the greater similarity between the BSA amino acids composition and that of microalgae. Fifty mg of freeze-dried sample of each culture were manually ground with pestle and mortar in 4 mL distilled water and then centrifuged at 3000 rpm for 10 minutes at temperature. The Folin-Ciocalteu reactive (Sigma Co.) was diluted in two volumes of distilled water (1:2) and 0.5 mL of the diluted reactive was added to 1 mL of sample, previously mixed with 5 mL of the reactive "C" [50 volumes of reactive "A" $(2.0\% \text{ Na}_2\text{CO}_3 + 0.1 \text{ N NaOH}) + 1 \text{ volume of}$ reactive "B" (1/2 volume of 0.5% CuSO₄ 5H2O +1/2 volume of 1.0% $C_4H_4NaO_6$ 4H₂O)]. After the addition of each reactive, samples were stirred for 2 s in a test tube stirrer. Thirty minutes after the start of the chemical reaction at room temperature (time needed for colour development), spectrophotometric readings were achieved at 750 nm and compared BSA standards which ranged between 25 and 200 µg mL⁻¹.

Carbohydrate concentrations were assessed by according to the phenol–sulphuric acid method (Dubois et al., 1956). To prepare the carbohydrate solution, 100 mg of dry weight cyanobacteria sample were hydrolysed with 2.5 N chloridic acid for 1 hour at 100 °C. A 0.2 ml aliquot of this solution is mixed with 1 ml of 5% aqueous solution of phenol (prepared immediately before use) in a test tube and then 5 mL of concentrated sulfuric acid were rapidly added to the mixture. After

10 minutes, the tubes are vortexed for 30 seconds and placed for 30 minutes in a water bath at room temperature for colour development. The samples were then analyzed at 490 nm using a spectrophotometer (Thermo Fisher Scientific, Massachusetts - USA) against control and compared with the standard glucose solution (concentration from 10 to 200 μ g mL⁻¹).

Phycobiliproteins determination

The quantification of phycobiliproteins (phycocyanin: PC, allophycocyanin: APC and phycoerythrin: PE) in the microalgal strain was carried out using the spectrophotometric method of (Bennett and Bogorod, 1973). After 5 minutes centrifugation of a 10 ml volume of fresh cultures at 3000 rpm, the pellet was washed with buffer solution (1 M Tris-Cl; pH = 8.1) and one volume of cell mass was subsequently re-suspended in five times of the volume of the same buffer. After that, continuous freezing at - 20 °C and thawing at +4 °C, and sonication (10 minutes with cycles of 30 seconds) were applied to all samples in order to allow the destruction of the microalgal cell walls. The cell fragments were then separated by centrifugation at 12000 rpm for 10 minutes before taking the supernatant spectrophotometric estimation phycobiliproteins using the buffer as a blank. Absorbance measurements were performed at 620, 652 and 562 nm wavelengths using an UV - visible spectrophotometer (Heaios Y, Thermo Scientific; Massachusetts - USA). The amount of PC, APC and PE in the sample was calculated using the following formula (Bennett and Bogorod, 1973; Horvath et al., 2013):

 $\begin{array}{ll} PC \ [mg/ml] = (A_{620}\text{-}0.474xA_{652})/5.34 \\ APC \ [mg/ml] = (A_{652}\text{-}0.208xA_{620})/5.09 \\ PE \ \ [mg/ml] \ = \ \ (A_{562}\text{-}2.41xPC\text{-}0.849xAPC)/9.62 \end{array}$

All experiments were done in triplicates and the data are presented mean of three independent experiments with standard deviation.

Table 1: Cyanobacterial strains isolated from Tunisian inland waters (river, lagoon, reservoir, spring water) with location and culture mediums.

Order	Strain		Sampling site	Culture medium	Sampled and isolated in	
Chroococcales	Chroococcus Chroo-CH	sp.	Chanchou river Lat (33.540° N); Long (9.432° E) S: 4.2 PSU	BG11	Present work	
	Arthrospira platensis Arthro-KB		Korba lagoon Lat (36.613° N); Long (10.884° E) S: 16 PSU	BG11	Present work	
	Leptolyngbya S	sp1.	Nabhena reservoir Lat (36.061° N); Long (9.865° E) S: 0.5 PSU	BG11	Fathalli et al. (2011)	
	Leptolyngbya s	sp2.	Chanchou river Lat (33.540° N); Long (9.432° E) S: 4.2 PSU	BG11	Present work	
Oscillatoriales	Limnothrix Osci-BM-01	sp.	Bir Mcherga reservoir Lat (36.509° N); Long (10.010° E) S: 1.6 PSU	BG11	Fathalli et al. (2011)	
	Planktothrix agardhii Plank-SS-01		Sidi Saad reservoir Lat (35.381° N); Long (9.689° E) S: 1.9 PSU	BG11	Fathalli et al. (2011)	
	Spirulina Spir-ML	sp.	Maltine river Lat (34.245° N); Long (10.194° E) S: 45 PSU	CONWAY	Present work	
	<i>Lyngbya</i> Lyng-ML	sp.	Maltine river Lat (34.245° N); Long (10.194° E) S: 45 PSU	CONWAY	Present work	
Nostocales	Anabaenopsis circularis Pseud-01		Siliana spring water Lat (35.911° N); Long (9.335° E) S: 0.2 PSU	BG11	Present work	
	Cylindrospermop raciborskii Cyl-NB-05	psis	Nabhena reservoir Lat (36.061° N); Long (9.865° E) S: 0.5 PSU	BG11	Fathalli et al. (2011)	

S: Salinity; Lat: Latitude; Long: Longitude.

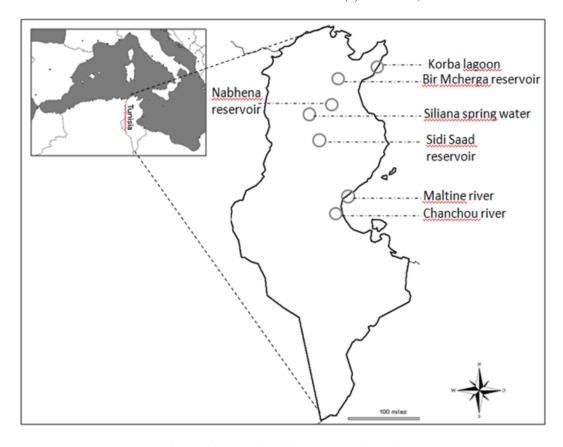


Figure 1: Map of Tunisia with sampling sites.

Table 2: Molecular identification of the cyanobacterial strains isolated from different Tunisian inland waters with % of similarity according to BLAST database and accession numbers of 16S rRNA or rpoC1 sequences.

Strain	Taxon according to BLAST	0/ of similarity	Accession numbers		
Strain	database	% of similarity	16S rRNA	rpoC1	
Osci-NB-01	Leptolyngbya sp.	99	MG762090	-	
Osci-BM-01	Limnotrix sp.	100	MG762091	-	
Plank-SS-01	Planktothrix agardhii	100	MG762092	-	
Cyl-NB-05	Cylindrospermopsis raciborskii	96	-	HQ389355	
Pseud-01	Anabaenopsis circularis	99	MG098078	-	
Arthro-KB	Arthrospira platensis	100	MG098079	-	
Lepto-CH	Leptolyngbya sp.	99	MG098077	-	
Spir-ML	Spirulina sp.	99	MG518486	-	

RESULTS Fatty acid profiles

Results showed that the investigated strains produced 14 types of fatty acids, with carbon chains ranging from C14 to C18 with odd and even carbon numbers as well as different degrees of unsaturations (Table 3). The percentage of unidentified fatty acids was always under 1% for all strains, with the exception of Lepto-CH which showed 3% of unidentified fatty acids.

Proteins and polysaccharides

In the present study, the percentage of total protein ranged between 4 and 30% of DW (Figure 2). Apart from Osci-BM-01 and Cyl-NB-05, which presented a relatively good protein amount (about 30% of DW each), the other strains exhibited lesser percentages, reaching a minimum of less than 5% DW in Chroo-CH.

As regards to total polysaccharides, the majority of the strains exhibited low carbohydrate contents (less than 6 mg/ml or 30% DW) (Figure 3). The only strains presenting a relatively high total carbohydrate content exceeding 6 mg/ml or 30% of DW were Lepto-CH, Cyl-NB-05, and Chroo-CH (9.3; 6.6 and 6.1 mg/ml or 46.6; 33.3 and 30.6% DW, respectively).

Phycobiliproteins

Quantitative evaluation of PC, APC, and PE content was performed in all investigated microalgal strains; the results are displayed in Figure 4. Among all the essayed isolates, Cyl-NB-05 contained maximum total phycobiliprotein content, exceeding 3150 µg/ml of fresh culture. Interestingly, the PC level in this Nostocales cyanobacterium occurs at more than ten-folds higher than in the other two phycobiliprotein groups taken together (2873 µg/ml of PC against 277 µg/ml for APC and PE combined) and at more than two-folds higher compared with the PC level in all the other 10 strains. Significantly smaller amounts of PC were found in the other tested strains which were in the range between 11 (Pseud-01) and 671 µg/ml (Plank-SS-01). In terms of APC production, the maximum yield was found in the two Oscillatoriales cyanobacterium strains Osci-NB-01 and Plank-SS-01 (362 and 326 µg/ml, respectively), while PE was present at lower quantities comparing with the other two constituents (between 20 and 126 µg/ml) in the investigated strains, except for Chroo-CH and Pseud-01, where it was the major phycobiliprotein produced.

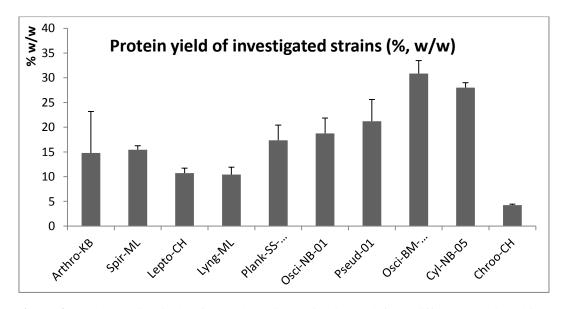


Figure 2: Total protein yields of cyanobacterial strains isolated from different aquatic habitats expressed in % of dry weight. Bars indicate the standard deviation of three replicates.

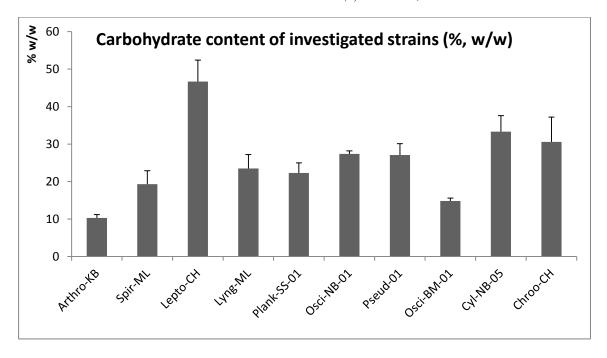


Figure 3: Total carbohydrate content of cyanobacterial strains isolated from different aquatic habitats expressed in % of dry weight. Bars indicate the standard deviation of three replicates.

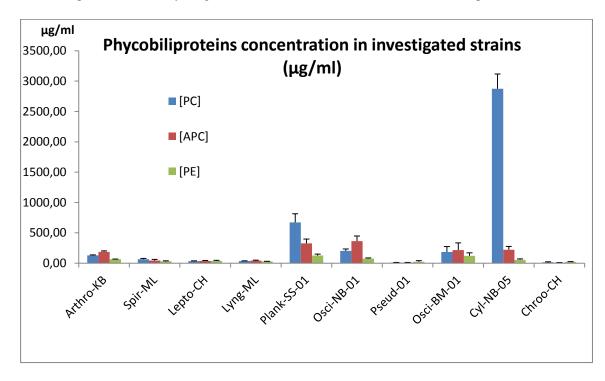


Figure 4: Phycobiliproteins concentration of microalgal strains isolated from different aquatic habitats expressed in μ g per ml of fresh culture. Bars indicate the standard deviation of three replicates (PC = Phycocyanin; APC = Allophycocyanin; PE = Phycocythrin).

Table 3: Fatty acid relative composition expressed in % of total fatty acids content of the investigated strains. Values are means of three replicates \pm standard deviation.

Fatty acids	Arthro-KB	Osci-NB-01	Lepto-CH	Lyng-ML	Osci-BM-01	Plank-SS-01	Spir-ML	Chroo-CH
C14:0	nd	nd	nd	0.62 ± 0.2	22.89 ± 2.5	nd	25.98 ± 2.5	nd
C15:0	nd	0.39 ± 0.1	1 ± 0.1	0.68 ± 0.2	nd	0.75 ± 0.1	1.23 ± 0.0	nd
C16:0	48.83 ± 4.5	29.56 ± 3.6	26.68 ± 5.1	38.08 ± 1.5	23.34 ± 0.9	32.42 ± 4.9	20.2 ± 1.9	60.79 ± 5.6
C17:0	nd	0.84 ± 0.3	nd	nd	nd	nd	nd	nd
C18:0	nd	1.26 ± 0.3	2.61 ± 0.5	1.57 ± 0.2	nd	2.56 ± 0.9	1.26 ± 0.4	1.04 ± 0.2
Σ Saturated	48.83 ± 4.5	32.05 ± 4.5	30.29 ± 5.6	40.95 ± 0.9	46.24 ± 2.03	35.74 ± 5.9	48.69 ± 1.1	61.83 ± 5.5
C14:1	nd	nd	nd	nd	17.02 ± 1.0	nd	nd	nd
C16:1 (n-7)	3.62 ± 0.7	25.27 ± 3.4	3.2 ± 0.2	44.54 ± 3.4	19.76 ± 0.6	21.84 ± 0.9	48.84 ± 1.3	10.09 ± 0.7
C16:1 (n-5)	nd	nd	nd	nd	12.55 ± 0.3	nd	nd	3.61 ± 0.5
C16:1 (n-3)	nd	nd	19.67 ± 1.0	nd	nd	nd	nd	nd
C17:1	0.12 ± 0.04	0.54 ± 0.1	nd	nd	nd	nd	nd	nd
C18:1 (n-9)	nd	23.32 ± 6.7	21.41 ± 3.3	0.65 ± 0.3	4.41 ± 1.1	20.32 ± 3.5	0.85 ± 0.1	4.5 ± 3.2
C18:1 (n-7)	nd	4.03 ± 1.7	7.11 ± 1.0	2.33 ± 0.7	nd	4.55 ± 2.3	1.61 ± 0.1	1.85 ± 0.6
C18:1 (n-6)	nd	nd	1.36 ± 0.3	nd	nd	nd	nd	nd
Σ MUFA	3.74 ± 0.4	53.17 ± 1.4	52.75 ± 2.7	47.52 ± 2.7	53.76 ± 2.03	46.72 ± 2.7	51.3 ± 1.1	20.06 ± 2.6
C14:2	nd	nd	nd	nd	nd	nd	nd	7.05 ± 0.5
C16:2	nd	0.95 ± 0.1	nd	11.11 ± 2.5	nd	nd	nd	3.43 ± 0.9
C18:2 (n-6)	19.91 ± 2.5	13.81 ± 3.0	13.92 ± 3.1	nd	nd	17.53 ± 4.5	nd	1.02 ± 0.6
C16:3	nd	nd	nd	nd	nd	nd	nd	6.59 ± 1.8
C18:3	26.84 ± 3.2	nd	nd	nd	nd	nd	nd	nd
Σ PUFA	46.75 ± 3.1	14.76 ± 2.9	13.92 ± 3.1	11.11 ± 2.5	nd	17.53 ± 4.5	nd	18.1 ± 2.9
PUFA/MUFA	12.5 ± 1.6	0.27 ± 0.04	0.26 ± 0.04	0.23 ± 0.06	nd	0.37 ± 0.09	nd	0.9 ± 0.02

nd = not detected.

DISCUSSION

Fatty acid composition

As can be seen in Table 3, the cyanobacterium Arthro-KB distinguished from all other strains as it was the only producer of linolenic acid, (C18:3) with a percentage exceeding 25%, which approached the results reported by many authors working on A. platensis, such as Xue et al. (2002). From a pharmaceutical point of producing important amounts of linolenic acid is of particular interest since several findings reported on the antimicrobial activity of this tri-unsaturated fatty acid, particularly against a group of pathogenic germs, such as Staphylococcus aureus, Candida albicans, and Aspergillus niger (Xue et al., 2002; Herrero et al., 2006).

Moreover, the percentages of the major fatty acids in Arthro-KB found in this study were in accordance with those in previous works (Xue et al., 2002; Colla et al., 2004), indicating that A. platensis was characterized by a fatty acid profile dominated mainly by palmitic (C16:0), linoleic (C18:2), and linolenic acid (C18:3) and that fatty acids with longer chains (> 20 carbons) were almost not detected. This profile represents a remarkable added-value to the therapeutic potential of this strain since linoleic acid (about 20%) is additionally efficient against human pathogens and also against some aquatic pathogenic bacteria, even at relatively low concentrations (Benkendorff et al., 2005).

As regards to oleic acid (C18:1), it was recorded in all strains, with the exception of Arthro-KB. However, only three strains, namely Plank-SS-01, Lepto-CH and Osci-NB-01, contained a high percentage of oleic acid (25%, 28%, and 27%, respectively). Several studies on this monounsaturated acid have demonstrated its powerful antioxidant effect (Ceron et al., 2007) and its anti-platelets aggregation property (Xu et al., 2018), the acid thus presenting a promising potential as an accessible and safe alternative to synthetic molecules. In addition, Harada et al. (2000) demonstrated the larvicidal activity of this fatty acid extracted from P. agardhii, especially against mosquito larvae. The latter functionality gives the top oleic acid producer strains characterized in the present work (Plank-SS-01, Lepto-CH, and Osci-NB-01) even more valuable potential as they can be used as natural sources of insecticides that are both effective and environmentally friendly.

Otherwise, palmitoleic acid (C16:1) was found in our study at significant percentages in Plank-SS-01, Osci-NB-01, Lepto-CH, and Osci-BM-01 (between 20 and 30% of total fatty acid content) and ultimately even in a much higher amount in Spir-ML (about 50%) and Lyng-ML (around 45%). Many authors reported the remarkable and significant bacteriolytic activity of palmitoleic acid-even at relatively low concentrationsagainst human pathogens and pathogenic bacteria as well as its ability to reduce the risk of certain heart diseases (Benkendorff et al., 2005; Plaza et al., 2009). Interestingly, among all these strains that are particularly rich in palmitoleic acid, only Lepto-CH appeared to produce omega-3 fatty acids (around 85% of total amount of palmitoleic acid) which are widely known for protective role against their diabetes, rheumatoid arthritis. rheumatism. diseases, digestive and metabolic diseases, neuro-cognitive disorders, and colorectal cancer (Shahidi and Ambigaipalan, 2018).

Proteins and polysaccharides content

Compared to the present study, higher protein amounts determined with the same analyzing method have been reported from an Indian strain of C. turgidus (Chinnu et al., 2014) and A. platensis (Lopez et al., 2010). We infer that the differences are probably due-apart from the strain effect—to the variations in culture conditions, such as the level of illumination (18:6 versus 12:12 light:dark cycle), the availability of nutrients (BG11 medium versus CFTRI and Zarrouk's medium), the culture technique (Batch versus continuous culture), or the growth phase. Indeed, these various culture conditions can lead to differences in protein production levels and can affect the quantity and quality of microalgal biochemical composition (Lopez et al., 2010; Salama et al., 2013).

In the present study, the two non-toxic cyanobacteria strains Osci-BM-01 and Cyl-NB-05 (Fathalli et al., 2011) showed a remarkably high protein productivity, thus representing excellent candidates to be valued in fast growing markets, such as animal feed and aquaculture. Furthermore, our protein best producing strains might be exploited beyond their nutritional values in several other interesting applications, such as nutraceuticals (Chacon-Lee and Gonzalez-Marino, 2010), cosmeceuticals (Sekar and Chandramohan, 2008), and pharmaceuticals (Samarakoon and Jeon, 2012).

In relation to total polysaccharides analyzed in the present work in terms of total sugars, results showed that most strains displayed relatively modest carbohydrate amounts, not exceeding 6 mg/ml. The lowest amount was found in Arthro-KB (2 mg/ml or 10.3% DW), which is in agreement with Becker (2007), who reported a similar value (11% DW) for this species.

On the other hand, it can be seen that the oscillatoriales strain Lepto-CH was most productive of total carbohydrates probably to the presence of a mucilaginous sheath covering its single trichome which is clearly visible under the microscope. Indeed, this mucilaginous sheath with high contents in pectic acids and acid mucopolysaccharides was also reported by Mandal et al. (2011). Moreover, some authors were able to induce an increase in the thickness of the mucilaginous sheath layer of some cyanobacterial species (for example the Indian strain Lyngbya majuscula) and consequently to improve the carbohydrate productivity by applying stressing culture conditions, such as prolonged exposure to UV-B dose (Mandal et al., 2011).

Phycobiliproteins

According to our results, the *Cylindrospermopsis* strain Cylind-NB occupies an outstanding position among all investigated isolates by being by far the most productive of total phycobiliproteins, with a total amount exceeding 3000 µg/ml. With regard to PC, and_compared to results reported

by other authors, the Cylindrospermopsis strain essayed in the present study appears to be an extraordinarily rich source of this pigment-protein complex. valuable instance, a much lower PC amount (about 5 ug/ml) in a Cylindrospermopsis raciborskii strain isolated from surface water samples of Lake Balaton (Hungary), using the same extraction technique (repeated cycles of freezing and thawing combined sonication), was reported by Horvath et al. (2013).

Aside from the strain effect—which may be the main factor explaining the huge difference observed between the factors isolates—other may also have intervened, especially the high light intensity (40 μmol m⁻² s⁻¹) applied in Horvath et al. (2013) compared to the present work (27µmol m⁻² s⁻¹). In fact, the negative effect of applied light intensities of more than 25 µmol m⁻² s⁻¹ on the PC production by cyanobacterial isolates was highlighted in several studies. For instance, De Oliveira et al. (2014) reported that the level of incident light is indeed one of most influence the factors that cyanobacteria metabolism and that lower light intensities were more advantageous in terms of total phycobiliprotein yields produced by two isolates of edaphic algae from the genus Nostoc. The authors suggested that when the light availability increased, the content of these pigments decreased as a strategy for prevention of photo-oxidative damage caused by the production of free radicals.

With PC such an exceptional productivity, the Cylindrospermopsis isolate investigated in the present study may indeed be considered as a superrich natural source of this highly valuable phycobiliprotein group which had been proved to possess numerous potent bioactivities with extremely health beneficial effects. Among the most interesting therapeutic and nutritional functions attributed to this natural, non-toxic substance that are especially sought for food and pharmaceutical industry is the antioxidant and radical scavenging activities (Samarakoon and Jeon, 2012). In fact, several synthetic commercial antioxidants, such as butylated hydroxyanisole

(BHA), butylated hydroxytoluene (BHT), tertbutylhydroquinone (TBHQ), and propyl gallate (PG) have been used widely to retard the oxidation and peroxidation processes. However, the use of these synthetic antioxidants is often associated with potential health hazards, making the search for natural antioxidants as safe alternatives in food and pharmaceutical industry even more appealing and urgent. Furthermore, it had been shown that the protective action of antioxidants may be valued on many levels. Firstly and most importantly, they have a positive effect on human health as they can protect the human body against damage by ROS (Reactive Oxygen Species) which attack macromolecules, such as membrane lipids, proteins, and DNA, leading to many health disorders, such as cancer, diabetes mellitus, aging, and neurodegenerative diseases. In addition, antioxidants can prevent food deterioration caused by lipid oxidation as well as the formation of undesirable secondary lipid peroxidation products by ROS, such as superoxide anion, hydroxyl radicals, and H₂O₂ which are known to be responsible for the decrease in the nutritional value of lipid foods hence affecting their safety and appearance (Pangestuti and Kim, 2011).

Some other studies also reported the effectiveness of PC as a proliferation inhibitor of cancer cells *in vivo* and *in vitro*. For example, Liu et al. (2000) reported the capacity of PC extracted from *Spirulina platensis* to significantly inhibit the growth of human chronic myelogenous leukemiablastcrisis K562 cells in a dose-dependent manner.

In addition, several epidemiological studies highlighted another very interesting therapeutic property of PC that may open very promising perspectives in the medical field, especially in the treatment of oxidative stress-induced neuronal injuries. For instance, the neuroprotective role of PC against cell death caused by 24 h potassium and serum (K/S) withdrawal in rat cerebellar granule cell (CGC) cultures was demonstrated by Rimbau et al. (2001).

Taken together, the PC super rich strain characterized in the present work, Cylind-NB, may be of particular interest in this field since it can be considered as a natural reservoir of effective and non-toxic drugs to be used in the fight against some of the most common neurodegenerative diseases, such as Alzheimer's and Parkinson's.

Conclusion

The extracts from the cyanobacterial strains, analyzed in the present study, represent a natural and available source of proteins and of some highly valuable fatty acids. Thus, the nutritional and functional complements provided by these extracts, particularly those from the Oscillatoriales strains Plank-SS-01, Arthro-KB, and Lepto-CH, can be used in the prevention of infectious diseases and of cardiovascular and coronary disorders and even as effective and environmentally friendly insecticides. Meanwhile. the Cylindrospermopsis raciborskii strain, Cyl-NB-05, characterized in the present study appears to be of huge and exceptional potential to be valued in therapeutic and medical sectors due to its extraordinary production and accumulation capacity of phycocyanin. Finally, we suggest that further studies be carried out on the produced protein's quality and especially on the absence of potential toxicity for human animal consumption. The themselves, their cultivation conditions, as well as the extraction and purification methods remain important basic elements to be taken into account in the biological activities sought.

COMPETING INTERESTS

The authors declare that they have no competing interests.

AUTHORS' CONTRIBUTIONS

KG designed and performed the experiments, derived the models and analysed the data. AF and CF contributed to sample preparation and verification of the analytical methods, as well as the interpretation of the results, especially with regard to the molecular

identification section. ABRJ and MSR supervised the project, helped conceiving the original idea, provided critical feedback and contributed in the shaping of the research, analysis and manuscript. All authors discussed the results and contributed to the final manuscript.

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