Synchronized nutrient stress conditions trigger the diversion of CDP-DG pathway of phospholipids synthesis towards de novo TAG synthesis in oleaginous yeast escalating biodiesel production

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**ABSTRACT**

In this study oleaginous yeast, *Rhodosporidium kratochvilovae* HIMPA1 grown in glucose synthetic medium containing different nitrogen (1 g/l and 0.1 g/l) and phosphorus (0.05 g/l and 0.1 g/l) limited conditions. Among various N and P-limited conditions, the highest lipid content (60.34 ± 0.69%) obtained under synchronized limitation of N and P (0.1 g/l N and 0.05 g/l P). Live fluorescent cell imaging of yeast cells after BODIPY505-515 nm staining endorses the results of triacylglycerol (TAG) accumulation in lipid droplets. The cells grown in synchronized limitation of N and P exhibited boosted cell size (6.76 ± 0.39 µm) and lipid droplet size (5.62 ± 0.28 µm). Under synchronized limitation of N and P, supersized irregular shaped lipid droplets (LD) coalesced to form big lobules in the cellular compartment of oleaginous yeast having 87.14% enhanced TAG accumulation as depicted by TLC. Synchronized nutrient limitation diverts the CDP-DG pathway of phospholipids synthesis towards de novo TAG synthesis. The maximum increment of oleic acid (C18:1) was reported in synchronized limitation of N and P that improve the biodiesel properties like oxidative stability, viscosity, cetane number and cold filter plugging point.

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1. Introduction

World research scenario has been focused on sustainable biofuel production due to hike in fossil fuel prices and critical environmental disputes on climate change [1]. Biodiesel is emerging as a major alternative to fossil fuels [2]. Its renewable, sustainable and eco-friendly nature make it a perfect vehicular fuel for polluted metro cities [3]. Production of biodiesel is totally reliant on the availability of triacylglycerol feedstocks such as vegetable oils (edible or non-edible), waste cooking oils and animal fats [4]. Its production from oils obtained from vegetable and food sources are no more sustainable and economical due to universal food securities reasons and it is necessary to search for novel resources to produce biodiesel. Microbial oils obtained from yeasts, bacteria, fungi, and algae could be a technologically feasible choice for biodiesel production as they have higher lipid yield under controlled environmental conditions than the plants and can be easily scaled up [5–7]. Only a minor population of yeasts are known to synthesize more than 20% of total lipid in its cellular compartment in the form of lipid droplets (LDs) after expoliating organic carbon sources [8]. *Rhodosporidium, Rhodotorula, Yarrowia, Cryptococcus, Candida, Lipomyces and Trichosporon* are some of the oleaginous yeast genera which accumulate more than 65% of lipid in their cellular compartment. Among these oleaginous yeast genera, *Rhodosporidium* has been recorded to produce the highest amount of lipid (67.5%) in its cellular compartment [6]. Oleaginous yeast has the unique property to grow well and utilize a large number of waste products from industries and agricultural residues for the synthesis of storage lipids (triacylglycerols) in its cellular compartment [1,9,10].

Industrial production of biodiesel still faces hurdles as it is totally dependent on the feedstock used and downstream of the fermentation process. The high costs associated with the biodiesel production are mainly due to the cost of raw materials and their fuel conversion routes. The cost of raw material imparts 60–80% of the total cost of biodiesel production, however, it can be reduced if glucose based renewable substrates (sugarcane bagasse or waste products) are used as carbon source [11]. In order to reduce the overall production cost, an alternative to conventional solvent
Biodiesel production from yeast: An overview of the recent advances.

Rhodosporidium toruloides DSM 4444 was grown in presence of various nutrients and the nutritional composition of the cultivating medium significantly increased the lipid content by 44.9% [21]. Altering the nutritional composition of the cultivating medium also controls the FAME composition. Recently an oleaginous yeast Rhodosporidium toruloides strain Y4 that showed 62.2% lipid accumulation, Wu et al. performed an experiment on oleaginous yeast strain Rhodospiridium toruloides Y4 that showed 62.2% of lipid content accumulation under high C/P molar ratio of 9552 along with low C/N molar ratio of 6 [26]. The p-limitation strategy seems to be very useful for microorganisms where the feedstocks are full of nitrogen sources. Although the effect of N and P starvation alone in the medium has been studied in much detail by several authors, little is known about the effect of both N and P deficiency together and therefore its role in lipid metabolism of R. rarochitovae has been deciphered in this study. The lipid content of synchronized N and P limitation was (60.34± 0.69%, w/w) higher than N and P limitation alone. A significant question addressed in this study was the morphological variations in the structure of yeast lipid droplets (LDs) in response to changes in culture conditions, especially to the cultivation of cells in double limitation of nutrients (N and P), LDs of various sizes have been observed in different conditions (N and P limitations). During the last decade, the research on lipid droplets increases tremendously due to their role as lipid buffer, oil storage and their involvement in human diseases. However, we are only at the beginning to understand metabolic functions which are attributed to this compartment. This study extends our recent knowledge about the molecular mechanism of changes in structure and size of lipid droplets based on the medium or nutritional requirement of the cell. The maximum lipid content (60.34± 0.69%, w/w) was observed in the condition where both essential nutrients (N and P) were in limited condition, further the mechanism of lipid accumulation in P-limited condition has been deciphered by TLC densitometer that showed the decrease in phospholipids content. Biodiesel obtained after down streaming processing were rich in mono-unsaturated fatty acids (MUFAs) and low in polyunsaturated fatty acids (PUFAs) which provide greater performance at low temperature (cold flow plugging properties) and high oxidative stability. Moreover, our new strategy can be utilized for biodiesel production from low N and P containing feedstocks in their natural environmental condition like Cassia fistula L. fruit pulp containing 0.329 g/l N, and 0.0078 ± 0.43 g/l phosphorus [31], pulp and paper industry effluent containing 0.187 ± 29 g/l N and 0.05 g/l P [32]. Jerusalem
2. Materials and methods

2.1. Microorganism and cultivating condition

Earlier isolated red oleaginous yeast *R. kratochvilovae* HIMPA1 (Gene Bank Acc. No. KF772881) was used in this study to explore the effect of dual nutrient limitation of N and P on the growth and lipid accumulation [35]. The culture was maintained on YPD agar (2%) plates and subculture once in a month. For the seed culture, yeast was grown in 50 ml YEPD broth at 30 °C for 48 h and the cells were harvested followed by the washing with sterilized distilled water to remove the media components. The cells were suspended in 0.9% sterilized saline to attain cell density of 6.5–7.8 × 10^9 cells/ml.

2.2. Preparation of nutrients limited media for lipid production

The glucose synthetic media (GSM) used in this experiment as control contains (g/l): Glucose, 70; Salt solution: KH₂PO₄, 1; MgSO₄, 0.5; (NH₄)₂SO₄, 5.0; CaCl₂, 0.1; Trace elements (mg/l): boric acid, 0.5; CuSO₄, 0.04; KI, 0.1; FeCl₃, 0.2; MnSO₄, 0.4; NaMoO₄, 0.2; ZnSO₄, 0.4; Vitamin solution (mg/l): D-Biotin, 0.002; calcium pantothenate, 0.4; folic acid 0.002; inositol 0.20; niacin, 0.4; PABA, 0.2; pyridoxine HCl 0.4; riboflavin 0.2; thiamine, 0.4. For the optimization of nitrogen and phosphorus content in the glucose synthetic medium, spot test analysis was conducted using various agar plates containing different concentration of nitrogen sources such as Urea, NH₄Cl, NH₄NO₃, (NH₄)₂SO₄ (1, 5, 10 and 50 g/l) and phosphorus sources such as KH₂PO₄, Na₂HPO₄, KHPO₄, Na₂HPO₄ (1.5 and 10 g/l). All of these plates were supplemented with complete supplement mixture (CSM; 1.7 g/l) and YNB (0.79 g/l). The next day we made a serial dilution of fresh yeast cells in order to reach, 1 × 10⁻⁴, 1 × 10⁻³, 1 × 10⁻², 1 × 10⁻¹ cell/ml with Bürker chamber. From each one of these dilutions the yeast cell, we spotted 10 μl drops on plates containing the different growth media with different carbon, nitrogen, and phosphate sources at the indicated concentration. We serially plated 1 × 10⁻⁴, 1 × 10⁻³, 1 × 10⁻², 1 × 10⁻¹ cells per spot. To study the effects of nitrogen and phosphorus under limiting condition on the growth and lipid synthesis in cellular compartment of *R. kratochvilovae* HIMPA1, (NH₄)₂SO₄ (showing best growth) concentrations were reduced from 5.0 g/l to 0.1 g/l and KH₂PO₄ from 1 g/l to 0.05 g/l, respectively in distinct experiments. To investigate the combined effect of both nutrients limitation, the nitrogen and phosphorus concentration were 0.1 g/l and 0.05 g/l, respectively. Batch cultivations were performed in 500 ml Erlenmeyer flasks at 30 °C for 240 h with 180 rpm. The glucose content and amount of nitrogen and phosphorus in the medium of batch cultivation experiments were estimated by using the protocol of Patel et al. [35].

2.3. Estimation of cell dry weight and lipid content

In order to estimate the cell dry weight, 50 ml culture was harvested by centrifugation and the obtained pellets were kept on preweighed filter paper. The filter paper along with biomass was oven dried overnight at 60 °C and weighed using an analytical balance. The cell dry weight of *R. kratochvilovae* HIMPA1 was expressed g/l. For the lipid content (% w/w), the harvested cells were used for lipid extraction by the protocol described earlier in Patel et al., 2014 [35]. Lipid yield (g/l) was determined by gravimetric analysis of extracted lipids from the cellular biomass while lipid content was calculated by the following equation—

\[
\text{Lipid content(%, w/w)} = \frac{\text{Lipid yield(g/l)}}{\text{Cell dry weight(g/l)}} \times 100
\]

2.4. Analysis of morphological variation and TAG accumulation in *R. kratochvilovae* HIMPA1

The morphological variation during various nutrient limitations and TAG accumulation in the form of LDs in cellular compartments of *R. kratochvilovae* HIMPA1 were observed by live fluorescence microscopy protocol described in Patel et al. [35]. In brief, 1 ml of cell culture was harvested by centrifugation at 5000 rpm for 5 min and washed with saline water three times to remove media components followed by adding the 2 μl of BODIPY 555/590 nm (0.1 mg/ml DMSO) and incubated for 5 min at room temperature. The fluorescence microscopy was carried out using a digital inverted fluorescence microscope (EVOS- FL, AMG, USA) equipped with EVOS light cube GFP. The morphological variations and lipid droplets accumulation were examined by ImageJ 1.48a software with the help of collected images having statistically significant data.

2.5. Lipid composition analysis by thin layer chromatography (TLC)

Compositional analysis of the extracted lipids were carried out by TLC using 0.25-mm-thick silica gel G-60 F254 plates (Merck, India) and chromatograms were developed by double development method of TLC [36]. The plate was initially developed to 2.5 cm from the origin with hexane: tert-butyl methyl ether: acetic acid (50:50:0.5, v/v/v), and after air dried, redeveloped to 8 cm from the origin with hexane: tert-butyl methyl ether: acetic acid (97:3:0.5, v/v/v) as a developing agent. The samples were visualized by spraying sulphuric acid 50% (w/w) and then heating at 135 °C. After staining, the plate was scanned with Canon Scanner at a fix resolution of 300 dpi and obtained image was processed with Image J 4.18a software for measurement of optical density in the areas of different spots.

2.6. FAME analysis by TLC

Extracted lipids were converted into fatty acid methyl esters (FAME) with transesterification reaction by using the protocol of Patel et al. [35]. Transesterified samples were loaded on the Silica gel TLC-plate (Silica gel 60 F₂₅₄, Merck, Germany) [37]. The standards for triacylglycerol (Triolein) and FAME (Palmitic acid methyl esters) were spotted in lane 1. The development was carried out in TLC chamber with 15% ethyl acetate in hexane as solvent. After drying, the plates were further developed in water for homogeneous saturation. The plate was immersed in 100 ml of 1% solution of KMnO₄ in 4% NaOH for 20 s followed by washing with water for 3–4 times. After staining, the plate was scanned with Canon Scanner at a fix resolution of 300 dpi and obtained image is processed with Image J 4.18a software for measurement of optical density in the areas of different spots.

2.7. Determination of fatty acid methyl ester composition by GC-MS analysis and biodiesel properties

FAME samples were analyzed by GC-MS (Agilent, Santa Clara, CA, USA) equipped with the capillary column (DB- 5MS; dimensions 30 m × 0.25 mm ID and 0.25 μm film thicknesses). The column was initially set for 1.5 min at 50 °C and the temperature was increased to 180 °C for 1 min with the rate of 25 °C/min. It was further increased to 280 °C for 1 min with the rate of 10 °C/min. Finally, the temperature was ramped to 250 °C for 3 min with the...
rate of 15 °C/min. The sample (1 μl) was injected in splitless mode at 250 °C. All the samples were analyzed with the help of electron ionization (70 eV) in scan mode (50–600 m/z). Library search report and retention time were used to identify FAME presence in TAG-derived from nutrient stressed Rhodosporidium kratochvilovae HIMPA1. The empirical formulas were derived for estimation of biodiesel properties such as SV (saponification value; mg KOH), IV (iodine value; gI2/100 g), CN (cetane number), DU (degree of unsaturation; % wt), (LCSF long chain saturation factor; % wt), CFPP (cold filter plugging point; °C), HHV (high heating value; MJ Kg–1), KV (kinematic viscosity, μ; 40 °C in mm2 s–1), density (ρ; 20 °C in g cm–3) and OS (oxidative stability; h) [38–41].

SV = \sum 560(\% FC)/M

IV = \sum 254 DB \times \%FC/M

CN = 46.3 + 5458/SV - (0.255 \times IV)

DU(%) = MUFA + (2 \times PUFA)

LCSF = (0.1 \times C16) + (0.5 \times C18)

CFPP = (3.417 \times LCSF) - 16.477

HHV = 49.43 - 0.041(SV) - 0.015 (IV)

ln(KV) = -12.503 + 2.496 \times ln(\sum M) - 0.178 \times \sum DB

Density = 0.8463 + 4.9/\sum M + 0.0118 \times \sum DB

OS = 117.9295/(wt \% C18 : 2 + wt \% C 18 : 3) + 2.5905

where M = molecular mass of each fatty acid component, DB = number of double bonds, FC = % of each fatty acid component, MUFA = weight % of monounsaturated fatty acids, PUFA = weight % of poly unsaturated fatty acid.

2.8. Experimental determination of SV and IV

The saponification value (SV) of the oleaginous yeast oil was determined by AOCS method [42]. In brief, 1 g yeast oil sample was mixed in 12.5 ml of 0.5 N ethanolic potassium hydroxide. The oil droplets were dissolved in the ethanolic KOH after refluxing of mixture for 20 min. The mixture was cooled at room temperature (25 °C) followed by addition of Phenolphthalein indicator. The obtained solution was titrated with 0.5 N HCl until the pink color disappears. A blank titration was also carried out in the same manner except that no oil was added. It was calculated using the formula:

SV = \frac{56.1 (c - t) \times N}{W}

where, c = Volume (ml) of 0.5 mol/l HCl consumed in the blank test, t = Volume (ml) of 0.5 mol/l HCl consumed in the test, N = Normality of HCl, W = Weight of oil sample.

The Iodine Value (IV) of the oleaginous yeast oil was determined by AOCS method [42]. 0.1 g oil sample was mixed with 20 ml carbon tetrachloride in a conical flask and sealed with stopper. The mixture was subjected to sonication in a sonicator bath followed by addition of 25 ml Hanus solution. After vortexing, the mixture was kept in dark at 20 °C for 30 min. 15% potassium iodide (10 ml) and 100 ml water were added into mixture, sealed and vortex for 30 s. The mixture was titrated with 0.1 mol/l sodium thiosulfate to obtain iodine value. Similarly, blank test was performed to obtain blank level.

IV = \frac{127(\text{c} - \text{t}) \times \text{N}}{10\text{W}}

where; c = Volume (ml) of 0.1 mol/l sodium thiosulfate consumed in the blank test, t = Volume (ml) of 0.5 mol/l sodium thiosulfate consumed in the test, N = Normality of sodium thiosulfate, W = Weight of sample.

2.9. Statistical analysis

The data values are means ± standard deviation of three independent recorded values. One-way analysis of variance (ANOVA) using Microsoft Office Excel 2013 (Microsoft, USA) with p < 0.05 was used for data acceptance.

3. Results and discussion

3.1. Effect of synchronized nutritional stress conditions on the growth and lipid accumulation of Rhodosporidium kratochvilovae HIMPA1

Lipid accumulation in the cellular compartment of oleaginous microorganisms is totally dependent on the metabolic accessibility of provided nutrients. Oleaginous yeasts showed enhanced lipid accumulation while grown in a condition of nutrient limitation with excess of carbon sources [43]. It has been previously reported that medium components such as carbon, nitrogen, phosphorus, sulfur and their ratios (C/N, C/P, C/S) revealed significant influence on the growth and lipid accumulation of oleaginous yeasts [26,43,44]. The synthetic medium used in this study for the cultivation of Rhodosporidium kratochvilovae HIMPA1 was also optimized for maximum biomass production using various nitrogen (Urea, NH4Cl, (NH4)2SO4, NH4NO3) and phosphorous (KH2PO4, NaH2PO4, K2HPO4, Na2HPO4) sources by implying spot test method (Supplementary data Fig. 1). It was observed that among the different sources tested, the growth on (NH4)2SO4 at 5 g/l for nitrogen (Supplementary data Fig. 1) and KH2PO4 at 1 g/l for P were optimum for maximum biomass formation under normal conditions in glucose synthetic medium (GSM). The present study deals with the effect of nutrients (nitrogen and phosphorus) limitation alone and in synchronized manner on the growth and lipid accumulation of oleaginous yeast R. kratochvilovae HIMPA1 and compared with the conditions of optimized nutrients in GSM used as control (Table 1). The other Rhodosporidium species grown on different carbon sources are also compared for their cell dry weight (g/l), total lipid yield (g/l) and lipid content (% w/w) (Table 2). The batch cultivation experiments for cell dry weight, total lipid yield, lipid content, residual glucose, nitrogen and phosphorus under nutritional stress conditions in R. kratochvilovae HIMPA1 is presented in Fig. 1. The carbon source for each batch experiments was glucose (70 g/l or 7%; w/v). The cell dry weight was reported to decrease constantly with the depletion of nitrogen concentration in the medium from 5 g/l to 0.1 g/l. The highest cell dry weight (13.26 g/l) obtained when cells were grown in 5 g/l nitrogen while at 0.1 g/l N the cell dry weight was minimum (9.23 g/l). R. kratochvilovae HIMPA1 cells showed reduced cellular metabolic activity (as measured by XTT assay) and number of cell divisions under nitrogen limited condition (data not shown). However, it was interesting to note that the cells exhibit lipid accumulation in the
opposite manner regardless of their growth. The highest lipid content (59.69%) was observed with the nitrogen (0.1 g/l) limited condition. The TLC chromatogram of extracted lipids from oleaginous yeast grown under various nutrient stress conditions is presented in Fig. 2A. Quantitative measurement of optical density in the areas of different spots is presented in Fig. 2B. The amount of TAG, FFA, St, PE and PC (% w/w) in obtained total lipid (g/l) were calculated according to their corresponding area represented in

Table 1
Comparative study of cell dry weight (g/l), total lipid yield (g/l) and lipid content (% w/w) of R. kratochvilovae HIMPA1 grown in various nutrient-limited conditions and GSM.

<table>
<thead>
<tr>
<th>Cultivation medium containing (N and P)</th>
<th>Cell dry weight (g/l)</th>
<th>Total lipid yield (g/l)</th>
<th>Lipid content (% w/w)</th>
<th>Total cell dry weight (g/g)</th>
<th>Lipid yield (g/g)</th>
</tr>
</thead>
<tbody>
<tr>
<td>5 g/l N; 1 g/l P (control)</td>
<td>13.26 ± 0.98</td>
<td>6.2 ± 0.58</td>
<td>46.76 ± 0.61</td>
<td>0.189</td>
<td>0.088</td>
</tr>
<tr>
<td>1 g/l N; 1 g/l P</td>
<td>12.17 ± 0.34</td>
<td>5.92 ± 0.76</td>
<td>48.64 ± 0.45</td>
<td>0.173</td>
<td>0.084</td>
</tr>
<tr>
<td>0.1 g/l N; 1 g/l P</td>
<td>9.23 ± 0.65</td>
<td>5.51 ± 0.32</td>
<td>59.69 ± 0.65</td>
<td>0.131</td>
<td>0.078</td>
</tr>
<tr>
<td>5 g/l N; 0.1 g/l P</td>
<td>12.89 ± 0.37</td>
<td>6.32 ± 0.45</td>
<td>48.33 ± 0.53</td>
<td>0.184</td>
<td>0.090</td>
</tr>
<tr>
<td>5 g/l N; 0.05 g/l P</td>
<td>12.65 ± 0.12</td>
<td>6.54 ± 0.87</td>
<td>51.7 ± 0.81</td>
<td>0.180</td>
<td>0.093</td>
</tr>
<tr>
<td>0.1 g/l N; 0.05 g/l P</td>
<td>7.54 ± 0.65</td>
<td>6.45 ± 0.54</td>
<td>60.34 ± 0.69</td>
<td>0.117</td>
<td>0.066</td>
</tr>
</tbody>
</table>

*a g of total cell dry biomass produced per g of substrate consumed.

*b g of cellular lipids produced per g of substrate consumed.

Table 2
Rhodosporidium species grown on different carbon sources for lipid production.

<table>
<thead>
<tr>
<th>S. No.</th>
<th>Oleaginous Yeasts</th>
<th>Medium/mode of cultivation</th>
<th>Dry cell weight (g/l)</th>
<th>Total lipid yield (g/l)</th>
<th>Lipid content (% w/w)</th>
<th>References</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Rhodosporidium toruloides 21167</td>
<td>Cassava starch/2-l co-culture bioreactor</td>
<td>20.1</td>
<td>13.04</td>
<td>64.9</td>
<td>[69]</td>
</tr>
<tr>
<td>2</td>
<td>Rhodosporidium toruloides 2F5</td>
<td>Inulin</td>
<td>15.82</td>
<td>9.83</td>
<td>62.14</td>
<td>[70]</td>
</tr>
<tr>
<td>3</td>
<td>Rhodosporidium fluviale DMKU-RK253</td>
<td>Crude glycerol-YM (yeast extract- malt extract) medium/5 L jar fermenter</td>
<td>14.09</td>
<td>8.99</td>
<td>63.8</td>
<td>[71]</td>
</tr>
<tr>
<td>4</td>
<td>Rhodosporidium toruloides AS 2.1389</td>
<td>Acetic acid/carbon-to-nitrogen ratio (C/N) of 200 in a batch culture</td>
<td>4.35</td>
<td>2.1</td>
<td>48.2</td>
<td>[72]</td>
</tr>
<tr>
<td>5</td>
<td>Rhodosporidium diobovatum (08–225)</td>
<td>Pinewood pyrolytic sugars 100%</td>
<td>17.53</td>
<td>4.08</td>
<td>23.27</td>
<td>[73]</td>
</tr>
<tr>
<td>6</td>
<td>Rhodosporidium fluviale DMKU-SP314</td>
<td>Glucose and xylose</td>
<td>14.3</td>
<td>7.9</td>
<td>55.24</td>
<td>[74]</td>
</tr>
<tr>
<td>7</td>
<td>Rhodosporidium toruloides DEBB 5533</td>
<td>Sugarcane juice/batch cultivation (1000 L working volume)</td>
<td>24.5</td>
<td>10.1</td>
<td>41.22</td>
<td>[18]</td>
</tr>
<tr>
<td>8</td>
<td>Rhodosporidium toruloides ATCC 10788</td>
<td>Crude glycerol media.</td>
<td>21.16</td>
<td>11.27</td>
<td>53.28</td>
<td>[75]</td>
</tr>
<tr>
<td>9</td>
<td>R. kratochvilovae HIMPA1</td>
<td>Aqueous extract of Cassia fistula L. (CAE) fruit pulp</td>
<td>8.9</td>
<td>4.86</td>
<td>53.18</td>
<td>[31]</td>
</tr>
<tr>
<td>10</td>
<td>R. kratochvilovae HIMPA1</td>
<td>Pulp and paper industry effluent</td>
<td>13.87</td>
<td>8.56</td>
<td>61.71</td>
<td>[32]</td>
</tr>
<tr>
<td>11</td>
<td>R. kratochvilovae HIMPA1</td>
<td>GSM containing 5 g/l N; 1 g/l P (control)</td>
<td>13.26</td>
<td>6.2</td>
<td>45.76</td>
<td>This study</td>
</tr>
<tr>
<td>12</td>
<td>R. kratochvilovae HIMPA1</td>
<td>GSM containing 0.1 g/l N; 1 g/l P</td>
<td>9.23</td>
<td>5.51</td>
<td>59.69</td>
<td>This study</td>
</tr>
<tr>
<td>13</td>
<td>R. kratochvilovae HIMPA1</td>
<td>GSM containing 5 g/l N; 0.05 g/l P</td>
<td>12.65</td>
<td>6.54</td>
<td>51.7</td>
<td>This study</td>
</tr>
<tr>
<td>14</td>
<td>R. kratochvilovae HIMPA1</td>
<td>GSM containing 0.1 g/l N + 0.05 g/l P</td>
<td>7.54</td>
<td>4.65</td>
<td>60.34</td>
<td>This study</td>
</tr>
</tbody>
</table>
get converted into lipid droplets in cellular compartment [28]. However, in the case of oleaginous yeast, the excess carbon source the amount of mannans and glucans contents increases. but when they grow in nitrogen-limited medium with excess car-

Fig. 2. A. Separation of extracted lipids from R. kratochvilovae HIMPA1 grown in various nutritional starved conditions on a TLC-plate. Lane 1, a standard mixture of TAG (triolein), PC, sterol (St), FFA, phosphatidylethanolamine; lane 2, 5 g/l N and 1 g/l P; lane 3, 0.5 g/l N and 1 g/l P; lane 4, 0.1 g/l N and 1 g/l P; lane 5, 5 g/l N and 0.1 g/l P; lane 6, 5 g/l N and 0.05 g/l P; lane 7, synchronized limitation of N (0.1 g/l) and P (0.05 g/l). B. Optical densities of the individual spots were determined within the same lane, sequentially applied to all lanes. Areas of the spots plotted as the functions of the corresponding masses.

Fig. 2B (Table 3). The data of lane 3 and 4 depict increased TAG content in LDs (31.77% and 64.83%) in N-limited condition (1 g/l and 0.1 g/l) respectively as compared to normal (without limitation, lane 2) while the scanty amount of FFA, PE and PC were present. These results validate the enhanced TAG accumulation in N-limited condition. Similar results was shown by Li et al. where R. toruloides Y4 accumulated more than 70% lipid content of their dry biomass in N-limited condition (C/N molar ratios higher than 570) however, in the lower C/N molar ratios (<100), lipid contents were decreased to 20% [45]. Mortierella isabellina, an oleaginous fungus observed to accumulate more than 50% lipid content under the high (150—300) C/N molar ratios [46]. Hassan et al. showed that oleaginous yeast Cryptococcus curvatus accumulate more lipid content with stearic acid (14%) under the N-limited condition, whereas, it was boosted by the combined effect of Fe and N-limited condition (19%) [47]. Brauer et al. described the N regulation in yeast cells where the cells had reduced ribosomal synthesis and translation, that results in reduced ribosomal synthesis and translation, that results in slow growth and enlargement of the G1 phase of the cell cycle under N-limited conditions [48]. Non-oleaginous yeast such as Saccharomyces cerevisiae and food yeast (Candida ultilis) cannot accumulate lipid content more than 10% of their total biomass [49], but when they grow in nitrogen-limited medium with excess carbon source the amount of mannans and glucans contents increases. However, in the case of oleaginous yeast, the excess carbon source get converted into lipid droplets in cellular compartment [28]. When nitrogen exhausts from the culture medium, Adenosine monophosphate deaminase gets activated and catalyzes the reaction of AMP to Inosine 5’-monophosphate (IMP) and Ammonium [43]. The Adenosine monophosphate deaminase enzyme is present in oleaginous yeast but no such absolute dependency in non-oleaginous yeast occurs [10]. Isocitrate dehydrogenase (ICDH) is inactivated when the concentration of AMP decreases and Iso-

Table 3

<table>
<thead>
<tr>
<th>S. No.</th>
<th>Cultivation medium containing (N and P)</th>
<th>Total lipid yield (g/l)</th>
<th>TAG (% w/w)</th>
<th>FFA (% w/w)</th>
<th>St (% w/w)</th>
<th>PE (% w/w)</th>
<th>PC (% w/w)</th>
</tr>
</thead>
<tbody>
<tr>
<td>a</td>
<td>5 g/l N; 1 g/l P (control)</td>
<td>6.2 ± 0.58</td>
<td>20.46</td>
<td>11.97</td>
<td>9.06</td>
<td>19.32</td>
<td>39.19</td>
</tr>
<tr>
<td>b</td>
<td>1 g/l N; 1 g/l P</td>
<td>5.92 ± 0.76</td>
<td>30.51</td>
<td>18.88</td>
<td>13.65</td>
<td>11.20</td>
<td>25.76</td>
</tr>
<tr>
<td>c</td>
<td>0.1 g/l N; 1 g/l P</td>
<td>5.51 ± 0.32</td>
<td>45.53</td>
<td>23.45</td>
<td>8.03</td>
<td>9.76</td>
<td>13.23</td>
</tr>
<tr>
<td>d</td>
<td>5 g/l N; 0.1 g/l P</td>
<td>6.32 ± 0.45</td>
<td>62.26</td>
<td>0.90</td>
<td>21.17</td>
<td>13.46</td>
<td>2.21</td>
</tr>
<tr>
<td>e</td>
<td>5 g/l N; 0.05 g/l P</td>
<td>6.54 ± 0.87</td>
<td>98.05</td>
<td>0.68</td>
<td>0.69</td>
<td>0.19</td>
<td>0.39</td>
</tr>
<tr>
<td>f</td>
<td>0.1 g/l N; 0.05 g/l P</td>
<td>4.65 ± 0.54</td>
<td>87.46</td>
<td>12.14</td>
<td>nd</td>
<td>nd</td>
<td>0.4</td>
</tr>
</tbody>
</table>

nd: not detected.
The studies on the biosynthesis of intracellular carbohydrate and its relation to intracellular lipid synthesis in Cryptococcus curvatus NRRL Y-1511 have opened the fresh area of research for metabolites synthesis. This oleaginous yeast synthesized high quantities of intracellular carbohydrate (up to 68% w/w) at the starting of fermentation process when grown in lactose or sucrose under nitrogen limiting condition while the amount was decreased to 20%, w/w at the end of the process. Complementary to the decline in carbohydrate level led to increase in lipid content under continuous intake of extracellular lactose. However, this situation was different in the presence of excess quantities of nitrogen in the medium higher quantities of intracellular carbohydrate was produced [50]. Boosted lipid accumulation is also reported in oleaginous yeast cells under N-limited condition; however, it is difficult to attain high cell dry weight in such condition.

Another target for lipid accumulation in R. kratochvilovae HIMPA1 was phosphorus limitation in the medium that was effective irrespective of the presence of high amount of nitrogen in the medium. For the growth and lipid accumulation in oleaginous yeast cells, not only nitrogen but also phosphorus is an important factor that is assimilated for phospholipids, nucleic acid, and coenzymes formation in yeast cells. Additionally, phosphorus has the main role in the generation of metabolic energy in the form of ATP and NADP [51]. Batch cultivation experiments for phosphorus-limited conditions (0.1 and 0.005 g/l) without altering the concentration of nitrogen (5 g/l) was represented in Fig. 1. The cell dry weight of R. kratochvilovae HIMPA1 was reported to be almost constant (6.32 ± 0.45 g/l and 6.54 ± 0.87 g/l) for both phosphorus starved conditions (0.1 and 0.05 g/l). Whereas R. kratochvilovae HIMPA1 was grown in the medium containing 0.1 g/l P, the cell dry weight, and lipid yield were 12.89 g/l and 6.32 g/l respectively while at 0.05 g/l P, the cell dry weight was 12.65 g/l along with 6.54 g/l lipid yield. Moreover, the lipid content has increased from 48.33% to 51.7% with the decreasing amount of P (Table 1). The results obtained from TLC of extracted lipids in lane 5 and lane 6 specifically showed 34.70% and 78.55% increased TAG contents of LDs in phosphorus starved conditions (0.1 and 0.05 g/l) than the normal condition (lane 2) while the content of FFA, PE (35.47%) and PC (56.81%) were reduced (Fig. 2B).

These results prove our hypothesis that amount of phospholipids were reduced under P-starved condition (0.05 g/l P) while the amount of TAG was increased due to the diversion of CDP-DG pathway for phospholipids synthesis towards de novo TAG biosynthesis pathway as shown in Fig. 3 and confirmed by TLC densitometry. It has also been studied previously that the content of TAG was increased from 6.5% up to 39.3% of total lipids while the phospholipid content was reduced in the phosphate-starved Arabidopsis thaliana cells whereas the synthesis of galactolipids compensates for the decrease in phosphatidylcholine (PC) under phosphate limited condition [29]. Earlier, it was reported that phosphate limitation causes the change of membrane phospholipids into non-phosphorus betaine lipids in photosynthetic bacteria, Rhodobacter sphaeroides [30] and, thus, represents an effective phosphate-conserving mechanism. The strategy to increase the lipid accumulation under P limitation is not only compatible with yeast species but also, several algae species also accumulate high lipid content under P limitation (high C/P ratio in medium) e.g. Isochrysis galbana U4 accumulated almost 50% lipid content under P-starved condition [52].

The other oleaginous yeast Rhodosporidium toruloides Y4 also produced higher amount of lipids under P-limited condition in the growth medium [45]. Under the phosphate-limited condition, C. utilis synthesized more intracellular non-polar lipid content but...
polar lipid remained relatively constant [53]. It has been suggested already that under the N-limited condition, the batch culture of R. glutinis showed meager growth in wet biomass and totally inhibits cell divisions on N consumption [54]. However, the condition was completely different under phosphate exhaustion condition where the cells showed exponential growth with increased lipid-free biomass.

Interestingly, the medium containing synchronized nitrogen and phosphorus limitations had 7.54 ± 0.54 g/l cell dry weight and 4.65 ± 0.67 g/l lipid yield along with 13–14% higher lipid content (60.34± 0.69%) than grown under the single limitation of N or P (Table 1). TLC of synchronized nutrient limitations showed highest amount of TAG (87.14% in comparison with normal condition of GSM) with no PE and reduced amount of PC (98.36%) in lane 7 (Fig. 2B). Sterols (St) were also absent which enhance the purity of biodiesel produced. It suggests that dual limitation of N and P causes diversion of metabolic flux from CDP-DG pathway of phospholipids (PI, PC and PE) synthesis toward that of TAG biosynthesis. This will ease TAG production without even creating mutants or inhibitors in the specific pathway to boost de novo TAG biosynthesis. These results are also comparable with the recent research work done by Bellou et al. on Y. lipolytica grown under the double limitation of nitrogen and magnesium [55]. The double limitation of nitrogen and magnesium support the lipid accumulation (47.5% w/w) in this yeast and synthesize 12.2 g/l cell dry weight and 5.8 g/l of nitrogen and magnesium supports the lipid accumulation (47.5% w/w) in this yeast and synthesize 12.2 g/l cell dry weight and 5.8 g/l lipid yield respectively when glucose was provided as carbon substrate [55]. Furthermore, the extension of cultivation period after glucose depletion from the medium was carried out to examine the effect of nutrient limitation on lipid biodegradation. In the case of N limitation (1 g/l), the lipid content was reported to slightly decrease from 48.64% to 48.32% when culture was cultivated from 144 h to 240 h, while at 0.1 g/l N the lipid content was decreased from 59.69% to 58.69% (Fig. 1). These results showed that N limitation promotes lipid degradation after glucose depletion from the medium. Interestingly, the lipid content was constant at both P-starved condition (0.1 g/l and 0.05 g/l) after the extended growth period. While under synchronized limitation of N and P, the biodegradation of lipid was observed and the lipid content was slightly changed from 61.71% to 61.36% after glucose exhaustion from the medium (from 168 h to 240 h).

3.2. Morphological changes during the nitrogen and phosphorus limitation

The TAG accumulation in N and P-limited conditions were evaluated by morphological and structural variations in lipid droplet (LD) formation of yeast cellular compartment. The results of average variation in cell size and lipid droplets were statistically verified and presented in Fig. 4A and B. Maximum lipid droplets size (5.62 ± 0.28 μm) in R. kratochvilovae HIMPA1 were observed when cells were grown under combination of both N and P limitations in the medium, which is comparable to lipid droplets size of N limitation (1 g/l N; 4.32 ± 0.64 μm and 0.1 g/l N; 4.68 ± 0.21 μm) and P limitation (0.1 g/l P; 3.91 ± 0.32 μm and 0.05 g/l P; 3.96 ± 0.19 μm) alone provided to the yeast cells. The obtained results showed a direct correlation between lipid droplet size with the TAG accumulating ability of R. kratochvilovae HIMPA1 on consumption of nutritional variation of N and P (Fig. 4A and B) as the dye (BODIPY) stains TAG specifically.

In N-limited condition, the perfectly round LD is surrounded by distinct tough phospholipid layer (Fig. 4B, lane 2 and 3) while it seems to be little diffused and leaky as in lane 4 and 5 (Fig. 4B) of P-limited condition. Furthermore, lack of phospholipid synthesis clearly disrupts the distinct round LD into diffused and irregular shaped LD lacking membrane layer in synchronized limitation of N and P. In this case the TAG storage in round LDs coalesced into an irregular multilobular large mass without a distinct LD cover with enhanced TAG content per unit cell as compared to the normal N and P condition (5 g/l N and 1 g/l P).

TAG is not appropriate for membrane constituents as they are the neutral lipids and serves mainly as an energy source. Biogenesis of LD in the cellular compartment defined by two distinct models ‘flesming model’ and the ‘bicelle formation model’. In flesming model, LD bud off from ER through the help of only outer leaflet while in bicelle formation model, monolayer of phospholipids form over LD with the help of both leaflets of ER. This monolayer contains PC, PI, and PE [56,57]. One more possibility of LD formation is secretory vesicle model in which TAG are filled in secretory vesicles [58] (Fig. 3). The data from the morphological studies of LD in N and P synchronized limitation supports the secretory vesicle model as the LD coalesce into one multilobular mass similar to secretory vesicles coalesce with in outer cell membrane during secretion process.

3.3. Effect of nutritional stress conditions on fatty acid profile and biodiesel properties

Effect of nitrogen and phosphorus limitation on fatty acid profile of R. kratochvilovae HIMPA1 is represented in Table 4. It has been already reported that R. kratochvilovae HIMPA1 grown on control (GSM) containing optimum 5 g/l N and 1 g/l P, synthesized mainly myristic acid (C14:0) 4.24%, palmitic acid (C16:0) 34.79%, stearic acid (C18:0) 21.32%, oleic acid (C18:1) 23.15%, along with 2.23% linoleic acid (C18:2) [32]. The data showed that nitrogen limitation (0.1 g/l N) imparts the increment of mono-unsaturated fatty acids (MUFA) from 23.15± 0.53% to 75.79± 0.64% while the amount of saturated fatty acid (SFA, from 63.35% to 12.43%) were reduced from that obtained with optimum amount of N (5 g/l N) as shown in Table 4. The P limitation affects the fatty acids content in a similar manner. The amount of MUFA was increased from 23.15% to 66.79% while SFA content was decreased from 63.35% to 28.52% when the yeast cells grown in P-starved condition (0.05 g/l P). However, a significant increment in oleic acid (C18:1; 81.52%) was observed when both nutrients (0.1 g/l N and 0.05 g/l P) were in limited condition together. It has been earlier reported that the amount of linoleic acid (C18:2) present in high quantity during the lag phase of a growth period that incorporated in the cell membrane while the amount of stearic acid (C18:0) and oleic acid (C18:1) increased in stationary phase or lipid accumulating phase. Later on, Δ9 desaturase converted C18:0 into C18:1 by adding double-bond [59]. Desaturation in the later stage of growth ensues by an oxidative process that requires oxygen, NADH, NADPH and substrate [60]. This process prevents the accumulation of reactive oxygen species under stress condition. Various researchers previously described that the amount of MUFA was increased by unsaturated fatty acid mutants (USFA) of A. curvatu with mutagenic treatment [61] or by using sterculia oil as Δ9 desaturase inhibitor [62] that block the conversion of stearic acid (C18:0) to oleic acid (C18:1). The fatty acid profile of oleaginous yeast can be altered according to provided medium e.g. Y. lipolytica ACA-DC 50109 showed enhanced intracellular C18:0 when cultivated on saturated fatty substrates and desired cocoa butter substitute was obtained after adding donor of C18:1 as a co-substrate in the provided medium [15]. In another example, this strain synthesizes intracellular fatty acids similar to cocoa butter when grown on mixtures of stearin and biodiesel by-products glycerol [24].

The conversion of total extracted lipid into FAME confirmed complete conversion (97.33%) of TAG by TLC (Fig. 5, lane 3–7) is demonstrated in TLC chromatogram (lane 3–7). However, 91.12% TAG conversion is reported in Fig. 5, lane 2. In the FAME profile, the
Fig. 4. A. Estimation of cell size and lipid droplets size in oleaginous yeast cells stained with BODIPY505/515 nm using digital inverted LED equipped fluorescence microscopy at resolution of 60×. B. Representative images for lipid droplets formation in single cell of R. kratochvilovae HIMPA1 when grown on various N and P starved condition; (1) GSM, 5 g/l N and 1 g/l P (2) 0.5 g/l N and 1 g/l P (3) 0.1 g/l N and 1 g/l P (4) 5 g/l N and 0.1 g/l P (5) 5 g/l N and 0.05 g/l P (6) Synchronized limitation of N (0.1 g/l) and P (0.05 g/l). Images were taken when cells achieved its stationary phase using fluorescence microscopy of live cells stained with BODIPY505/515 nm.
degree of unsaturation and fatty acids chain length are the two main features that determine the fuel properties. The empirical formulas were designed to explain the biodiesel properties of FAME obtained after \textit{R. kratochvilovae} HIMPA1 was grown in various N and P limitation conditions and various important biodiesel properties such cetane number (CN), degree of unsaturation (DU), iodine value (IV), heating value (HHV)/calorific value, saponification value (SV), and oxidative stability (OS) were evaluated as per ASTM D6751 (United States) and EN 14214 (European Union) fuel standards. FAME profile obtained after transesterification of extracted lipid directly influences the quality of biodiesel. The N and P-limited condition affect the fatty acids profiles as under N-limited conditions, high quantity of MUFA (monounsaturated fatty acids) were obtained (1 g/l N, 67.27% and 0.1 g/l N, 75.78%) whereas under P limitation the quantity of MUFA also raised (0.1 g/l P; 48.9% and 0.05 g/l P 66.79%) as compared to normal condition of N and P. Moreover, the combination of both nutrients limitation contributed high quantity of MUFA (0.1 g/l N and 0.05 g/l P; 81.52%) that causes the enhancement of cold flow plugging properties (−10.897 °C). High SFA (saturated fatty acids) present in FAME mitigate the biodiesel to auto-oxidation and thereby increasing its shelf-life while UFA (unsaturated fatty acids) quantities determine its cold flow plugging properties. Therefore, it is necessary to control the fuel properties with the optimum ratio of SFA to UFA. The important property of biodiesel to perform at low temperature is determined by the degree of unsaturation. The data showed that degree of unsaturation was increased from 27.61% to 99.77% in N-starved condition (0.1 g/l N) while 87.67% increment was reported in P-starved condition (0.05 g/l P) \textbf{Table 5}. Fatty acid unsaturation is also depicted by iodine value (IV). Biodiesel obtained from \textit{R. kratochvilovae} HIMPA1 grown under N (0.1 g/l N) and P (0.05 g/l P) limited condition showed 83.1616 and 75.3527 gI$_2$/100 g of iodine value, respectively while in normal condition of N and P (5 g/l and 1 g/l), it was 23.7184 gI$_2$/100 g. The experimental value of IV of different oil samples obtained from oleaginous yeast grown in GSM containing (5 g/l N and 1 g/l P) control; (1 g/l N and 1 g/l P; 0.1 g/l N and 1 g/l P) nitrogen limitation; (5 g/l N and 0.1 g/l P; 5 g/l N and 0.05 g/l P) phosphorus limitation; (0.1 g/l N and 0.05 g/l P) synchronization limitation of N and P were 20.13, 11.29, 74.65, 59.65, 69.43, 78.32 gI$_2$/100 g respectively. European Union standard (EN 14214) has set a limit of 120 g I$_2$/100 g which excludes sunflower, soya bean, grape seed oil as potential feedstocks for biodiesel production. As reported in the literature that high iodine value results in polymerization of glycerides and on heating it has the problem of gum formation [63,64]. Another important biodiesel property the saponification value (SV) is used to measure the

\begin{table}[h]
\centering
\caption{Fatty acid methyl ester profiles of oleaginous yeast (OY) grown in various nutrient limited condition.}
\begin{tabular}{|c|c|c|c|c|c|c|c|c|c|}
\hline
FAME composition (%) & 
\multicolumn{9}{|c|}{Cultivation medium containing (N and P)}
\hline
 & 5 g/l N & 1 g/l P & 1 g/l N & 1 g/l P & 0.1 g/l N & 1 g/l P & 5 g/l N & 0.1 g/l P & 5 g/l N & 0.05 g/l P & 0.1 g/l N & 0.05 g/l P
\hline
SFA* & & & & & & & & & & & & \\
Lauric acid methyl ester (C$_{12}$:0) & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd \\
Myristic acid methyl ester (C$_{14}$:0) & 4.24 ± 0.23 & 0.91 ± 0.43 & nd & nd & 0.19 ± 0.31 & nd & nd & nd & nd & nd & nd \\
Palmitic acid methyl ester (C$_{16}$:0) & 34.79 ± 0.34 & 16.44 ± 0.12 & 9.1 ± 0.13 & 18.24 ± 0.34 & 7.8 ± 0.29 & 8.63 ± 0.27 & nd & nd & nd & nd & nd & nd \\
Margaric acid methyl ester (C$_{17}$:0) & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd \\
Stearic acid methyl ester (C$_{18}$:0) & 21.32 ± 0.11 & 8.37 ± 0.87 & 3.33 ± 0.63 & 17.6 ± 0.54 & 10.36 ± 0.59 & 1.54 ± 0.56 & nd & nd & nd & nd & nd & nd \\
Acetic acid methyl ester (C$_{18}$:0) & 23.15 ± 0.53 & 67.27 ± 0.75 & 75.79 ± 0.64 & 48.9 ± 0.81 & 66.79 ± 0.21 & 81.52 ± 0.53 & nd & nd & nd & nd & nd & nd \\
MUFA** & & & & & & & & & & & & \\
Oleic acid methyl ester (C$_{18}$:1) & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd \\
\hline
PUFA*** & & & & & & & & & & & & \\
Linoel acid methyl ester (C$_{18}$:2) & 2.23 ± 0.76 & 8.38 ± 0.91 & 10.49 ± 0.73 & 13.35 ± 0.61 & 10.48 ± 0.43 & 7.98 ± 0.21 & nd & nd & nd & nd & nd & nd \\
α-Linolenic acid methyl ester (C$_{18}$:3) & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd & nd \\
\hline
\end{tabular}
\end{table}

\begin{figure}[h]
\centering
\includegraphics[width=\textwidth]{image}
\caption{Estimation of TAG conversion into FAME after transesterification of obtained lipid from various nutrient limited condition- \textbf{Lane 1}, a standard mixture of TAG (triolein) and FAME (palmitic acid methyl esters); \textbf{Lane 2}, 5 g/l N and 1 g/l P; \textbf{Lane 3}, 0.5 g/l N and 1 g/l P; \textbf{Lane 4}, 0.1 g/l N and 1 g/l P; \textbf{Lane 5}, 5 g/l N and 0.1 g/l P; \textbf{Lane 6}, 5 g/l N and 0.05 g/l P; \textbf{Lane 7}, synchronized limitation of N (0.1 g/l) and P (0.05 g/l).}
\end{figure}

SFA* Saturated fatty acids.
MUFA** Monounsaturated fatty acids.
PUFA*** Polyunsaturated fatty acids.
molecular weight or chain length of fatty acids and defined by the amount of KOH in mg required to saponify one g of fat under a specific condition [65]. SV of oil obtained from R. kratochvilovae grown under N (0.1 g/l N) and P (0.05 g/l P) starved condition showed high value (190.22 mg KOH and 183.65 mg KOH, respectively) than 160 mg KOH as obtained under optimum condition of (5 g/l) N and (1 g/l) P (Table 5). The experimental value of SV of different oil samples obtained from oleaginous yeast grown in GSM containing (5 g/l N and 1 g/l P) control; (1 g/l N and 1 g/l P; 0.1 g/l N and 1 g/l P) nitrogen limitation; (5 g/l N and 0.1 g/l P; 5 g/l N and 0.05 g/l P) phosphorus limitation; (0.1 g/l N and 0.05 g/l P) synchronized limitation of N and P were 158, 189, 178, 179, 163 and 178 mg KOH respectively. The obtained SV value were practically similar with the value obtained from empirical formula. The flow of any type of fuel in a conventional engine is defined by their density and viscosity. Under N and P-starved condition the kinematic viscosities were 4.60324 mm²/s and 4.60324 mm²/s respectively, while under synchronized nutrient limited condition (0.1 g/l N and 0.05 g/l P) the KV was 4.52329 mm²/s that show similar result under normal condition of (5 g/l N and 1 g/l P). The ASTM D6751 viscosity specification of 1.9–6.0 mm²/s is satisfied by these results and KV majorly affects the injection into the engine because higher viscosity leads to larger droplet sizes, poorer vaporization, and narrower injection spray angle [66]. Density is also an important parameter of biodiesel that is correlated with another parameter for engine performance such as cetane number and heating value. Fuel injection property is totally dependent on its density as higher or lesser volume of fuel is injected. It affects the pumping of fuel by its volume not by mass. It can be defined as the mass of a substrate divided by its volume or weight per unit volume. Denser oil has more energy and biodiesel is usually having higher density than petroleum diesel. The density of the fuel is correlated with other properties such as HHV, viscosity, cetane number which depends on temperature, water content, and the presence of free fatty acid content in FAME. The data obtained for density after synchronized nutrient limitation of N and P showed 0.872103 g/cm³ that was similar with data obtained in the normal condition of N and P (0.872103 g/cm³). Oxidative stability is another essential fuel property which gets influenced by the presence of high amount of UFA in biodiesel. Increasing the number of double bonds in UFA leads to autoxidation. The data showed that without limitations of nutrients in the medium, the biodiesel obtained was more stable (55.4737 h) than the limitations of N (1 g/1 N, 16.6632 h) and 0.1 g/l N, 13.8326 h) and limitations of P (0.1 g/l P, 11.4242 h and 0.05 g/l P, 13.8433 h). Cetane number (CN) is again the significant property of fuel that decides the ignition characteristics of fuel i.e. how any fuel got ignite and combust. It affects the various parameters of engine performance such as noise, emissions of CO and stability. Higher CN imparts the better ignition of biodiesel than the conventional diesel fuel ensuring better cold start behavior, smooth engine run and complete combustion leading to reduced gaseous and particulate emissions [66,67]. The data showed in this study suggest that the medium containing sufficient quantity of N and P (without any limitations) has higher CN (74.3076) than CN obtained after N and P-limited condition (53.6484 and 52.639 respectively). The data also indicate that the biodiesel obtained after normal and nutrient limited condition as listed in Table 1 satisfy the CN limit describe by both ASTM D6751-02 and EN 14214 standard. Cetane number has both its lower and higher limits as lower cetane number of biodiesel causes difficulty of engine starting in cold environmental and generation of noise and pollution (emissions of hydrocarbons) without proper combustion of biodiesel while higher cetane number causes instant ignition without proper mixing of air that results in decrease in efficiency of fuel [39,67]. The heating energy released during the combustion of the unit value of fuels is considered as the heating value (HV) of fuels and it is also known as calorific value or heat of combustion. The elements of fuel such as O₂, H, C, N, and S after burning generates gaseous CO₂, NO, SO₂, and water along with heat. Biodiesel has low energy content due to approximately 12% less HV than that of diesel (39.57–41.33 MJ/kg) and consumption rate in a diesel engine is also high in order to attain similar yield as that of conventional diesel [67,68]. The result concluded that biodiesel obtained after synchronized N and P-starved condition had 40.3748 MJ/kg HHV while the value was high (42.5033 MJ/kg) under normal condition of N and P. From the above discussion, biodiesel properties such as oxidative stability, viscosity, cetane number and Cold filter plugging point; KV- kinematic viscosity; IV- iodine value; SV- saponification value; CN- cetane number; DU- degree of unsaturation; LCSF- long chain saturation factor; HHV- high heating value; CFPP- cold filter plugging point; where IV- iodine value; SV- saponification value; CN- cetane number; DU- degree of unsaturation; LCSF- long chain saturation factor; HHV- high heating value; CFPP- cold filter plugging point; KV- kinematic viscosity; 

<table>
<thead>
<tr>
<th>Quality parameters of biodiesel</th>
<th>5 g/l N and 1 g/l P</th>
<th>1 g/l N and 1 g/l P</th>
<th>0.1 g/l N and 1 g/l P</th>
<th>5 g/l N and 0.1 g/l P</th>
<th>5 g/l N and 0.05 g/l P</th>
<th>0.1 g/l N and 0.05 g/l P</th>
<th>Standard fuel parameters</th>
</tr>
</thead>
<tbody>
<tr>
<td>SV (mg KOH)</td>
<td>160.266</td>
<td>196.806</td>
<td>191.141</td>
<td>191.054</td>
<td>183.65</td>
<td>190.22</td>
<td>–</td>
</tr>
<tr>
<td>IV (g l/100 g)</td>
<td>23.7184</td>
<td>14.4797</td>
<td>83.1616</td>
<td>65.0288</td>
<td>75.3527</td>
<td>83.7415</td>
<td>–</td>
</tr>
<tr>
<td>CN</td>
<td>74.3076</td>
<td>70.3406</td>
<td>53.6486</td>
<td>58.2855</td>
<td>56.8046</td>
<td>53.639</td>
<td>47 min</td>
</tr>
<tr>
<td>DU (% wt)</td>
<td>27.61</td>
<td>16.76</td>
<td>96.77</td>
<td>75.6</td>
<td>87.67</td>
<td>97.48</td>
<td>–</td>
</tr>
<tr>
<td>LCSF (% wt)</td>
<td>14.139</td>
<td>5.649</td>
<td>2.575</td>
<td>1.864</td>
<td>5.96</td>
<td>1.63</td>
<td>–</td>
</tr>
<tr>
<td>HHV (Mg/kg)</td>
<td>42.5033</td>
<td>41.1438</td>
<td>40.3458</td>
<td>40.6214</td>
<td>40.7701</td>
<td>40.3748</td>
<td>–</td>
</tr>
<tr>
<td>KV (mm³/s)</td>
<td>4.52329</td>
<td>4.31376</td>
<td>4.60324</td>
<td>4.93235</td>
<td>4.60324</td>
<td>4.52329</td>
<td>1.9–6.0</td>
</tr>
<tr>
<td>Density (g/cm³)</td>
<td>0.872103</td>
<td>0.887722</td>
<td>0.870393</td>
<td>0.888883</td>
<td>0.870393</td>
<td>0.872103</td>
<td>0.86–0.90</td>
</tr>
</tbody>
</table>

Where IV- iodine value; SV- saponification value; CN- cetane number; DU- degree of unsaturation; LCSF- long chain saturation factor; HHV- high heating value; CFPP- cold filter plugging point; KV- kinematic viscosity.

**4. Conclusion**

In summary, R. kratochvilovae HIMPA1 has the ability to accumulate high levels of storage lipid enriched with monounsaturated fatty acids under certain nutrient stress conditions. Among various N and P limitation conditions, the highest lipid content (60.34± 0.69%) was obtained when both nutrients were in limited condition (0.1 g/l N and 0.05 g/l P). However, P limitation under...
nitrogen-rich condition gives high lipid content without compromising its biomass that was contrary in N limitation where the lipid accumulation compensates with biomass. P limitation also causes decrease in phospholipid (PE and PC) tremendously while increase in 78.55% TAG as depicted by TLC. An important finding of this study is the yeast cell morphology and lipid droplets size in response to change of cultivation conditions, especially in nutrient stress condition. Under synchronized limitation of N and P super-sized, coalesced irregular shaped lipid lobules were formed in the cellular compartment of oleaginous yeast along with 87.14% enhanced TAG accumulation. Both N and P limitation diverts the CDP-DG pathway of phospholipids synthesis towards de novo TAG synthesis. Moreover, the accumulated lipids under these conditions were rich in MUFA that provide biodiesel having greater performance at a lower temperature (CFPP) and enhanced oxidative stability.

Acknowledgement

The present study was financially supported by DBT, Govt. of India, Bio Care Programme, DBTSanction No. 102/IFD/SAN/3539/2011-2012 (Grant No.: DBT-608-BIO) and SRF to Alok Patel from UGC, India (Grant No.: 6405-35-044).

Appendix A. Supplementary data

Supplementary data related to this article can be found at http://dx.doi.org/10.1016/j.energy.2017.08.052.

References


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