

Fatty Acid Content and Composition of Triacylglycerols of *Chlorella kessleri*

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[Abstract] Triacylglycerols (TAGs) are esters formed from one glycerol and three fatty acids. TAGs are induced to accumulate in algal cells under environmental stress conditions including nutrient-limitation, hyperosmosis, and low temperature, for the storage of metabolic energy and carbon, and also for the consumption of excess energy (e.g., Hirai *et al.*, 2016; Hayashi *et al.*, 2017). Beside their physiological significance, the commercial utilization of algal TAG has been expected for the production of biodiesel, the methyl esters of fatty acids, from the aspect of carbon-neutral conception. The amounts of TAGs can be determined through quantitative measurement of their constituent fatty acids. This protocol consists of the following three parts: the first is the extraction of total lipids from algal cells with the use of organic solvents, chloroform and methanol, according to the method of Bligh and Dyer (1959), the second is the separation of TAG from the other lipid classes by thin-layer chromatography (TLC), and the third is the production of methyl-esterified derivatives of their constitutive fatty acids and subsequent quantitation of them by capillary gas-liquid chromatography (GLC). This protocol adapted from Sato and Tsuzuki (2011) is used for TAG analysis in a green alga, *Chlorella kessleri*.

Keywords: *Chlorella kessleri*, Gas-liquid chromatography, Green algae, Lipids, Thin-layer chromatography, Triacylglycerols

[Background] Several methods have been used for determination of the fatty acid content of TAG. Simple and convenient protocols, e.g., include conversion of TAG to glycerol on treatment with a lipase, and subsequent measurement of the glycerol content through enzymatic generation of a product that reacts with a color- or fluorescence-generating probe (McGowan *et al.*, 1983; Mendez *et al.*, 1986). However, this enzymatic reaction based quantitation of TAG, inevitably, gives no information about the composition of constituent fatty acids. Meanwhile, HPLC provides information on TAG molecular species through their separation based on the numbers of carbon atoms and double bonds of constituent fatty acids, and enables their respective quantitation when combined with tandem mass spectrometer like in LC-MS/MS (Mu *et al.*, 2000; Dorschel, 2002; MacDougall *et al.*, 2011). The LC-MS/MS instrument, however, is very expensive. In this context, TLC/GLC based protocol for the measurement of the fatty acid content of TAG is introduced here, in view of the requirement of less expensive equipment than LC-MS/MS and definite information that can be obtained on quality and quantity of the constituent fatty acids.

Materials and Reagents

1. 50 ml polypropylene centrifuge tubes with conical bottom (Corning, Falcon®, catalog number: 352070)
2. 50 ml glass screw cap centrifuge tubes with PTFE lined phenolic caps (AGC Techno Glass, catalog number: 8422CTF50)
3. 9 inch glass Pasteur pipettes (AGC Techno Glass, catalog number: IK-PAS-9P)
4. TLC silica gel 60 glass plates 20 x 20 cm (Merck, catalog number: 105721)
5. Chromatography filter paper 1CHR 200 x 200 mm (GE Healthcare, catalog number: 3001-861)
6. Glass microcapillary pipette (Sigma-Aldrich, catalog number: Z543292)
7. 100 µl microsyringe (1710 RN, Hamilton, catalog number: 81030) with 22s/51/2 needle (Hamilton, catalog number: 7758-03)
8. 14 ml glass screw cap test tubes with PTFE lined phenolic caps (AGC Techno Glass, catalog number: TST-SCR16-125)
9. Inserts for large opening vials volume 0.15 ml (Sigma-Aldrich, catalog number: 24719)
10. 2 ml large opening vials with open-top screw cap (Sigma-Aldrich, catalog number: 29116-U)
11. ULBON HR-Thermon-3000B GLC capillary column I.D. 0.25 x 25 m (Shinwa Chemical Industries)
12. *Chlorella kessleri* 11 h, which corresponds to *Parachlorella kessleri* of NIES collection (<http://mcc.nies.go.jp/>) (National Institute for Environmental Studies, catalog number: NIES-2160)
13. Potassium chloride (KCl) (Wako Pure Chemical Industries, catalog number: 163-03545)
14. Butylated hydroxytoluene (BHT) (Wako Pure Chemical Industries, catalog number: 029-07392)
15. Methanol (CH₃OH) (Wako Pure Chemical Industries, catalog number: 138-06473)
16. Chloroform (CHCl₃) (Wako Pure Chemical Industries, catalog number: 033-08631)
17. Primuline (C₂₁H₁₅N₃O₃S₃) (Tokyo Chemical Industry, catalog number: P0603)
18. Acetone (CH₃COCH₃) (Wako Pure Chemical Industries, catalog number: 016-00346)
19. Tripalmitin (C₅₁H₉₈O₆) as TAG standard (Wako Pure Chemical Industries, catalog number: 200-03002)
20. N₂ gas
21. Hydrogen chloride-methanol reagent (5-10%) (Tokyo Chemical Industry, catalog number: X0041)
22. *n*-Hexane (C₆H₁₄) (Wako Pure Chemical Industries, catalog number: 084-03421)
23. Supelco 37 component FAME mix (Sigma-Aldrich, catalog number: 47885-U)
24. Gamborg's B5 medium salt mixture (Nihon Pharmaceutical, catalog number: 399-00621; Gamborg *et al.*, 1968)
25. Sorbitol (C₆H₁₄O₆) (Wako Pure Chemical Industries, catalog number: 198-03755)
26. Arachidic acid (C₂₀H₄₀O₂) (Tokyo Chemical Industry, catalog number: E0006)
27. 1/4 GB medium with or without 0.6 M sorbitol (see Recipes)

28. Arachidic acid solution as an IS (see Recipes)

Equipment

1. High-pressure steam sterilizer (TOMY SEIKO, model: LBS-245)
2. Tabletop centrifuge (Kubota, model: 5220) equipped with ST-720M swing rotor (16 x 50 ml)
3. Vortex mixer (Scientific Industries, model: Vortex-Genie 2)
4. Rotary evaporator (Tokyo Rikakikai, model: N-1110V-W) equipped with screw cap tube adaptor
5. UV transilluminator (UVP, model: LM-20)
6. Fume hood (Yamato Scientific, model: KFS)
7. Forced air flow oven (Tokyo Rikakikai, model: WFO-451SD)
8. Spectrophotometer (Beckman Coulter, model: DU 640)
9. Double trough TLC chamber for 20 x 20 cm plates (Camag, catalog number: 022.5256)
10. Capillary gas-liquid chromatograph (Shimadzu, model: GC-2025) equipped with split/splitless injector, flame ion detector (FID) and autoinjector
11. Chromatopac integrator (Shimadzu, model: C-R7A plus)
12. Glass chromatographic reagent atomizer (Corning, PYREX[®], catalog number: 2153-125)

Procedure

A. Total lipid extraction

1. Culture *C. kessleri* cells at 30 °C in 100 ml of 1/4 GB (see Recipes) without sorbitol, under illumination (30 W m⁻²) and aeration for 48-72 h, to the late exponential phase of growth (the values of optical density at 730 nm [OD₇₃₀] within ca. 0.4 to 0.6).
2. Harvest cells through centrifugation in 2 x 50 ml tubes at 1,500 x g for 15 min at room temperature (tabletop centrifuge). Discard supernatant by decantation and resuspend cells in 20 ml of 1/4 GB with 0.6 M sorbitol for the induction of TAG accumulation (Hirai *et al.*, 2016).
3. Harvest cells through centrifugation at 1,500 x g for 15 min. Discard supernatant through decantation and resuspend cells in 20 ml of 1/4 GB with 0.6 M sorbitol. Repeat this centrifugation-resuspension step twice, finally with the OD₇₃₀ value of cell culture adjusted to 0.3 in 50 ml of 1/4 GB with 0.6 M sorbitol. Culture cells for three days under the same growth conditions.
4. Harvest cells through centrifugation at 1,500 x g for 15 min at room temperature. Discard supernatant through decantation and resuspend cells in 2 ml of 0.1 M KCl.
5. Transfer cell suspension into a glass screw cap centrifuge tubes (50 ml vol.) with PTFE lined phenolic caps. Add 6.0 ml of methanol and 30 µl of 1.0% (w/v) BHT in methanol as an antioxidant agent to the sample, and agitate it by a vortex mixer for 30 sec at the maximum speed to destabilize cellular membrane systems.
6. Add 3.0 ml of chloroform to the sample. Agitate it by a vortex mixer for 30 sec at the maximum

speed and stand it for 10 min to extract lipids from the cells.

7. Add 3.0 ml of chloroform once again and agitate the sample for intensive extraction of lipids by a vortex mixer for 30 sec at the maximum speed.
8. Add 3.0 ml of distilled water and vortex the sample for 30 sec for emulsification. Centrifuge the mixture at 1,000 x g for 5 min at room temperature for its separation into three phases, *i.e.*, the upper phase of H₂O and methanol, the medium phase of cell debris, and the lower phase of chloroform. There is no cell debris at the bottom of the tube.
9. Transfer the lower phase including total lipids with a Pasteur pipette into a glass screw cap centrifuge tubes (50 ml vol.) with PTFE lined phenolic caps. Be careful not to pick up the cell debris or the upper phase. The remaining lower phase can be recovered by the subsequent procedure.
10. Add 3.0 ml of chloroform to the remaining upper and medium phases, and agitate the mixture for 30 sec with a vortex mixer. Centrifuge it at 1,000 x g for 15 min at room temperature.
11. Recover the lower phase to the glass tube at Step A9.
12. Repeat Steps A10 to A11 twice until lipids are fully recovered. Full recovery of lipids can be confirmed through observation of bleaching of the cell debris, which results from extraction of green chlorophylls into the lower phase.

Note: Do not mechanically disrupt C. kessleri cells before lipid extraction, since the mechanical disruption would not increase the TAG recovery, but would decrease the TAG content with appearance of free fatty acids, owing probably to the action of endogenous lipases.

13. Evaporate the solvent completely with a rotary evaporator, if necessary, with the addition of methanol or chloroform/methanol (2:1, v/v) for removal of residual H₂O through azeotropy. Weigh dry residue and dissolve lipids in *ca.* 200 µl of chloroform/methanol (2:1, v/v), and transfer this total lipid fraction into a glass vial to be stored at -20 °C until use. The value of dry residue weight will be helpful for optimization of loading of total lipids on TLC plate for better separation of TAG.

B. Separation of TAG from the other lipid classes by TLC

1. Set a TLC chamber for the separation of TAG by placing a piece of filter paper along its wall, and then by pouring a developing solvent of hexane/diethylether/acetate (70:30:1, v/v) up to *ca.* 1 cm in depth. Close the lid for vapor saturation, which is promoted by the filter paper that has been soaked in the solvent. Stand the chamber for an hour or more to saturate it with the solvent vapor prior to the start of development. In order to ensure repeatability, the solvent is prepared freshly for each analysis.
2. Heat a TLC silica gel 60 plate at 120 °C for a couple of hours in an oven to activate the silica gel just before development. Don't touch the plate with bare hands during this procedure.
3. Draw a horizontal line that is about 2 cm away from the bottom side of a TLC plate with a carbon pencil, and outline a horizontally long rectangle (*ca.* 2 x 3 cm) on the horizontal line of the plate. Lines have to be gently drawn to avoid scraping off silica gel. Don't use an ink pen,

otherwise, the lines would disappear.

Note: Depending on the number of samples, the TLC plate may be cut into two pieces (10 cm wide ones) before use.

4. Apply 45% of the stored total lipid fraction to the rectangle spot with a glass microcapillary pipette or 100 μ l microsyringe with 22s/51/2 needle, and air-dry it. Repeat this step several times until this fraction is completely transferred to this spot.
5. Apply 5-25 μ l of standard TAG stock solution (1 mg/ml in chloroform, w/v) on the horizontal line next to the spot of the analyte samples.
6. Place the TLC plate in the chamber, cover the chamber with its lid, and allow the plate to develop until the solvent goes up to about 2 cm below the top side of the plate.
7. Remove the plate from the chamber and air-dry the plate.
8. Spray the plate with primuline (0.01% in 80% acetone, w/v) in a fume hood, and illuminate it under 365 nm UV light to detect lipid compounds as exhibiting a pale color (see Figure 1). Outline the TAG spot with a pencil, judged from the position of TAG standard.

Note: To prevent the TAG spot from diffusing, avoid spraying excessive detection reagent.

9. Scrap off the silica gel of the TAG spot by a flat edge of micro-spatula. Silica gel from TAG should be immediately treated for the production of fatty acid methyl esters (FAMES, see below) to prevent the oxidation of its constituent fatty acids.

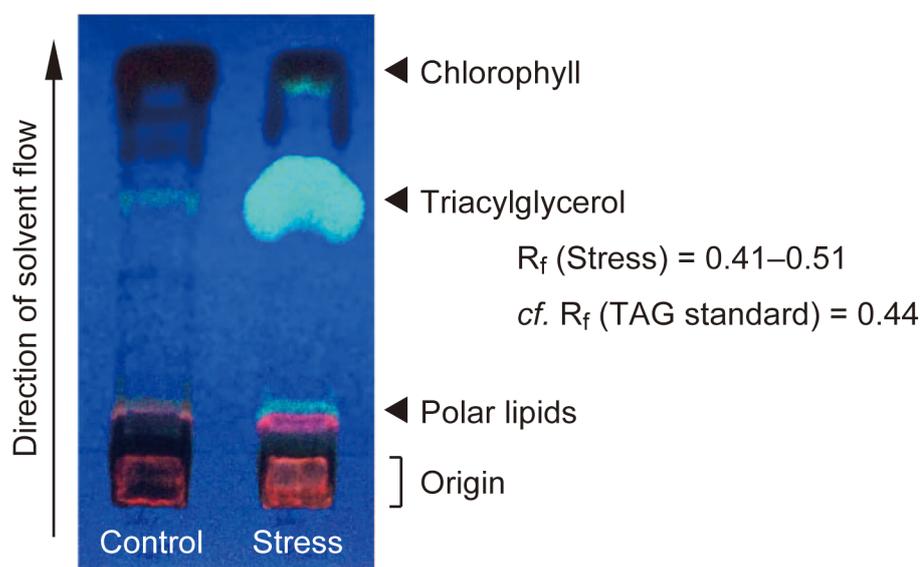


Figure 1. TLC separation of TAG in total lipid fraction in *C. kessleri*

C. Conversion of constituent fatty acids into fatty acid methyl esters

1. Take 50 μ l of arachidic acid solution as an internal standard (IS, see Recipes) into glass screw cap test tubes (14 ml vol.) with PTFE lined phenolic cap, and dry it up under the flow of N_2 gas.
*Note: Since *Chlorella kessleri* 11 h has no endogenous C20 series fatty acids, the authors use arachidic acid (C20:0) as an IS. Nonadecanoic acid (C19:0) can also be used as an IS.*

- Put scraped silica gel of TAG and 5% of the total lipid fraction into the test tubes, respectively. The latter lipid solution has to be air-dried under the flow of N₂ gas. The remainder of the total lipid fraction, *i.e.*, 50% of the initial level, can be stored at -80 °C for later analysis.
- Add 2.0 ml of 5-10% (w/v) hydrogen chloride-methanol reagent to each tube, cap it tightly, and agitate it by a vortex mixer for 30 sec at the maximum speed.
- Heat each tube at 95 °C for 2 h for methyl-esterification of the constituent fatty acids of TAG or total lipids.
- Cool down the tubes to room temperature, and add 2 ml of *n*-hexane to it. Agitate the tubes by a vortex mixer for 30 sec at a maximum speed for the extraction of FAMES.
- Stand the tubes for 1 min for the separation of its content into upper (*n*-hexane) and lower (methanol) phases.
- Transfer the upper phase into a glass screw cap centrifuge tube (50 ml vol.) with PTFE lined phenolic caps by a glass Pasteur pipette.
- Add 2.0 ml of *n*-hexane to the remaining lower phase and vortex it for 30 sec. Stand the tube for 1 min at a room temperature.
- Recover the upper phase to the glass tube at Step C7.
- Repeat Steps C8 to C9 additionally two times for intensive extraction of FAMES.
- Evaporate FAMES solution with a rotary evaporator, and dissolve the dried samples in a small volume (< 50 µl) of *n*-hexane. Transfer the concentrated FAMES solution into a 0.15 ml insert in a 2 ml glass vial, which is then capped for storage at 4 °C until measurement.
- Analyze fatty acid composition of the sample with a capillary GLC (see Figure 2). GLC is operated according to manufacturer's instructions. GLC operating conditions are shown in Table 1.

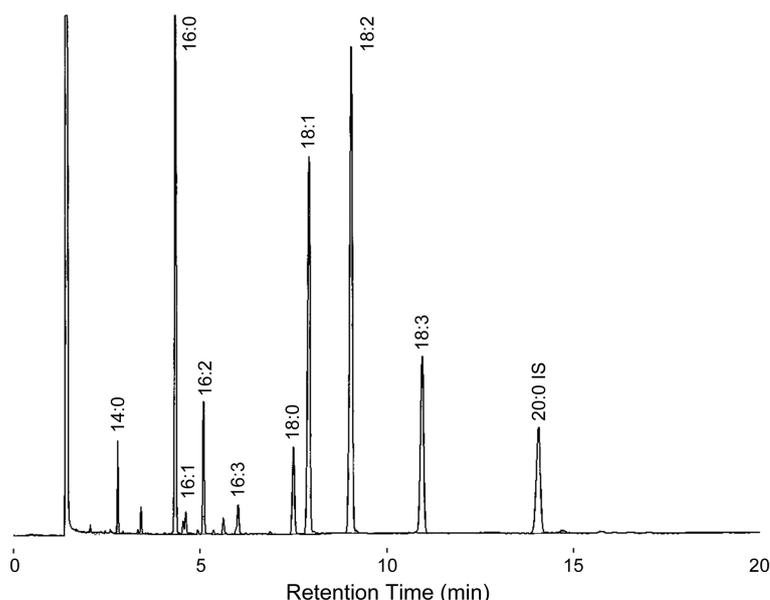


Figure 2. A gas chromatogram of FAMES from TAG of *C. kessleri*

Table 1. GLC operating conditions for determination of fatty acid composition

	Specification
Column	ULBON HR-Thermon-3000B 0.25 x 25 m
Injection volume	1 μ l
Carrier gas flow	1 ml/min Helium, 115 kPa
Injector temperature	250 °C, split
Split flow	50 ml/min (split ratio: 1:50, v/v)
Oven temperature	180 °C isothermal
Detector	250 °C, Flame Ionization Detector (FID)

Note: FID detects ions that are generated through burning of carbon compounds (Holm, 1999).

Data analysis

1. Identify and calculate the peak areas of respective FAMES on chromatogram with an integrator according to manufacturer's instructions. FAMES identification is based on the retention times as compared with those of the standard FAME mixture (Supelco 37 component FAME Mix, 100 μ g/ml solution diluted with *n*-hexane). Check that chromatogram is being integrated properly with the appropriate baseline being shown.
2. Quantitate each molecular species of FAME by using following Formula [1] with arachidic acid (20:0) as an IS. The values, 160 and 312, indicate the content of 20:0 (nmol) taken at Step C1 and the molecular weight of 20:0, respectively. The molar content of total lipids can be expressed on the basis of fatty acids that are included (Formula [2]).
3. The TAG content is estimated as 1/3 of the summed molar contents of its constituent fatty acids (Formula [3]). Be careful that obtained values correspond to those included in respective samples that are analyzed. Alternatively, the content of TAG relative to that of total lipids is estimated on the basis of fatty acids (Formula [4]). As shown here, multiplication of 'summed content of FAME from TAG' by 1/9 is necessary in accordance with the quantitative ratio of the total lipid fraction used for TAG analysis (45%) to that for total lipid analysis (5%). The quantitative results should be presented by mean values \pm standard deviations from at least three experiments.

$$\text{FAME content (nmol)} = 160 \frac{\text{FAME peak area / FA molecular weight}}{20:0\text{-ME peak area} / 312} \quad [1]$$

Note: The peak area of FAME is conveniently divided by the molecular weight of FA, but not by that of FAME, in view of full responses of FAME carbon atoms except its carbonyl one in FID (Ackman and Sipos, 1964).

$$\text{Total lipids content (nmol fatty acids)} = \text{summed content of FAME from total lipids} \quad [2]$$

$$\text{TAG content (nmol)} = \frac{\text{summed content of FAME from TAG}}{3} \quad [3]$$

The content of TAG relative to total lipids (mol%) = $1/9 \frac{\text{summed content of FAME from TAG}}{\text{summed content of FAME from total lipids}}$ [4]

Recipes

- 1/4 GB medium with or without 0.6 M sorbitol
 - Dissolve Gamborg's B5 medium salt mixture (3.3 g/packet, Table 2) in distilled water for preparation of normal Gamborg's B5 medium (GB; final volume, 1 L)

Table 2. Chemical composition of Gamborg's B5 medium salt mixture

Chemical	(mg/L)	Chemical	(mg/L)
KNO ₃	2,500	MnSO ₄ ·H ₂ O	10
MgSO ₄ ·7H ₂ O	250	H ₃ BO ₃	3.0
NaH ₂ PO ₄ ·H ₂ O	150	ZnSO ₄ ·7H ₂ O	2.0
CaCl ₂ ·2H ₂ O	150	KI	0.75
(NH ₄) ₂ SO ₄	134	Na ₂ MoO ₄ ·2H ₂ O	0.25
Na ₂ -EDTA	37.3	CuSO ₄ ·5H ₂ O	0.025
FeSO ₄ ·7H ₂ O	27.8	CoCl ₂ ·6H ₂ O	0.025

- Add 750 ml of distilled water to 250 ml of GB for preparation of 1/4 GB
 - Otherwise, dissolve 109 g of sorbitol in 250 ml of GB with concomitant addition of distilled water for preparation of 0.6 M sorbitol containing 1/4 GB (final volume, 1 L)
 - 1/4 GB and 0.6-M sorbitol containing 1/4 GB have to be subjected to autoclave sterilization (120 °C, 20 min)
- Arachidic acid solution as an IS
 - Dissolve 10 mg of arachidonic acid in 10 ml 2-propanol
 - Store the solution at 2-8 °C

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