



Nutrient Deprivation Mobilizes the Production of Unique Tocopherols as a Stress-Promoting Response in a New Indigenous Isolate *Monoraphidium* sp.

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Singh R, Paliwal C, Nesamma AA, Narula A and Jutur PP (2020) Nutrient Deprivation Mobilizes the Production of Unique Tocopherols as a Stress-Promoting Response in a New Indigenous Isolate Monoraphidium sp. Front. Mar. Sci. 7:575817. doi: 10.3389/fmars.2020.575817 The economic feasibility of the next-generation algal biofuels rely largely on the simultaneous production of high-value-added bioproducts (HVABs) in these photosynthetic cell factories. In this study, a newly isolated indigenous microalga belonging to the selenastraceae family *Monoraphidium* sp. (designated as CABeR41) has been isolated and characterized for its relevant biofuel precursors along with other HVABs, i.e., tocopherols. Our preliminary data sets demonstrate accumulation of higher amounts of lipids (266.6 mg g⁻¹ dcw) along with antioxidant molecules, such as α - and δ -tocopherols (456 μ g g⁻¹ dcw and 974 μ g g⁻¹ dcw), simultaneously when subjected to nutrient deprivation. The total tocopherol content has been increased in nitrogen-deplete conditions to 1450.24 μ g g⁻¹ dcw from 638.2 μ g g⁻¹ dcw. In conclusion, bioprospecting of new isolates from natural habitats and fine-tuning of specific conditions without compromising growth has potential for developing into a sustainable industrial strain for concomitant production of biofuels and bioproducts.

Keywords: Monoraphidium, nutrient deprivation, tocopherols, microalgae, environmental stress

INTRODUCTION

Global population demands and rapid industrialization have led to remarkably high consumption of natural petroleum, creating urgency to look for alternative, renewable sources (Gao et al., 2019). Microalgae are photosynthetic microorganisms fixing ~9% of solar energy to generate biomass efficiently compared to terrestrial plants, sequestering ~513 tons ha⁻¹ year⁻¹ of atmospheric CO₂ without competing for arable land with food crops (Chen et al., 2018). Microalgae are sustainable and potential feedstock for biofuels and other high-value-added bioproducts (HVABs), such as carotenoids, polyunsaturated fatty acids (PUFAs), and vitamins (Durmaz, 2007; Borowitzka, 2013; Skjånes et al., 2013; Paliwal et al., 2016). The primary barrier for commercialization of algal-derived biofuels is presumably the production costs. Henceforth, the concurrent processing of co-products from microalgae would be offset for the production of biofuels (Stephens et al., 2010; Wijffels and Barbosa, 2010). Strategies for the accumulation of lipids and carotenoids can be triggered by stress conditions, such as nutrient deprivation and/or other external factors (Sun et al., 2018).

Macronutrients, such as carbon, nitrogen, and phosphorous along with other micronutrients (mainly comprising iron, manganese, and zinc), play an important role in microalgal growth. The effect of such nutrients are well documented in strains such as Chlorella sorokiniana, Chlorella vulgaris, Dunaliella tertiolecta, and Tetraselmis suecica (Ghafari et al., 2018). Nitrogen, for example, is a key element of cellular biomolecules, including DNA, protein, and chlorophyll, which affects growth and other biosynthetic pathways (Simionato et al., 2013). Previous studies on nitrogen deprivation demonstrate the accumulation of reactive oxygen species (ROS), such as H_2O_2 , superoxide (O_2^-), and hydroxyl (OH) radicals that lead to growth hampering, reduced protein, and chlorophyll synthesis preceded by concomitant increase in the lipid and/or carbohydrate contents (Pérez-Pérez et al., 2012). ROS molecules are mainly involved in cellular signaling processes, but excess production may lead to apoptosis. In order to control and mitigate stress, the cell produces biomolecules, such as tocopherols, polyphenols, carotenoids, flavonoids, and other enzyme scavengers, like superoxide dismutase (SOD), catalase, and ascorbate peroxidase (APX), as an integral part of the internal defense mechanism (Cirulis et al., 2013).

Tocopherols are possibly regarded as efficient microalgal HVABs that defend fatty acids against the oxidation process and ensure the stability of the lipid membrane (Fritsche et al., 2012; Lv et al., 2015). α-tocopherol has the greatest antioxidant potential and is one of the leading tocopherols present in both algae and plants (Mudimu et al., 2017). Biosynthesis of tocopherol exists in chloroplasts, and because human beings cannot synthesize it, it is provided as a dietary supplement (Munné-Bosch and Alegre, 2002). These tocopherols can prevent and inhibit oxidative damage and, hence, are used as cardioprotective additives as well as UVprotectants and antiaging creams in cosmetics (Poljšak and Fink, 2014). A study indicates the existence of another type of tocopherol, i.e., δ -tocopherol, with unique antioxidant and anti-inflammatory properties (Ahsan et al., 2014). However, a number of algal strains, such as Nannochloropsis oculata (1.4 mg g⁻¹), Haematococcus pluvialis (0.8 mg g⁻¹), Euglena gracilis (2.6 mg g⁻¹), and Coccomyxa sp. (0.42 mg g⁻¹) were identified as potential α-tocopherol producers (Ogbonna et al., 1998; Durmaz, 2007; Mudimu et al., 2017).

In the present study, our focus is to identify and optimize the strain for the production of biofuels and HVABs simultaneously; thus, it could be categorized as an efficient strain for sustainable algal biorefinery. The stress-promoting responses on newly isolated indigenous microalga *Monoraphidium* sp. was studied for understanding their physiology and regulation of carbon flux into major metabolic pathways, such as lipid, carbohydrate, and protein biosynthesis when subjected to nutrient deprivation. We further evaluate the impact of the stress phenomenon on photosynthetic activity and also characterized few HVABs, such as tocopherols, to understand their role in oxidative damage within these microalgal cells.

MATERIALS AND METHODS

Isolation of Strain(s)

Microalgae was isolated from freshwater springs near the Dharamshala ($32^{\circ}14'40.1''N76^{\circ}20'03.2''E$) in the Kangra district of Himachal Pradesh, India. Samples (2 mL) collected from the springs were inoculated in BG-11 medium (50 mL) initially and grown for 14 days following 16:8 light/dark periods under light intensity ($80 \ \mu E \ m^{-2} \ s^{-1}$) and temperature of $25^{\circ}C$ with constant shaking at 150 rpm. Isolation of axenic cultures was carried out by serial dilution followed by the spread-plate technique until unialgal colonies were obtained (Lee et al., 2014) and the isolated stock cultures were maintained on agar slants for further experiments.

Molecular Characterization of Newly Isolated Indigenous Strain

The unialgal/axenic culture was maintained in Erlenmeyer flasks (50 mL) containing minimal medium (BG-11) at 80 μ E m⁻² s⁻¹ light intensity, 16:8 h light/dark photoperiods and 25°C at 150 rpm. Genomic DNA was extracted from the cells ($\sim 10^9$ cells) in the logarithmic phase using the Zymo Research kit (Quick-DNA Fungal/Bacterial Microprep Kit, Catalog No. D6007, United States) as per the manufacturer's instructions. The DNA amplification was carried out with Internal Transcribed Spacer (Fritsche et al., 2012) region primers as follows: forward (5'-GGACAGAAAGACCCTATGAA-3') and reverse (5'-TCAGCCTGTTATCCCTAGAG-3'). The polymerase chain reaction (PCR) performed in the BioRad T100TM Thermal Cycler using the program: preheating at 95°C for 3 min followed by 35 cycles of denaturation for 1 min at 95°C, annealing at 58°C for 1 min, and extension at 72°C for 90 min followed by another 10 min extension at 72°C. Later, the PCR products were visualized by electrophoretic separation on 1.5% agarose gels and sequenced. The ITS rRNA sequences were searched for homology in the NCBI database employing nucleotide BLAST¹. All the retrieved sequences were aligned by multiple sequence comparison by log-expectation (MUSCLE) alignment (Edgar, 2004) followed by phylogenetic analysis using molecular evolutionary genetics analysis (Hall, 2013; Lee et al., 2014) employing the neighbor joining method with 1000 rounds of bootstrapping (Saitou and Nei, 1987). The sequence is deposited to the GenBank with accession no. MT622543.

Culture Conditions

A total number of cells ($\sim 10^8$) were inoculated at a logarithmic phase containing 1.5 g L⁻¹ of NaNO₃ (termed nitrogen replete) and 0 g L⁻¹ of NaNO₃ (nitrogen deplete) in BG-11 medium, respectively, i.e., approximately 0.056 g L⁻¹ DCW biomass with an initial optical density (OD₇₅₀) of 0.1. The cells were grown in BG-11 medium for 10 days at 25°C in 16:8 h light/dark photoperiods and a light intensity of 150 µE m⁻² s⁻¹ with constant shaking at 150 rpm. Sampling was carried out at regular time intervals of 0, 2, 4, 6, 8, and 10 days.

¹http://blast.ncbi.nlm.nih.gov

The cell growth was calculated using following equation (Kareya et al., 2020):

$$\mu = \ln(N_2/N_1)/(t_2 - t_1)$$

where μ is the specific growth rate and N_1 and N_2 are the biomass at times (t_1 and t_2), respectively. Doubling time was also calculated by the following equation:

Doubling time =
$$\ln(2)/\mu$$

Biochemical Characterization

Lipid estimation was done by the sulfophosphovanillin assay (Mishra et al., 2014). Briefly, 1 mL of culture was pelleted down into 100 μ L of distilled water. Then, 2 mL of 98% H₂SO₄ was added to cells and incubated at 100°C for 10 min. After cooling for 5 min in ice, 5 mL of phosphovanillin reagent (0.6 g vanillin in 10 mL of ethanol in 90 mL distilled water and 400 mL 85% phosphoric acid) were added, and the samples were incubated for 10 min at 37°C, absorbance measured at 530 nm.

A modified phenol-sulfuric acid method was used for total carbohydrate estimation (Paliwal et al., 2015). Then, 0.2 mL of 98% H_2SO_4 was added to 100 μ L of cells and hydrolyzed at room temperature for 1 h; subsequently, 5% phenol with 1 mL of H_2SO_4 was added and incubated at room temperature for 20 min after vortexing. The absorbance was measured at 490 nm.

A modified biuret method used for quantification of protein content. Approximately 1 mL of cell suspension was pelleted down and 1 ml of extraction buffer (25% NaOH in methanol) added. The reaction mixture was incubated for 15 min at 80°C. The sample was cooled to room temperature and centrifuged to remove debris. Then, 300 μ L of the supernatant was treated with 150 μ L of CuSO₄ solution (0.21% CuSO₄ in 30% NaOH) and incubated at room temperature for 10 min, and further absorbance was measured at 310 nm.

Nile Red Staining

For visualization of triacylglycerols (TAGs) in microalgae, cells were stained with fluorescent lipid-sensitive Nile Red fluorescent dye (Janchot et al., 2019). A Nile Red concentration of 0.5 μ g mL⁻¹ in 25% of dimethyl sulfoxide (DMSO) was added to 1 mL of cell culture and stained for 20 min at 37°C (Cooksey et al., 1987). Lipid bodies were visualized by fluorescence microscopy, and images were captured through a Nikon 80i epifluorescence microscope (Nikon, Japan) equipped with a camera Q-Imaging Micropublisher 3.3 RTV (Real Time Viewing), and Image-Pro Plus v6.0 software (Media Cybernetics, Inc., Bethesda, MD, United States).

Scanning Electron Microscopy

Morphological changes in the *Monoraphidium* sp. CABeR41 when subjected to nutrient deprivation was captured by scanning electron microscopy (SEM). Samples were prepared by washing $\sim 10^7$ cells with 0.1 M sodium phosphate buffer (pH 7.2). Later, the samples were fixed overnight with 2.5% of glutaraldehyde and incubated at 4°C for SEM analysis (Lambrecht et al., 2015).

Estimation of Nitrate Utilization

Nitrate uptake by the cells was carried out spectrophotometrically (Collos et al., 1999). Briefly, 1 mL of culture was pelleted down at room temperature, and supernatant was diluted 50-fold with deionized water. The residual nitrate content was determined by measuring the absorbance at 220 nm.

PAM Measurements

A dual-PAM100 fluorometer (Heinz Walz, Effeltrich, Germany) measured the fluorescence signals and microalgal cells were darkadapted for 30 min before measuring fluorescence. Maximum fluorescence F_m was determined by directing a saturation pulse (6000 µmol photons m⁻² s⁻¹; $\lambda = 660$ nm) and quantum efficiency of PS II calculated by $(F_v/F_m = (F_m - F_o)/F_m)$ (Agarwal et al., 2019). Further quantum yields of photochemical quenching Y, non-photochemical quenching Y (NPQ), and thermal energy dissipation in the form of heat and fluorescence occurring during closed PSII system Y (NO) was calculated from the fluorescence values obtained (Kareya et al., 2020).

Fatty Acid Methyl Esters (FAME) Quantification by GC-MS

Quantification of FAME was done by the method described previously (Shaikh et al., 2019). Briefly, $\sim 10^8$ cells were hydrolyzed and methyl *trans*-esterified at 80°C with 50 µg heptadecanoic acid as an internal standard in 300 µl of 2% H₂SO₄ and methanol solution for 2 h. Then, 300 µl of 0.9% NaCl and 300 µl of hexane were added simultaneously after *trans*-esterification and mixed thoroughly for 20 s. Centrifugation was done at 3000 × g for 3 min and the clear upper hexane layer taken into vials. Then, 1 µl of sample was injected into an Agilent 6890 gas chromatograph (GC) coupled to a mass spectrometer (MS) equipped with Omega Wax 250 column (30 m × 0.25 mm × 0.25 µm). The FAME was detected by an electron ionization detector. The running conditions for GC-MS were described by Agilent's RTL DB Wax method.

Quantification of Pigments and Tocopherols

For pigment estimation, 1 mL of culture was centrifuged at 7500 × *g* for 10 min at 4°C. Pigment extraction was performed by adding 95% absolute ethanol and glass beads to the pellet followed by vortexing for 10 min at room temperature. The debris was pelleted down, and the supernatant was used for measuring absorbance at 648, 664, and 470 nm. The concentration of Chlorophyll a (C_a), Chlorophyll b (C_b), and Carotenoids (C) were calculated by using the following equation (Lichtenthaler and Buschmann, 2001):

 $C_a(\mu gmL^{-1}) = 13.36A_{664} - 5.19A_{648}$

 $C_b(\mu gmL^{-1}) = 27.43A_{648} - 8.12A_{664}$

 $C(\mu gmL^{-1}) = (1000A_{470} - 2.13C_a - 97.64C_b)/209$

Tocopherols were estimated by a previously described method (Moltó-Puigmartí et al., 2009) with slight modifications. Briefly,

 $\sim 10^7$ cells were centrifuged at 10,000 \times g for 10 min, and 500 μ L of absolute ethanol and glass beads were used for extraction of tocopherols. The solution was vortexed for 10 min with 2 min intermittent ice bath incubation. For phase separation, 500 µL of hexane and 200 µL of distilled water were added after extraction followed by centrifugation $(2200 \times g)$ for 10 min at 4°C. The upper hexane layer was collected and evaporated to dryness under a nitrogen stream. Finally, the dried extract was resuspended into 200 μ L of dichloromethane:methanol (2:1) and analyzed by an Agilent HPLC system (1260 Infinity II LC System, United States). Then, 10 µL of samples were injected and eluted in isocratic mode through an Eclipse plus C18 column $(95 \text{ Å}, 4.6 \times 150 \text{ mm}, 5 \mu\text{m}; \text{Agilent Technologies, United States})$ using acetonitrile:methanol (60:40, v/v) at a constant flow rate of 0.6 mL min⁻¹. Tocopherols were detected using a fluorescence detector with an excitation/emission wavelength of 297/328 nm, respectively. Tocopherols were quantified based on peak area ratio using analytical standards.

Statistical Analysis

All the experiments were carried out in biological triplicate and the analysis of variance done by Fischer's least significant difference (LSD) (Meier, 2006; Di Rienzo et al., 2011). The values were represented as average \pm standard error (SE) and the statistical differences were calculated at p < 0.05.

RESULTS

Identification of New Isolate (Microalgae) From Natural Environments

With the growing world population and increasing demand for high-quality nutritional supplements, new microalgae strains need to be explored from various natural environments providing a promising resource for HVABs in the near future. The indigenous strain isolated from Dharamshala (India) was morphologically identical to the Selenestraceae family with spindle shape and tapering toward apices. It is not possible to determine the identity of the strain by morphology alone due to the vast diversity existing among algal strains. Therefore, ITS rRNA sequencing was performed to identify the strain, and the obtained sequence was submitted to the NCBI GenBank with accession number (MT622543). The phylogenetic analysis of the ITS rRNA sequence showed close proximity with Monoraphidium contortum with a resemblance of 86.5% (Figure 1) and, hence, termed Monoraphidium sp. #CABeR41 ([#]local strain identifier).

Growth Profiles and Biochemical Changes Under Nitrogen Deprivation

At optimal conditions, with an initial inoculum of $\sim 10^6$ cells mL⁻¹, the *Monoraphidium* sp. CABeR41 achieved a biomass of 0.682 g L⁻¹ within 10 days, presenting better growth with a specific growth rate 0.20 d⁻¹ and doubling time of 78.0 h (**Figure 2A**). An adequate supply of macronutrients along with other micronutrients play a crucial role in the growth of these



FIGURE 1 | Phylogenetic analysis of new isolate using ITS regions by MEGA's neighbor joining method with 1000 rounds of bootstrapping.



TABLE 1 | Biochemical yields and productivity in replete and deplete conditions of

 Monoraphidium sp. CABeR41 on 10th day of cultivation.

Parameters	Nitrogen replete	Nitrogen deplete
Biomass (mg L ⁻¹)	647.5 ± 6.19^{a}	350 ± 3.28^{b}
Biomass productivity (mg $L^{-1} d^{-1}$)	62.56 ± 0.62	29.31 ± 0.31
Specific growth rate (d^{-1})	0.20 ± 0.01	0.11 ± 0.03
Lipid (mg g ⁻¹)	165.7 ± 3.59^{b}	266.3 ± 7.57^{a}
Lipid (mg L ^{-1} d ^{-1})	7.77 ± 0.05	5.87 ± 0.06
Protein (mg g ⁻¹)	520.7 ± 10.86^{a}	$138.4\pm4.08^{\rm b}$
Protein (mg L ⁻¹ d ⁻¹)	21.70 ± 0.04	0.33 ± 0.02
Carbohydrate (mg g ⁻¹)	238.3 ± 5.54^{a}	247.2 ± 15.95^{a}
Carbohydrate (mg L ⁻¹ d ⁻¹)	10.57 ± 0.38	4.78 ± 0.26
Total chlorophyll (mg g ⁻¹)	$14.5\pm0.65^{\text{a}}$	3.1 ± 1.75^{b}
Carotenoid (mg g ⁻¹)	$3.77\pm0.24^{\text{a}}$	$1.2\pm0.19^{\rm b}$

Statistical analysis was performed using one-way ANOVA with Fisher's LSD test, and significance was determined if p < 0.05. a and b both denotes the significant difference between the two conditions (a, maximum; b, minimum).

microalgae. When subjected to nitrogen starvation (**Figure 2A**), a severe effect on the growth was observed, as it starts retarding after the 4th day of cultivation and cells succumb on the 8th day, showing signs of early senescence. The biomass yield on 10th day of cultivation in nitrogen deplete medium was 0.35 g L⁻¹ with a specific growth rate 9.1 d⁻¹ and doubling time of 219.0 h. In addition, we also estimated utilization of nitrate uptake for 10 days in these cells under replete conditions, and it seems to be approximately 0.38 g L⁻¹ (**Figure 2B**).

The biochemical constituents of these photosynthetic cell factories were estimated to evaluate the changes occurring during nitrogen deplete (**Table 1**). Among these, the protein content rapidly decreased after the onset of stress from the 4th day

and was reduced from 358.8 mg g^{-1} dcw to 138.3 mg g^{-1} dcw, i.e., decreased by nearly twofold. When cells are in stress, the overall protein content reduces and is catabolized to form precursor molecules. During replete conditions, there was increase in protein content until the 4th day of cultivation (486.9 mg g^{-1} dcw), and a decline was observed after the 4th day until the 8th day (Figure 2C), which may be due to nutrient deficiency within the media but recovers to maximal on the 10th day. For carbohydrates, the overall content declines initially from 117.3 mg g⁻¹ dcw to 96.1 mg g⁻¹ dcw on the 2nd day (Figure 2D), but increases substantially from 187.1 mg g^{-1} dcw to 238.5 mg g^{-1} dcw until the 10th day of cultivation. However, carbohydrate yield was increased in nitrogen-deplete conditions from 117.1 mg g⁻¹ dcw to 361.0 mg g⁻¹ dcw (by nearly 1.5fold) during the first 6 days and later started to decrease on the 8th and 10th days of cultivation. Similarly, the total lipids in the nitrogen deplete condition increased from 13.9 mg g^{-1} dcw to 266.3 mg g^{-1} dcw until the 10th day of cultivation, i.e., increased by eightfold (Figure 2E), whereas in nitrogen replete (Figure 2E), the lipid content was increased to 83.3 mg g^{-1} dcw from 14.0 mg g⁻¹ dcw by the 4th day followed by increase up to 165.7 mg g⁻¹ dcw by the 10th day. This phenomenon was also confirmed by fluorescence staining using NR uniform under both nutrient-replete and -deplete conditions in which a significant amount of lipid bodies was observed in deplete conditions (Figure 3).

Microscopy Reveals Morphological Changes During Nitrogen Stress

The light microscopy analysis of *Monoraphidium* sp. CABeR41 displayed that the shape of the cells remains intact without any

change showing chloroplast density until the 8th day in nitrogenreplete conditions while the cells in the deplete state tend to be obese with significant changes in chloroplast structures. In starved conditions, morphological changes were observed after the 4th day itself, when cells became obese. Moreover, a massive change in chloroplast structures was observed after the 4th day with a gradual decline in the chloroplast density. NR-stained fluorescent light imaging displayed that the yellowstained globules called lipid bodies were prominently visible at the 10th day in replete conditions, whereas the globules remain smaller in size. However, under nitrogen stress conditions, the cells start accumulating lipid droplets on the 4th day, whereas the cells tend to get obese gradually from day 6 (**Figure 3**). SEM images depict a rigid shaped structure with no disruptions in the nitrogen-replete condition, whereas in nitrogen depletion, the cells were showing the signs of the weakened cell wall structure (**Figure 4**).

Changes in Photosynthetic Efficiency, Total Chlorophyll, and Carotenoid Contents

The total chlorophyll and carotenoid contents also followed a similar trend as observed in the growth profile of these cells. The total chlorophyll content in the control conditions almost doubled from 10.0 mg g⁻¹ dcw to 19.35 mg g⁻¹



FIGURE 3 | Time-course analysis for visualizing lipid droplets in *Monoraphidium* sp. CABeR41 subjected to nitrogen replete and deplete conditions by Nile Red fluorescence microscopy.



dcw in first 4 days: a ~0.93-fold increase. After the 4th day, the chlorophyll content declined, yielding 15.38 mg g⁻¹ dcw on the 10th day (**Figure 5A**). In nitrogen starvation, the total chlorophyll content declined rapidly from 10.27 mg g⁻¹ dcw to 0.63 mg g⁻¹ dcw on the 10th day, showing signs of reduced photosynthetic efficiency. Moreover, photosynthetic machinery, particularly in the case of nitrogen deficiency, is the major component affected during stress in these cell factories. Total carotenoid content in the replete condition increased from 1.0 mg g⁻¹ dcw to 7.90 mg g⁻¹ dcw by the 4th day and seems to have no change until the 10th day while in nitrogen-deplete cells they accumulate carotenoids until the 4th day up to 2.20 mg g⁻¹ dcw and later decline steadily to 0.90 mg g⁻¹ dcw by the 10th day (**Figure 5B**).

The maximum quantum efficiency of photosystem II (F_{ν}/F_m) in replete cells was 0.8 on the 2nd day and increased simultaneously to remain constant at 0.84 on the 4th day of cultivation (**Figure 6A**), demonstrating that the cells are in a healthy state and the photosynthetic machinery of cells were highly active although a slight decrease in the F_{ν}/F_m was observed after the 4th day even though it remained constant until the 10th day with an F_{ν}/F_m ratio of 0.68. However, in the nitrogen stress-deplete condition, F_{ν}/F_m between the 4th to the 10th day decreased from 0.61 to 0.24 (40.38%) demonstrating that cells were in stress, and photosynthetic efficiency declined drastically. Moreover, the growth profile also correlates with this phenomenon showing a decrease from the 4th day onward in nitrogendeplete conditions.



FIGURE 5 | (A) Line diagram representing the time-course dynamics of total chlorophyll content (in mg g^{-1} dcw) in *Monoraphidium* sp. CABeR41 subjected to stress condition; (**B**) time-course variation in the total carotenoid content (in mg g^{-1} dcw) of *Monoraphidium* sp. subjected to nutrient replete and deplete conditions. Statistical analysis was performed using one-way ANOVA with Fisher's LSD test and significance was determined if p < 0.05. a and b both denotes the significant difference between the two conditions (a, maximum; b, minimum).



FIGURE 6 | (A) Line diagram indicating maximum quantum yield of the PSII reaction centers (*F_V*/*F_m*) in replete and deplete conditions; **(B)** quantum yields of photochemical quenching Y[II], non-photochemical quenching Y[NPQ], and the non-light induced energy dissipated as heat or fluorescence Y[NO] in *Monoraphidium* sp. CABeR41 subjected to nutrient deprivation.

Time course changes with respect to quantum yields of photochemical quenching Y[II], non-photochemical quenching Y[NPQ], and thermal energy dissipation in the form of heat and fluorescence occurring during a closed PSII system Y[NO] were also calculated (**Figure 6B**). The Y[II] of both the replete and deplete cells remained approximately similar until the 4th day of cultivation, whereas a decrease was observed in deplete cells after the 4th day until the 10th day. The Y[II] of replete cells decreased after the 4th day but remained constant from the 6th to the 10th day as observed with the F_v/F_m ratio. In addition, a significant increase in Y[NPQ] was also observed in nitrogen-deplete cells on the 10th day of cultivation.

Fatty Acid Methyl Esters (FAMEs) Quantification and Profiling

The FAME quantification and profiling of new indigenous isolate *Monoraphidium* sp. CABeR41 exhibits the following compositions: 22.4% of saturated fatty acids (SFA), 21.4% of monounsaturated fatty acid (MUFA), and 56.32% of PUFA. In replete conditions, PUFA content slightly increases from 54.21% to 56.32% up to the 10th day. Alpha linolenic acid [C18:3; ALA] (30.8%), linoleic acid [C18:2] (18.9%), and hexadecatrienoic acid [C16:3] (20.2%) were prominent fatty acids (**Table 2**). Moreover, ALA is an important ω -3 fatty acid in nature and was the most abundant fatty acid in *Monoraphidium* sp. CABeR41. During the stress-deplete condition, the strain shows an increase in MUFA (29.28% to 38.71% of total FAME) and SFA (16.49% to 27.74% of total FAME) (**Table 2**). Moreover, ALA content of *Monoraphidium* sp. CABeR41 declined from 29.95% to 20.02% of total FAME

in deplete cells. It can be due to oxidative damage within the cells being stressed or membrane lipids (PUFAs) recycled into TAGs attributing toward membrane remodeling, further this was also confirmed by scanning electron images (SEM) in which irregular and ruptured cells observed on the 10th day of the nitrogen-deplete condition. However, oleic acid [C18:1] (29.5%) and hexadecanoic acid [C16:0] (25.7%) were

TABLE 2 | Total lipid profiles of *Monoraphidium* sp. CABeR41 subjected to nitrogen replete and deplete conditions on 10th day of cultivation.

Fatty acids methyl esters (FAMEs)	Nitrogen replete	Nitrogen deplete
Hexadecanoic acid (C16:0) (%)	$20.2\pm4.5^{\mathrm{b}}$	25.7 ± 2.20^{a}
9-Hexadecenoic acid (C16:1) (%)	2.4 ± 0.66	3.1 ± 1.30
7,10-Hexadecadienoic acid (C16:2) (%)	$1.3\pm0.94^{\text{a}}$	$0.5\pm0.32^{\rm b}$
7,10,13-Hexadecatrienoic acid (C16:3) (%)	5.8 ± 2.83	3.2 ± 1.42
Methyl 4,7,10,13-hexadecatetraenoate (C16:4) (%)	8.4 ± 2.85^{a}	$3.3\pm0.26^{\text{b}}$
Octadecanoic acid (C18:0) (%)	$2.3\pm0.44^{\text{b}}$	$8.3\pm4.52^{\text{a}}$
Oleic acid (C18:1) (%)	18.9 ± 2.81	29.5 ± 5.51
Linoleic acid (C18:2) (%)	10.0 ± 0.17	7.7 ± 0.30
γ-Linolenic acid (C18:3) (%)	$30.8\pm0.39^{\text{a}}$	$18.8\pm1.78^{\rm b}$
Saturated fatty acid (% of total FAMEs)	$22.44\pm5.04^{\text{b}}$	$33.96\pm9.64^{\text{a}}$
Monounsaturated fatty acid (% of total FAME)	$21.24\pm6.72^{\text{b}}$	32.56 ± 10.16^{a}
Polyunsaturated fatty acid (% of total FAMEs)	$56.32\pm3.04^{\text{a}}$	$33.48\pm0.13^{\text{b}}$
Total FAME (mg g^{-1})	4.8	6.9

Statistical analysis was performed using one-way ANOVA with Fisher's LSD test, and significance was determined if $\rho < 0.05$. a and b both denotes the significant difference between the two conditions (a, maximum; b, minimum).



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the major fatty acids during nitrogen depletion (**Figure** 7). In addition, the total FAME content in nitrogen-deplete conditions seems to be higher (6.9 mg g^{-1} dcw) than replete cells (4.8 mg g^{-1} dcw).

Qualitative and Quantitative Analysis of Tocopherols

 α -Tocopherol content in this strain was estimated to be around 113.6 μ g g⁻¹ of dcw initially (Figure 8A) and gradually increased to 607.8 μ g g⁻¹ of dcw by the 10th day in the replete conditions. However, almost similar trends observed with δ -tocopherol (reporting for the first time in *Monoraphidium* sp.) with increase to 496.5 μ g g⁻¹ of dcw by the 10th day. During nitrogen starvation, α -tocopherol levels are lower as compared to control conditions, but the trend remains similar. a-tocopherol during the nutrient stress-deplete condition initially declines from 113.6 $\mu g \: g^{-1}$ of dcw to 99.48 $\mu g \: g^{-1}$ of dcw in first 2 days and later increased to 450.15 μ g g⁻¹ of dcw on the 10th day. Whereas the δ -tocopherol levels reached up to 1000.09 µg g⁻¹ of dcw by the 10th day (Figure 8B). In addition, the totaltocopherol content increased from 632.8 $\mu g g^{-1}$ of dcw to 1115.5 μ g g⁻¹ of dcw by the 10th day of cultivation in the replete cells, whereas in the deplete condition, the levels reached up to 1400 μ g g⁻¹ of dcw with a fold change of 1.14 (Figure 8C and Table 3).

TABLE 3 | Tocopherol yields and productivity of *Monoraphidium* sp. CABeR41 in nutrient replete and deplete conditions on the 10th day of cultivation.

Types of tocopherols	Nitrogen replete	Nitrogen deplete
α -Tocopherol (µg g ⁻¹)	607.86 ± 8.03^{a}	450.15 ± 6.00^{b}
α -Tocopherol (μ g L ⁻¹ d ⁻¹)	28.44 ± 0.06	8.86 ± 0.01
δ-Tocopherol (μ g g ⁻¹)	507.18 ± 7.50^{b}	1000.09 ± 18.4^{a}
δ-Tocopherol (μ g L ⁻¹ d ⁻¹)	20.71 ± 0.06	17.93 ± 0.02
Total tocopherols ($\mu g g^{-1}$)	1115.05 ± 15.8^{b}	1450.24 ± 24.10^{a}
Total tocopherols (μ g L ⁻¹ d ⁻¹)	49.15 ± 0.12	26.79 ± 0.3

Statistical analysis was performed using one-way ANOVA with Fisher's LSD test, and significance was determined if p < 0.05. a and b both denotes the significant difference between the two conditions (a, maximum; b, minimum).

DISCUSSION AND CONCLUSION

The inherent ability of microalgae to synthesize HVABs makes these photosynthetic cell factories an important source of feed material. Moreover, such bioactive molecules are of great interest in the nutraceutical and cosmetics industries. In the current scenario, due to the rapid exploration of natural HVABs, the development of the health-management paradigm focuses on general immune system regulation and the improvement of specific immune functions. Microalgae are relatively high in content and diversity of HVABs, which are more potent than plants for the antioxidant efficacy of microalgae (Sansone and Brunet, 2019). HVAB accumulation is associated with variables, such as C, N, P, and S and may also contribute to a higher biomass and lower quality of HVABs. Similarly, discovering natural HVAB resources and improving yields are the future priorities for sustainable algal refineries.

In the current study, the microalgal strain was isolated from freshwater springs near Dharamshala, Himachal Pradesh, India, at the elevation of 6,831 feet above sea level. The sample was collected in autumn when temperatures range between 30° C and 0° C. The microalgae isolated from such environmental zones show higher adaptability unique to their habitats and genetic architecture producing specific metabolites, such as HVABs in adverse conditions to overcome and acclimatize to the biotic and/or abiotic stress conditions. Therefore, microalgal bioprospecting is an approach to evaluate and analyze the potential of these algal strains for sustainable development in biorefineries.

The newly isolated microalga *Monoraphidium* sp. is a fastgrowing strain with a growth rate of 0.20 d⁻¹ and biomass productivity of 60.56 mg L⁻¹ d⁻¹, which is higher than other reported strains of *Monoraphidium* sp. (*Monoraphidium* sp. QLY-1; 29.1 mg L⁻¹ d⁻¹, *Monoraphidium* sp. QLZ-3; 30.3 mg L⁻¹ d⁻¹, *Monoraphidium* sp. YLY-2; 10.5 mg L⁻¹ d⁻¹) (Zhao et al., 2016), *Monoraphidium* sp. KMN5; 13.0 mg L⁻¹ d⁻¹) (Tale et al., 2014), *Monoraphidium* sp. HDMA-20; 36.3 mg L⁻¹ d⁻¹) (Lin et al., 2019). A recent study by our group involving *Parachlorella kessleri* subjected to a nitrogen stress condition demonstrates that the photosynthetic activity ceases affecting the phosphorus absorption, and cells shut down energy-expensive metabolisms, such as protein synthesis, by diverting the carbon flux to energy storage metabolites, such as lipids and/or carbohydrates (Shaikh et al., 2019). Similarly, in the present study, growth was severely affected during the nitrogen-deplete condition, and a simultaneous decline in protein content was observed after the 4th day. The decline in growth was also accompanied by the loss of chloroplasts and massive damage to the photosynthetic machinery. The protein content was significantly reduced in nitrogen starvation, which may be due to reduced NADPH levels inhibiting the amino acid biosynthesis, specifically the conversion of α -ketoglutarate to glutamate (Kudahettige et al., 2018), similar to a study reporting the same effect of nitrate depletion on protein content in Acutodesmus dimorphus (Chokshi et al., 2017). Also, our data show a significant increase in carbohydrate content until the 6th day in deplete conditions (a threefold change) that decreased later due to diversion of carbons toward lipids under prolonged nutritional stress. Likewise, elevated lipid accumulation was also observed on the 10th day of cultivation in nitrogen-starved cells. An increase in lipid content when subjected to nitrate limitation has been exhibited in microalgal strains, such as P. kessleri, Monoraphidium sp., Chlorella zofingiensis (Feng et al., 2012; Zhao et al., 2018; Shaikh et al., 2019). The morphology of the cells in nitrogen stress indicates changes in the form and size after the 4th day of starvation as confirmation by the NR microscopy by dwindling and acquiring lipids. Perhaps the NR staining of lipid bodies in nitrogen starvation is reported to enhance cells' neutral lipids, which is quite well documented.

In a nutrient-deprived environment, the total amount of chlorophyll declined as stress continued during the days, which could show signs of cell senescence. Chlorophyll molecules are likely to contain four nitrogen molecules within their form, rendering it much tougher for organelles in the absence of nitrogen to synthesize certain compounds as previously reported in Chlorella vulgaris (Ikaran et al., 2015; Janchot et al., 2019). Moreover, the maximum quantum efficiency F_v/F_m of PSII reaction centers in the stress cells of the deplete condition also observed a sharp decline reaching to 0.24 F_{ν}/F_m on the 10th day, indicating that photosynthetic functioning inside the cells are highly deteriorated and stress might lead to decrease in F_{ν}/F_m ratio as demonstrated in *Chlorella* sp. (White et al., 2011). In addition, downregulation of the light-harvesting complexes in stressed cells also contributes to reduced photosynthetic efficiency (Miller et al., 2010). Also, the decrease in F_{ν}/F_m ratio is attributed by inhibition of the electron transport change from both the donor and acceptor sites of the PSII reaction center, further promoting the diversion of excitation energy toward PSI, resulting in elevated electron flow toward PSI. Moreover the decrease in the electron flow represents the adaptive acclimation to downregulate PSII so as to dissipate the excess energy from the photosystem to maintain the oxidative state of QA (Lu and Vonshak, 2002).

The energy dissipation Y[NO] also tends to be increasing more in deplete-condition cells after the 4th day, indicating that replete cells are functioning actively toward photochemical reactions. Moreover Y[NO] is defined as a simple indicator for reduced plastoquinone status within photosynthetic membranes, but for high-irradiance microalgae, it is seen that the lower plastoquinone status is minimized, suggesting that the photochemical reactions in normal cells are much more efficient and able to regulate the transportation of photosynthetic electrons compared to the nitrogen-starved cells (Grieco et al., 2012; Nawrocki et al., 2020). During a stress phenomenon the increase in Y[NO] is accompanied by inactivation of PSII reaction centers, which dissipate excitation energy into heat (Maxwell and Johnson, 2000). A decrease in the F_{ν}/F_m ratio is also triggered by photo inactivation leading to oxidative damage and loss of PSII reaction centers (Baker, 2008). Moreover, the significant increase in Y[NPQ] in the nutrient-deplete condition on the 10th day, reveals that harvested light is dissipated as thermal energy due to inefficient photosynthetic apparatus, which attributes toward the defensive mechanism of the cell (White et al., 2011).

Nitrogen starvation modulates the production of tocopherols and lipids and induces morphological changes in Monoraphidium sp. CABeR41. In addition, we hypothesize that fatty acids are differentially regulated in stress conditions with an increase in fatty acid content. Major changes occurred in the decline of PUFAs and increase in the SFAs and MUFAs making the profile ideal toward biodiesel properties (Huerlimann et al., 2010). With higher PUFAs, the longer the ignition delay, resulting in retarded combustion, moreover decreasing oxidation stability and raising NOx emission (Benjumea et al., 2011). However, there are a number of reports highlighting the biodiesel potential of the Monoraphidium sp. (Patidar et al., 2014) reported fatty acid profile rich in SFA (30.92%-68.94% of total FAME) with biomass productivity of $(80.3-303.8 \text{ mg L}^{-1} \text{ d}^{-1})$ and total lipid (22.08%-46.54% of total FAME) under mixotrophic conditions. Similarly, less than 0.36 g L^{-1} of the nitrate Monoraphidium sp. exhibited the lipid productivity of 0.18 g L^{-1} d⁻¹ with fatty acid compositions (SFA; 37.22%, MUFA; 39.19%, PUFA; 26.30% of total FAME), making it suitable toward biodiesel (Dhup and Dhawan, 2014). Also, M. contortum is reported with biomass productivity of 896.0 mg L⁻¹ d⁻¹ with neutral lipid accumulation of 20.4% of dcw under nitrogen deplete. Moreover, there are no studies based on tocopherol accumulation in other Monoraphidium sp. Few other strains are reported, such as *Chlorella* sp. with biomass productivity of 55.6 mg $L^{-1} d^{-1}$ and ~18% dcw of lipids (Kassim and Meng, 2017), Scenedesmus sp. CCNM 1077 with biomass productivity of 22.7 mg $L^{-1} d^{-1}$ and ~18.9% dcw of lipids (Pancha et al., 2015) and Nannochlropsis sp. with lipid productivity of 44.2 mg L^{-1} d⁻¹ and ~27.68% dcw of lipids (Meng et al., 2015). In the present study, Monoraphidium sp. CABeR41 showed biomass productivity of 62.6 mg L^{-1} d^{-1} with ~16.5% dcw of lipids, which is competitive in terms of previously defined potent strains for biofuel production, and the relationship also enhances the value with co-products, such as tocopherols.

During photoautotrophic growth and development, tocopherol levels are dramatically increasing as a result of the alteration in expression of pathway genes, degradation, and recycling in response to environmental stresses. In addition, it was also found that an increase in tocopherol accumulation contributes to stress tolerance by limiting ROS generation while a decrease in the tocopherol levels may cause oxidative damage (Hasanuzzaman et al., 2014). A study has shown a tocopheroldeficient mutant (*vte1*) in *Arabidopsis thaliana* shows similar growth and photosynthetic efficiency in the control condition, whereas during photo-oxidative stress, chlorophyll content and photosynthetic quantum yield were severely affected in (*vte1*), indicating the role of tocopherol in maintaining photosynthetic efficiency (Porfirova et al., 2002).

In N. oculata, reduced nitrogen levels contribute to an increase in accumulation of α -tocopherol with a decreased growth rate. Further, α -tocopherol content was dependent on factors like nitrogen source, concentration, and phase of development (Durmaz, 2007). Reports show accumulation of α-tocopherol up to 2.3 mg g⁻¹ dcw in N. oculata and 2.6 mg g⁻¹ dcw in Euglena gracilis (Fujita et al., 2008), respectively; however, until now, there have been no reports of δ -tocopherol production among microalgae. Our study is the first to report both aand δ-tocopherol accumulation in Monoraphidium sp. CABeR41 with an adequate amount of total tocopherol accumulation $(1.4 \text{ mg g}^{-1} \text{ dcw})$ in nitrogen-deplete conditions. Henceforth, we hypothesize that, in stressed cells, the carbon flux is majorly diverted toward 2-methyl-6-phytyl-1,4-benzoquinone (MPBQ) rather than 2,3-dimethyl-6-phytyl-1,4-benzoquinone (DMPBQ), which is utilized by tocopherol cyclase producing a 0.6-fold increase in δ -tocopherol compared to α -tocopherol. Furthermore, stressed cells show a decrease in δ -tocopherol content after the 6th day, and this might be due to the fact that tocopherols are degraded and recycled as the concentration of ROS increases in the chloroplasts (Munné-Bosch and Alegre, 2002). In addition, there is a decrease in productivity due to a declined growth rate, and previous reports observed that productivity of both α-tocopherol and total fatty acid increases dramatically when cultivations were gassed with 5% CO_2 (v/v) in Coccomyxa sp., Desmodesmus sp., and Muriella terrestris. Since our strain was aerated with atmospheric air, there is room for enhancement of FA and total tocopherols via enrichment of the overhead atmosphere with the external CO₂ supply. Additionally, optimization of cultivation conditions and the reduction of production cost strategies can be employed in the future with flue gas supply and two-stage cultivation for enhanced productivity. Though adequate stress conditions, like depletion of

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nitrogen, high light, and high salinity, stimulate the synthesis and accumulation of HVABs. Recent advances in integrated omics enhance research perspectives in understanding gene regulation and molecular mechanism in different stress conditions, which provide insights in further enhancement of HVAB accumulation through sustainable strategies.

DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found below: NCBI (accession number: MT622543).

AUTHOR CONTRIBUTIONS

RS, AAN, and PJ designed the experiments. RS and CP executed the experiments. AN and PJ supervised the project. RS, CP, and AAN wrote the manuscript with all the input from the authors. All authors read and approved the final manuscript.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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