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(54) NOVEL MICROALGAE AND USE FOR SAME

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(30) Foreign Application Priority Data

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CPC *C12N 1/12* (2013.01); *A23K 10/16*
(2016.05); *A23L 17/60* (2016.08); *A23L
33/105* (2016.08); *A61K 8/9717* (2017.08);
C12N 1/06 (2013.01); *C12N 15/74* (2013.01);
A23V 2002/00 (2013.01)

(57)

ABSTRACT

Provided is an alga belonging to Cyanidiophyceae having a
diploid cell and a haploid cell form. Also provided is a
nutrient composition containing an alga belonging to Cyano-
idiophyceae or an extract thereof.

Specification includes a Sequence Listing.

PCR product

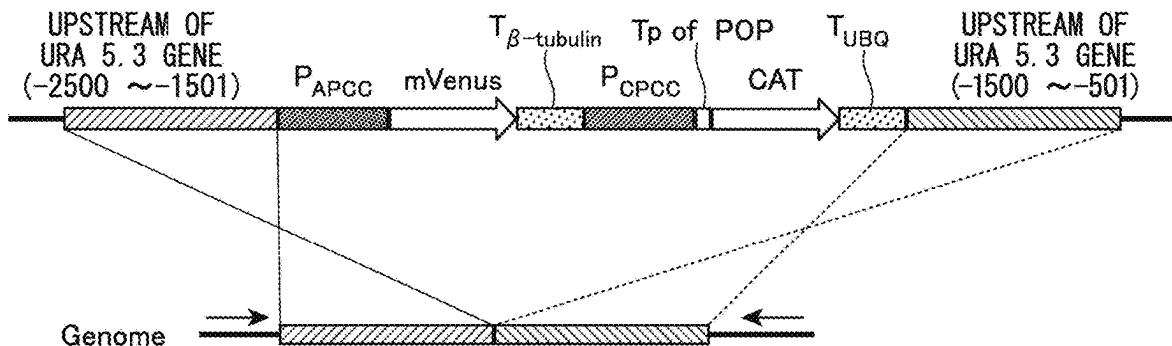


FIG. 1A

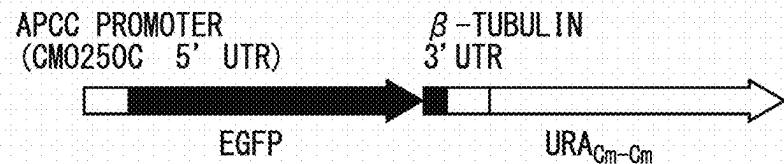


FIG. 1B

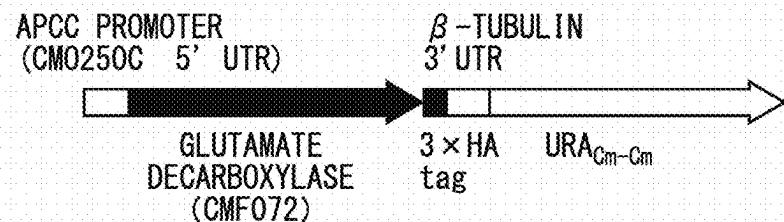


FIG. 1C

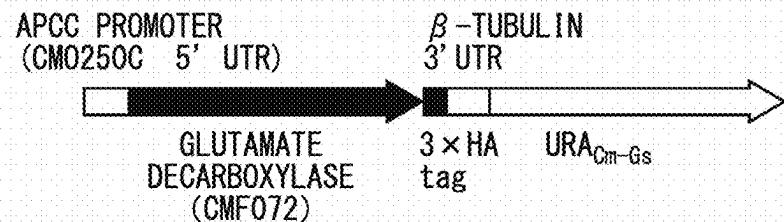


FIG. 2

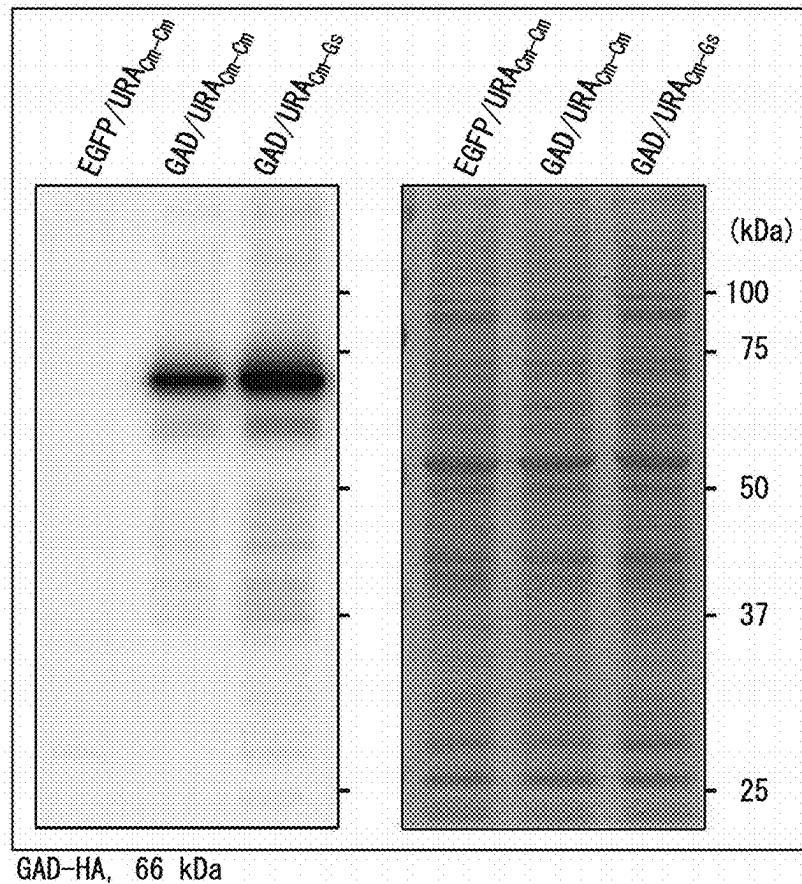


FIG. 3

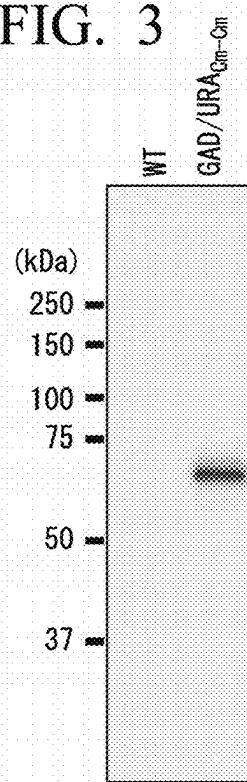
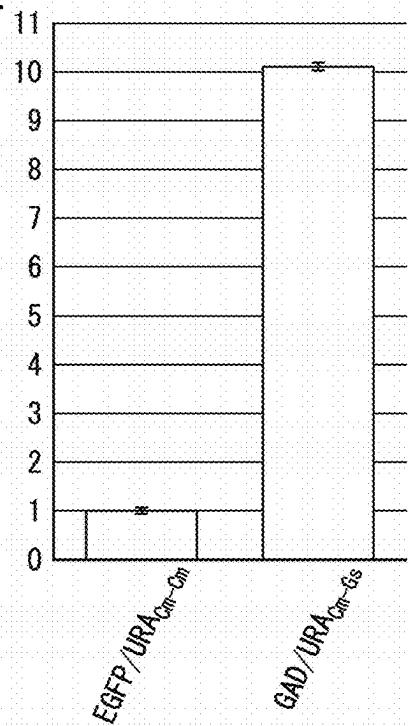


FIG. 4



(error bar = standard deviation: $n=3$)

FIG. 5

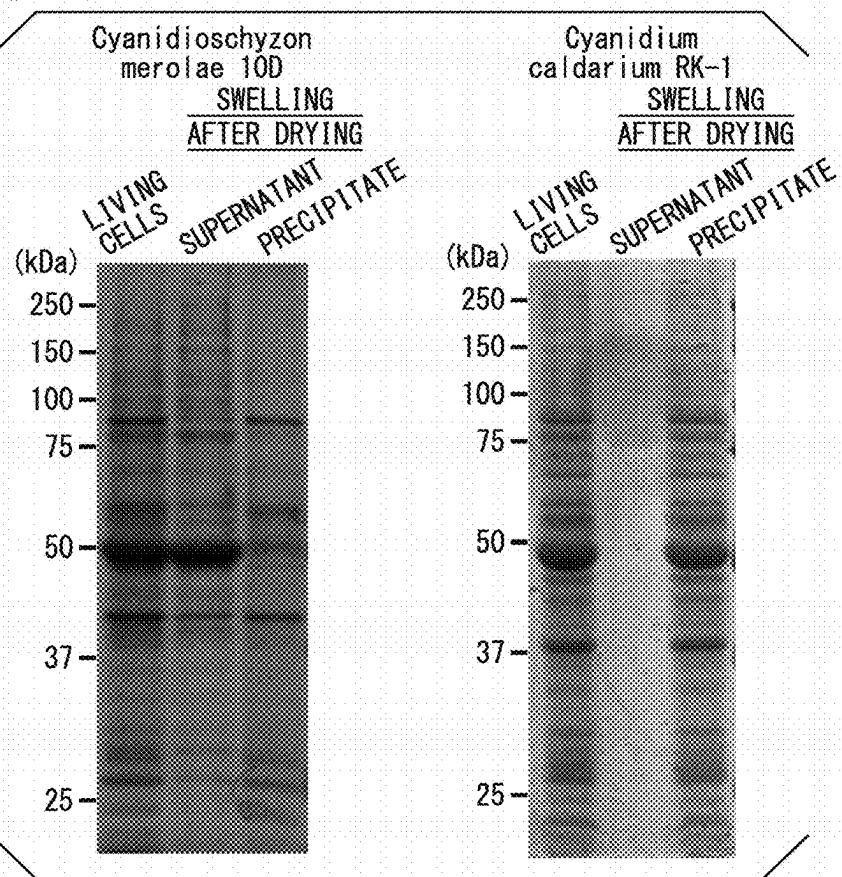


FIG. 6A

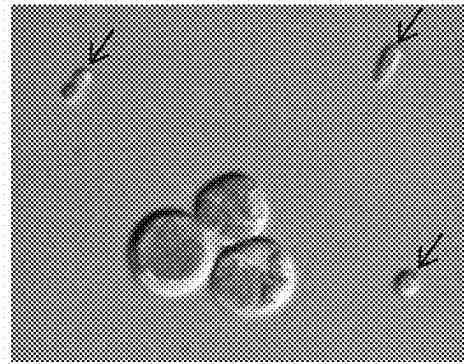


FIG. 6B

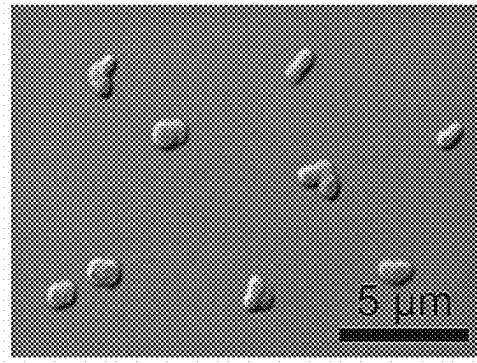


FIG. 7

ATTTTGTCACCTAACCTATGGATGGTAAACCAATTGGATGGTTCACCAAATGGATGGTACCCCTTACAGGAGCCGTCTCTCGTACGGTTGGT SEQ ID NO:12
HKN1_allele 1 ATTTTGTCACCTAACCTATGGATGGTAAACCAATTGGATGGTTCACCAAATGGATGGTACCCCTTACAGGAGCCGTCTCTCGTACGGTTGGT SEQ ID NO:12
ATTTTGTCACCTAACCTATGGATGGTAAACCAATTGGATGGTTCACCAAATGGATGGTACCCCTTACAGGAGCCGTCTCTCGTACGGTTGGT SEQ ID NO:13
HKN1_allele 2 ATTTTGTCACCTAACCTATGGATGGTAAACCAATTGGATGGTTCACCAAATGGATGGTACCCCTTACAGGAGCCGTCTCTCGTACGGTTGGT SEQ ID NO:13
ATTTTGTCACCTAACCTATGGATGGTAAACCAATTGGATGGTTCACCAAATGGATGGTACCCCTTACAGGAGCCGTCTCTCGTACGGTTGGT SEQ ID NO:14
HKN1_GHYZON-LIKE allele 1 ATTTTGTCACCTAACCTATGGATGGTAAACCAATTGGATGGTTCACCAAATGGATGGTACCCCTTACAGGAGCCGTCTCTCGTACGGTTGGT SEQ ID NO:14
HKN1_GHYZON-LIKE allele 2 ATTTTGTCACCTAACCTATGGATGGTAAACCAATTGGATGGTTCACCAAATGGATGGTACCCCTTACAGGAGCCGTCTCTCGTACGGTTGGT SEQ ID NO:14

FIG. 8A-8E

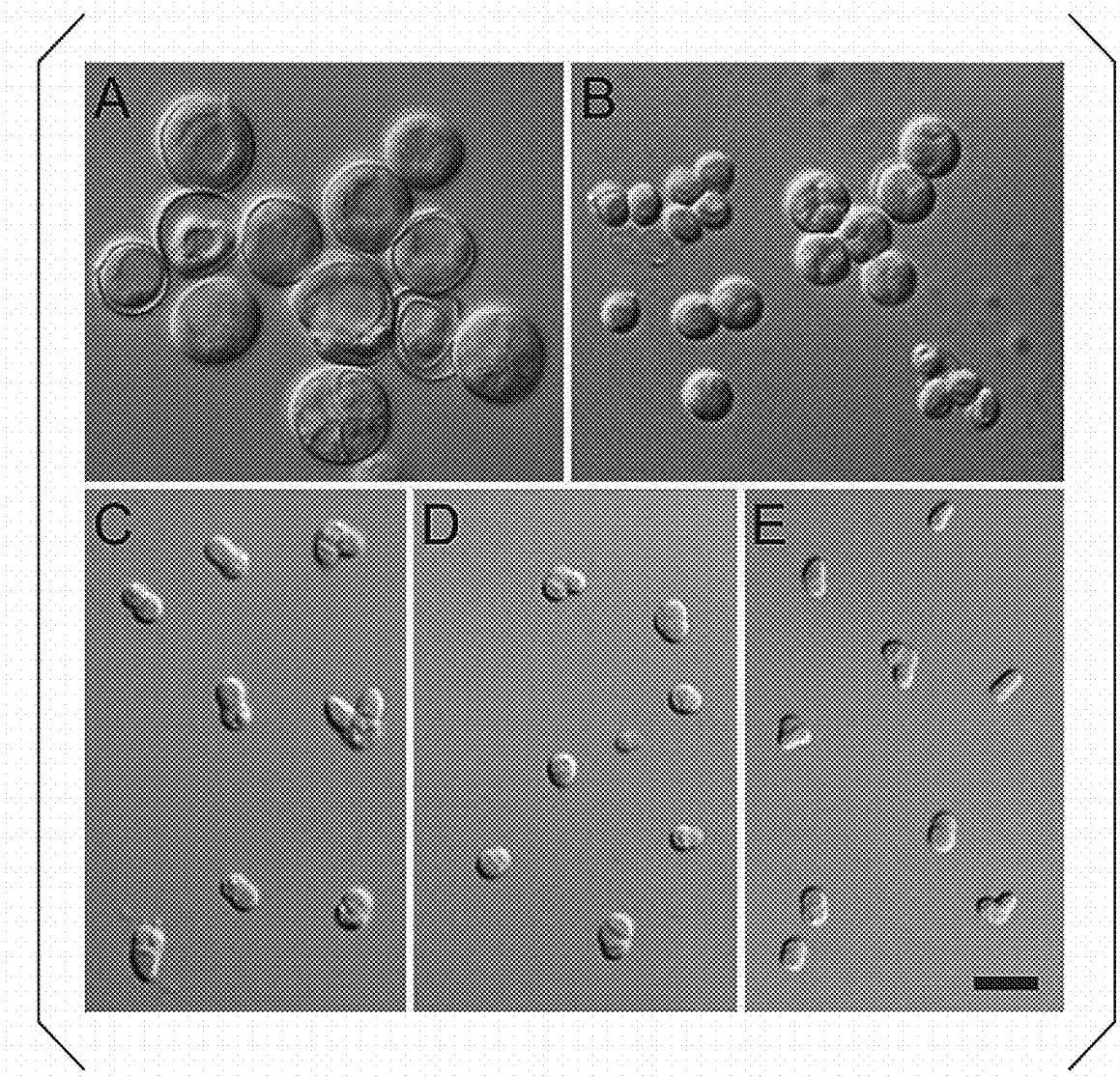


FIG. 9

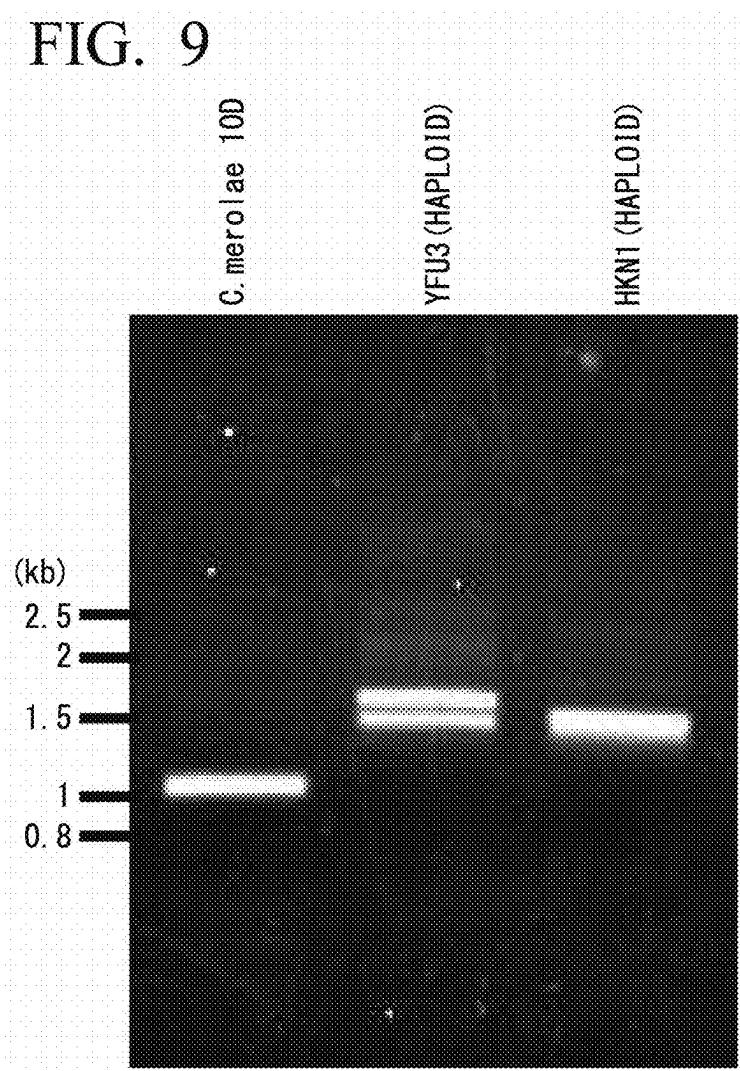


FIG. 10

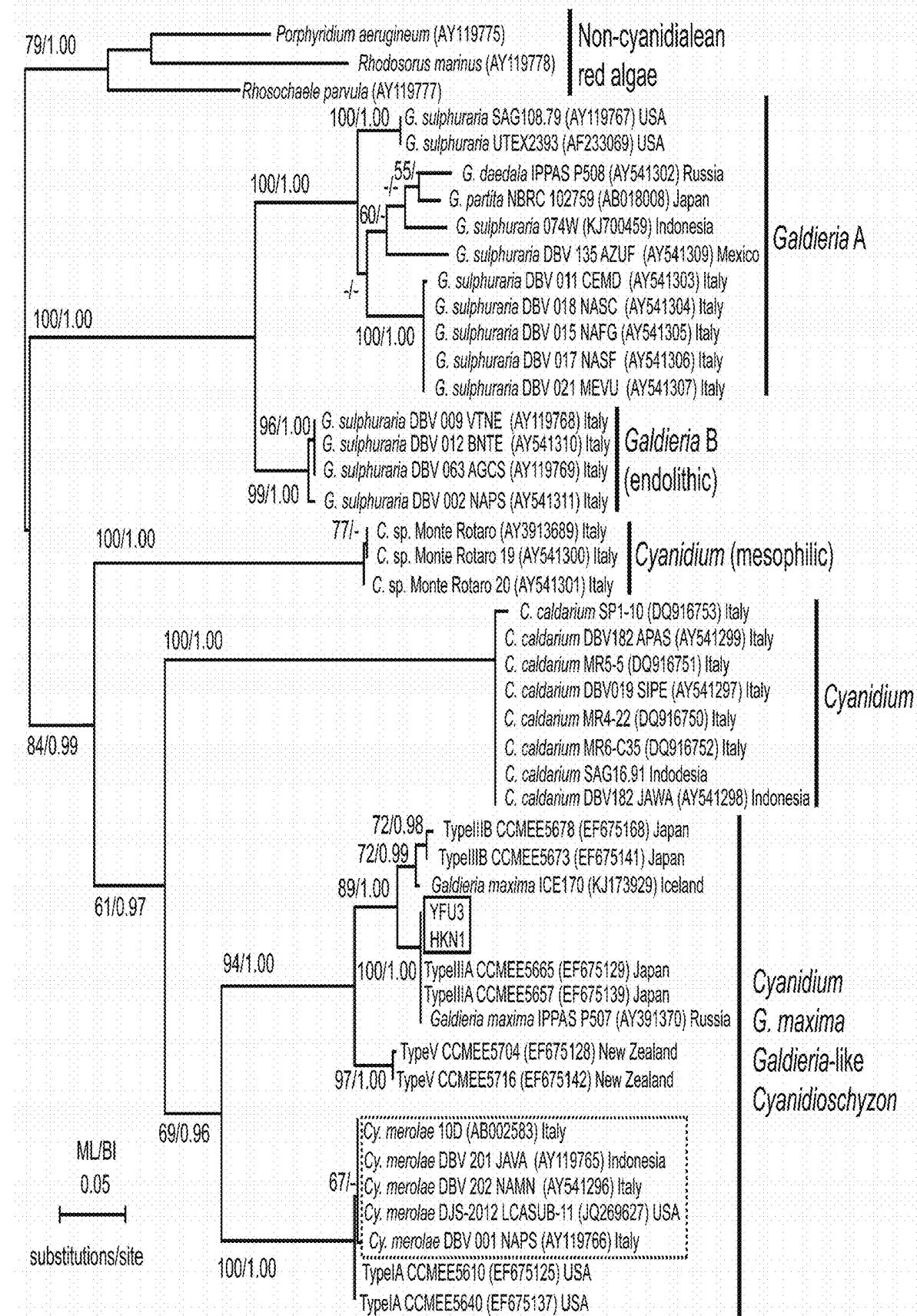


FIG. 11

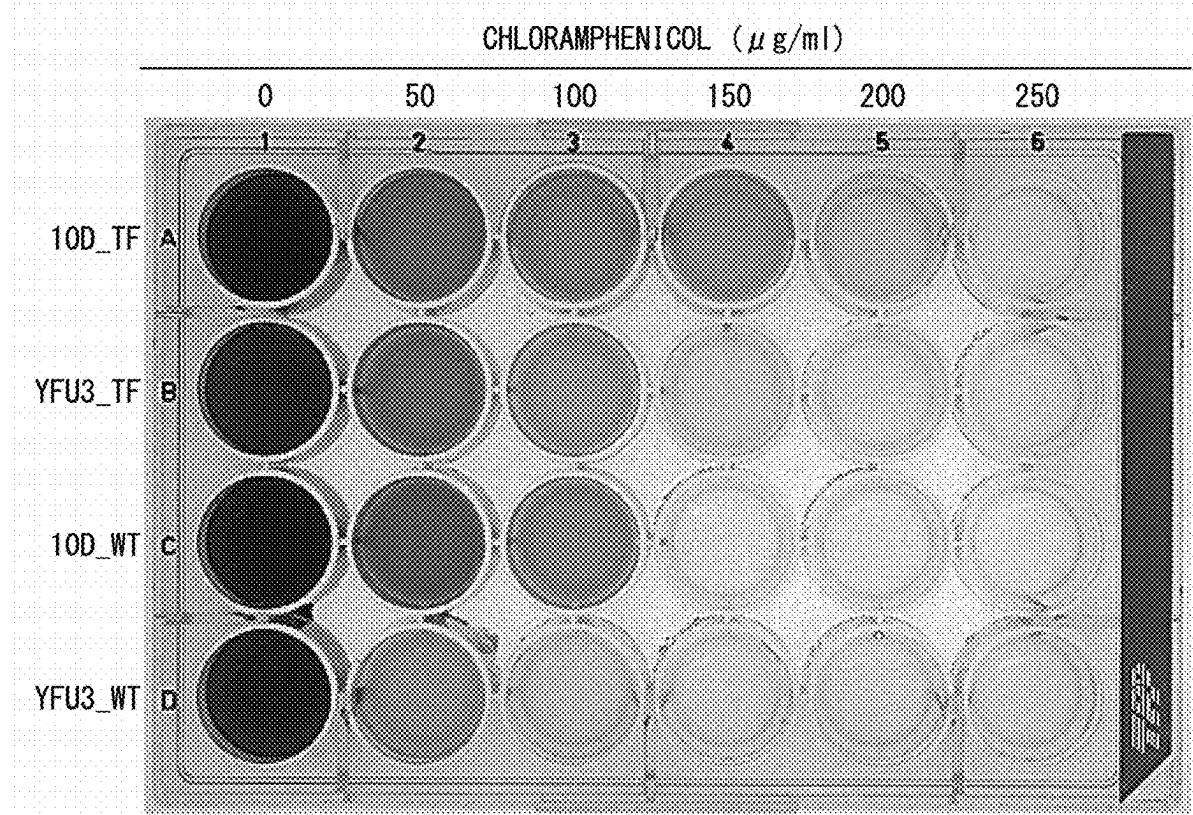


FIG. 12

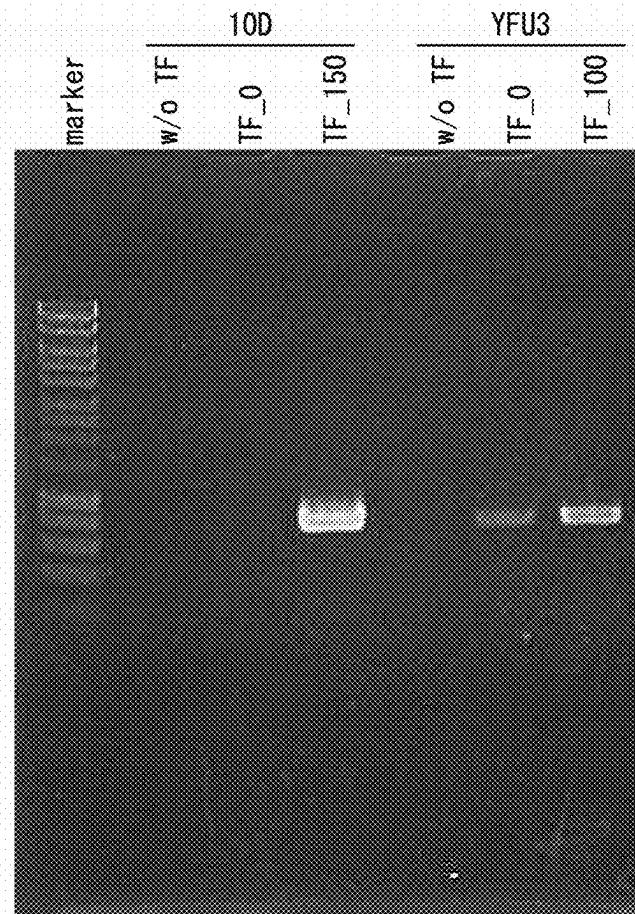


FIG. 13

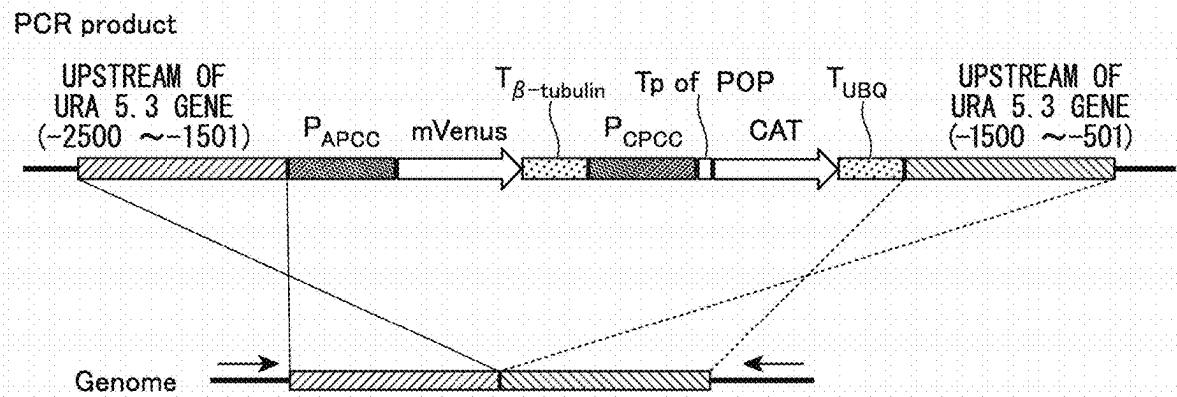


FIG. 14

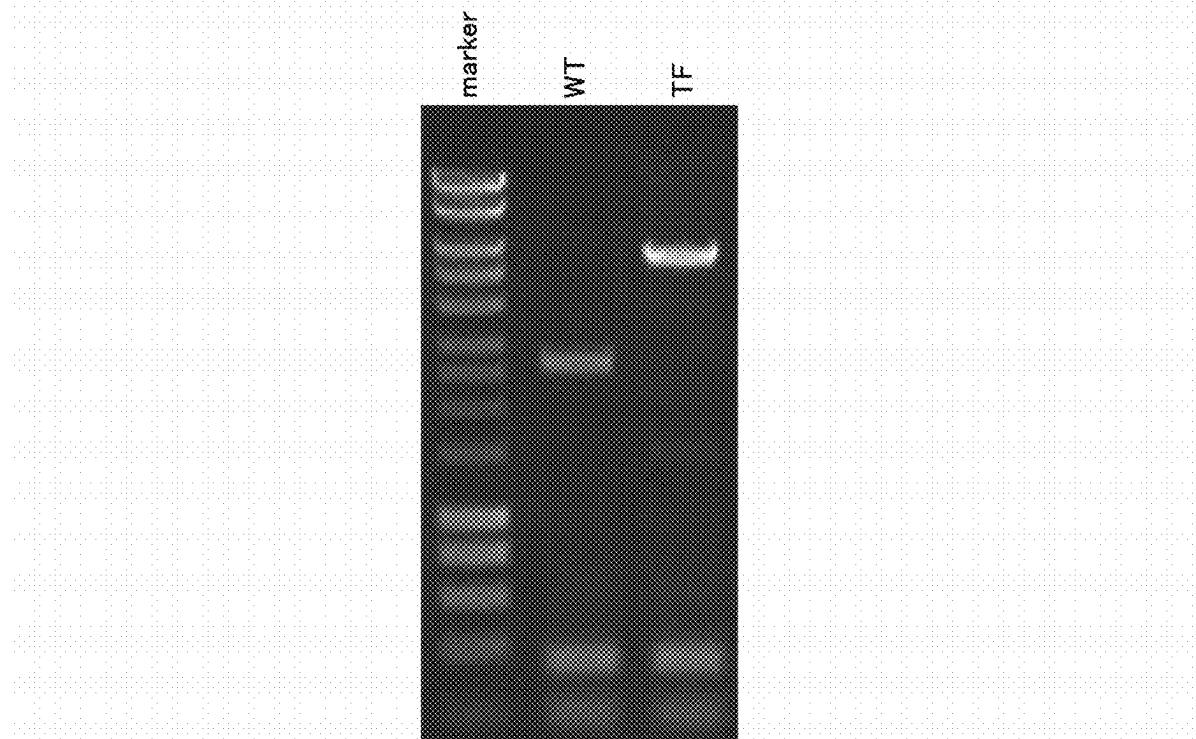


FIG. 15

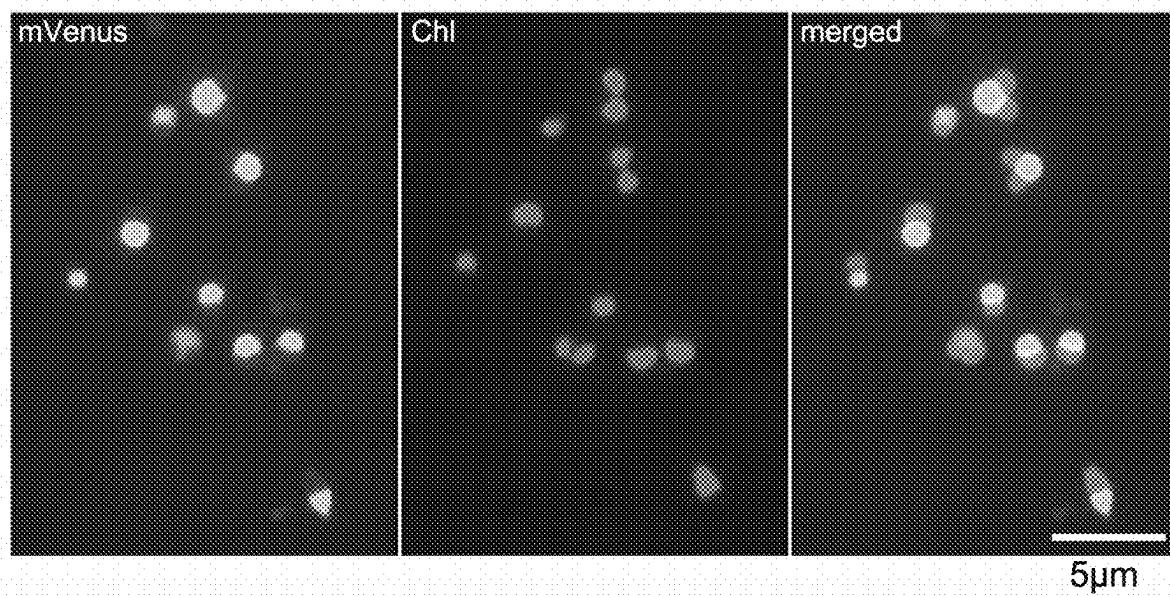


FIG. 16

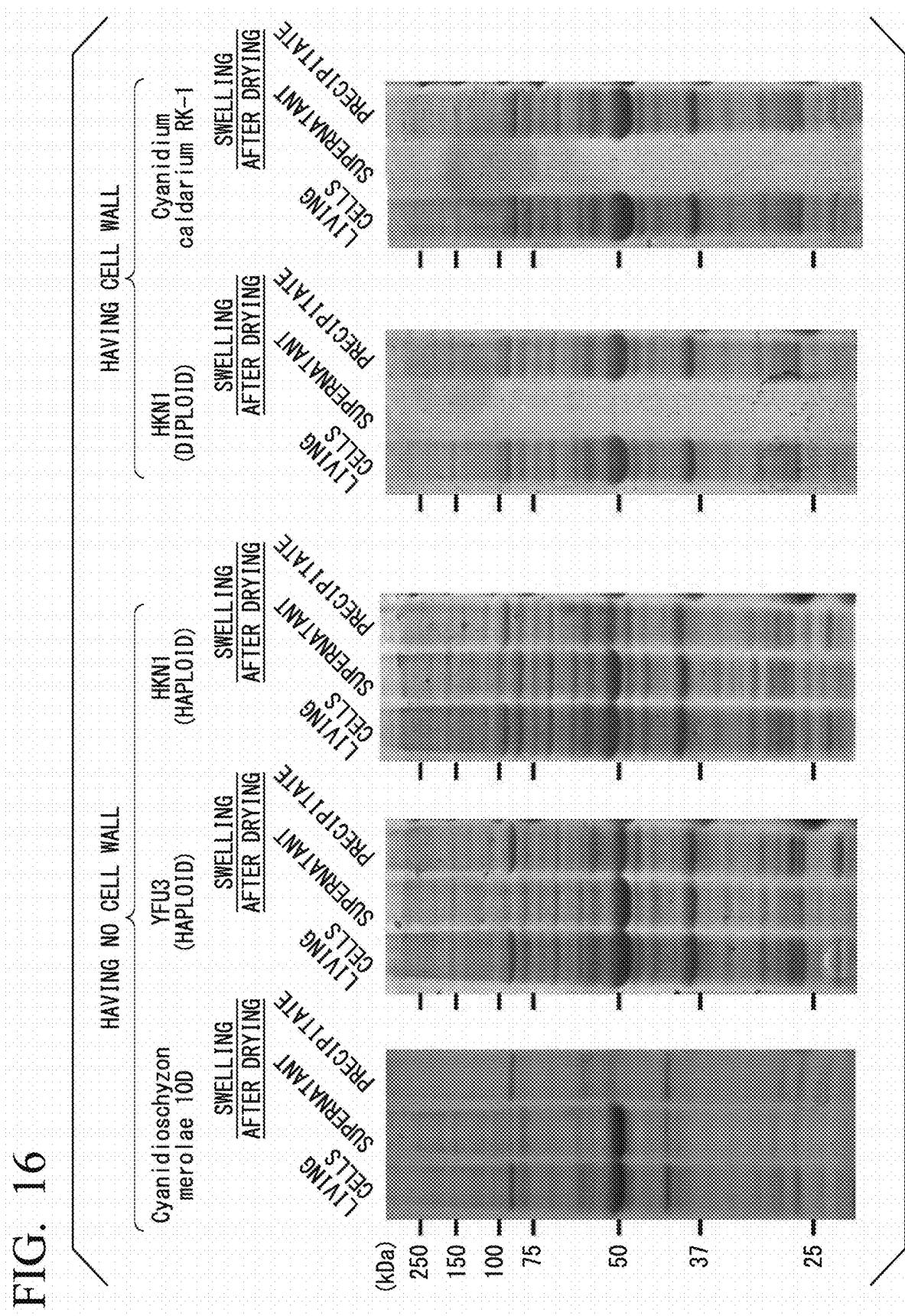


FIG. 17A

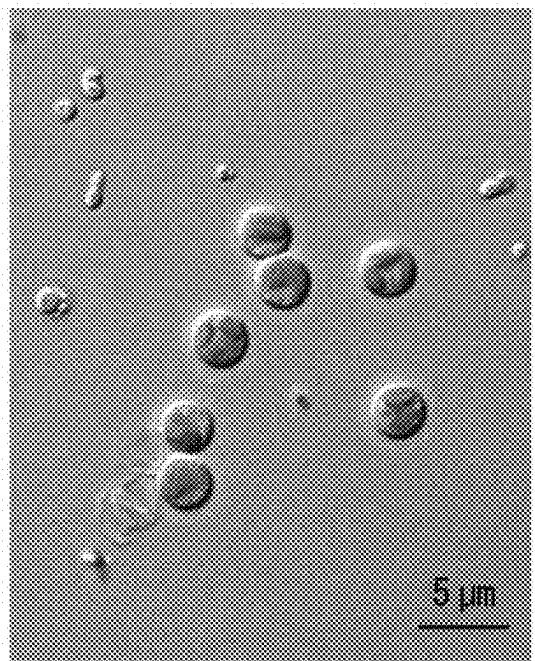


FIG. 17B

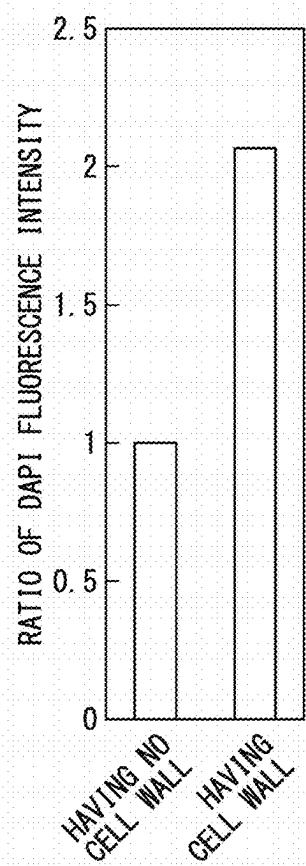


FIG. 18A

Galdieria sulphuraria SAG108.79

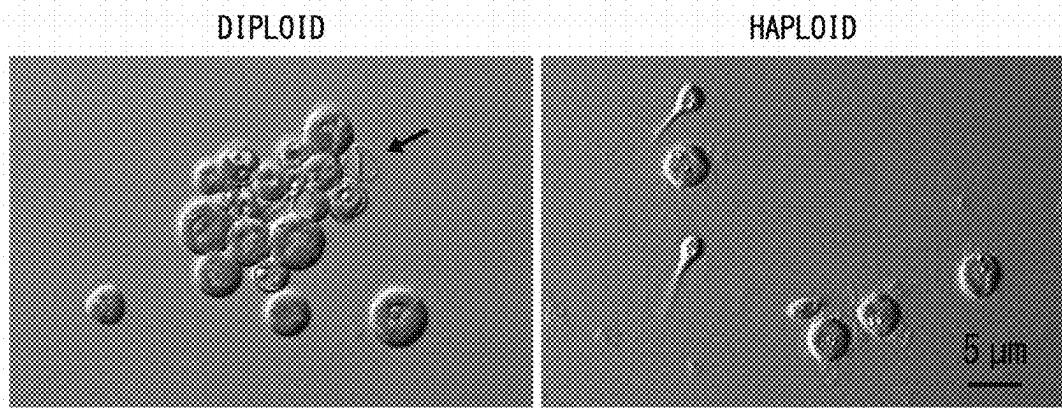


FIG. 18B

Galdieria partita NBRC 102759

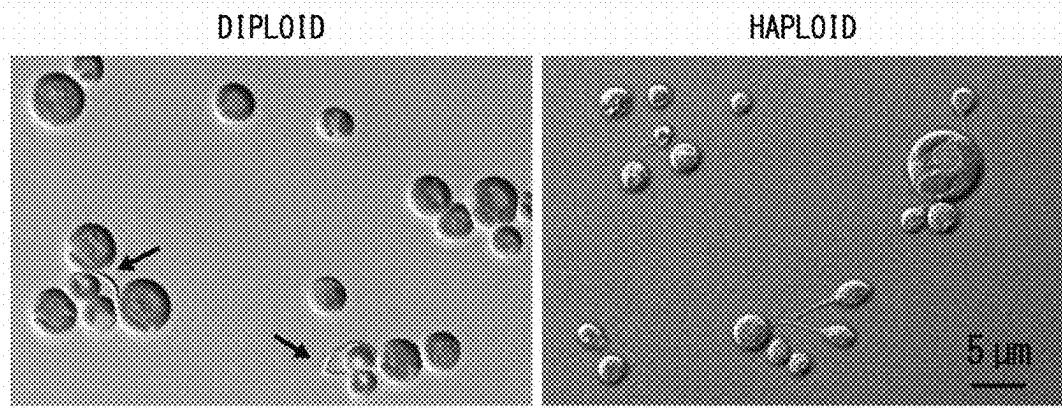


FIG. 19

2N_allele1 GTTGTAAACTCGGATGGCAAGTACTTTGGAAATTGACCATGAAAGATCCCTCACAAAGTGGAAACTTGCTCTCAGTTCTCCCTGT
2N_allele2 GTTGTAAACTCGGATTCGGATGGCAAGTACTTTGGAAATTGACCATGAAAGATCCCTCTCACAAAGTGGAAACTTGCTCTCAGTTCTCCCTGT
N_clone1 GTTGTAAACTCGGATTCGGATGGCAAGTACTTTGGAAATTGACCATGAAAGATCCCTCACAAAGTGGAAACTTGCTCTCAGTTCTCCCTGT
N_clone2 GTTGTAAACTCGGATTCGGATGGCAAGTACTTTGGAAATTGACCATGAAAGATCCCTCACAAAGTGGAAACTTGCTCTCAGTTCTCCCTGT
N_clone3 GTTGTAAACTCGGATTCGGATGGCAAGTACTTTGGAAATTGACCATGAAAGATCCCTCTCACAAAGTGGAAACTTGCTCTCAGTTCTCCCTGT
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100

FIG. 20

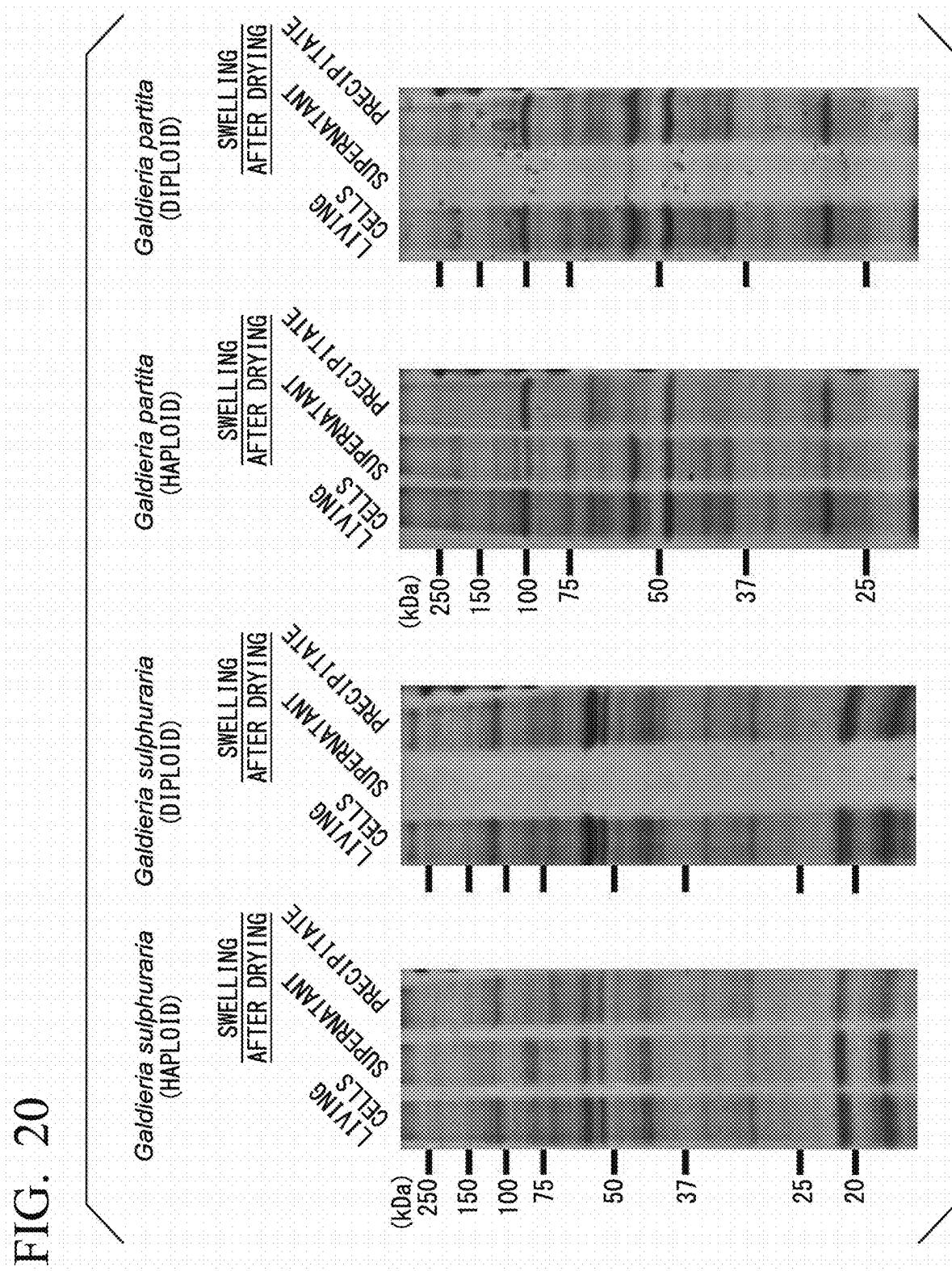


FIG. 21

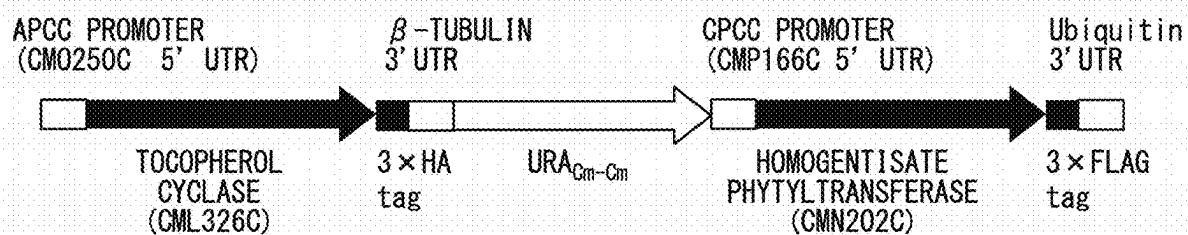
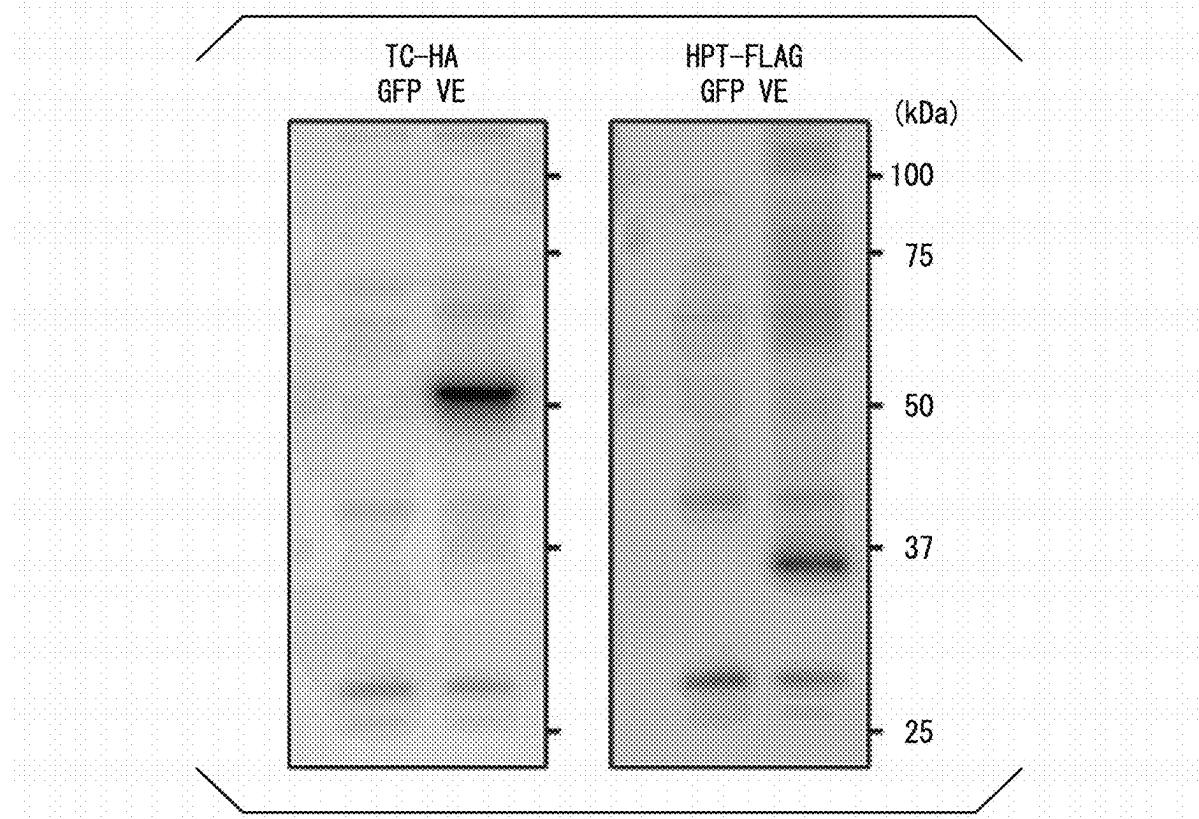


FIG. 22



NOVEL MICROALGAE AND USE FOR SAME

CROSS-REFERENCE TO RELATED APPLICATIONS

[0001] This application is a divisional application of U.S. Nonprovisional application Ser. No. 16/766,946 filed on Oct. 27, 2020, which is a US national phase application of International Application No. PCT/JP2018/043696 filed on Nov. 28, 2018, which claims the benefit of and priority to (1) Japanese Patent Application No. 2017-228396 filed on Nov. 28, 2017, (2) Japanese Patent Application No. 2017-228394 filed on Nov. 28, 2017, (3) Japanese Patent Application No. 2018-177416 filed on Sep. 21, 2018, and (4) Japanese Patent Application No. 2018-101753 filed on May 28, 2018, the contents of which are incorporated by reference herein in their entireties.

CROSS REFERENCE TO SEQUENCE LISTING

[0002] The genetic components described herein are referred to by sequence identifier numbers (SEQ ID NO). The sequence listing in written computer readable format (CRF) as an xml file named “51966-1060_Sequence_Listing.xml” created on Apr. 17, 2025, and having a size of 88,922 bytes, is incorporated by reference in its entirety.

TECHNICAL FIELD

[0003] The present invention relates to novel microalgae and use for the same. More specifically, the present invention relates to novel microalgae, a method for producing the microalgae as haploids, and the like. The present invention also relates to a nutrient composition and a method for producing a nutrient.

[0004] The present application claims priorities based on Japanese Patent Application Nos. 2017-228394 and 2017-228396 filed in Japan on Nov. 28, 2017, Japanese Patent Application No. 2018-101753 filed in Japan on May 28, 2018, and Japanese Patent Application No. 2018-177416 filed in Japan on Sep. 21, 2018, the contents of which are incorporated herein by reference.

BACKGROUND ART

[0005] Microalgae have a carbon dioxide fixation capacity higher than that of terrestrial plants and do not compete with agricultural products for their growing places. Therefore, some species of the microalgae are cultured in large quantities and used industrially as feed, functional foods, cosmetic materials, and the like.

[0006] In a case where the microalgae are used industrially, in view of costs and the like, microalgae that can be cultured outdoors in large quantities are desirable. However, the microalgae that can be cultured outdoors in large quantities need to meet requirements such as being resistant to environmental changes (light, temperature, and the like), being cultivable under conditions where other organisms cannot survive, and being able to grow to high density. Accordingly, hitherto, only a few species such as *Chlorella*, *Euglena*, *Dunaliella*, and *Spirulina* have been practically used in industry.

[0007] One of the characteristic of the aforementioned species of algae is that they can be cultured in an environment, such as an environment with a high salt concentration, high pH, and low pH, where other organisms are difficult to

grow. These species of algae are rich in amino acids and vitamins and are used as raw materials for functional foods or supplements.

[0008] Meanwhile, the algae belonging to Cyanidiophyceae, which are unicellular primitive red algae, grow preferentially in acidic hot springs containing sulfuric acid. Cyanidiophyceae includes *Cyanidioschyzon*, *Cyanidium*, and *Galdieria* (Non-Patent Document 1). As haploid algae, only *Cyanidioschyzon merolae* is known (Non-Patent Document 2). *Cyanidioschyzon merolae* has no rigid cell wall (Non-Patent Documents 1 and 2).

[0009] *Cyanidioschyzon merolae* is constituted with a very simple set of cell organelles, and the genome sequence thereof has been completely decoded. Therefore, *Cyanidioschyzon merolae* is used as a model organism for basic research on photosynthetic organisms, and the development of techniques for modifying the gene thereof is also making progress (Non-Patent Documents 3 and 4).

CITATION LIST

Non-Patent Literature

[0010] [Non-Patent Document 1]

[0011] Pinto, G. (2007) Cyanidiophyceae: looking back-looking forward, In: J. Seckbach (ed.) *Algae and Cyanobacteria in Extreme Environments*. Springer, Dordrecht, The Netherlands, pp. 389-397.

[Non-Patent Document 2]

[0012] Misumi O et al. (2005) *Cyanidioschyzon merolae* Genome. A Tool for Facilitating Comparable Studies on Organelle Biogenesis in Photosynthetic Eukaryotes. *Plant Physiol.* 137 (2): 567-585.

[Non-Patent Document 3]

[0013] Fujiwara T et al. (2013) Gene targeting in the red alga *Cyanidioschyzon merolae*: single- and multi-copy insertion using authentic and chimeric selection markers. *PLOS ONE*. September 5; 8 (9): e73608.

[Non-Patent Document 4]

[0014] Fujiwara T et al. (2015) A nitrogen source-dependent inducible and repressible gene expression system in the red alga *Cyanidioschyzon merolae*. *Front Plant Sci.* August 26; 6:657.

SUMMARY OF INVENTION

Technical Problem

[0015] As described above, hitherto, microalgae practically used in industry have been limited to several species of algae.

[0016] Therefore, an object of the present invention is to provide novel microalgae that can be used industrially and a method of using the novel microalgae. Another object of the present invention is to provide a nutrient composition using microalgae rich in nutrients, and a method for producing a nutrient using the microalgae.

[0017] Still another object of the present invention is to provide a method for preparing haploid cells from diploid

algal cells, haploid cells prepared by the method, and a haploid algal cell population obtained by culturing the haploid cells.

[0018] Yet another object of the present invention is to provide a method for preparing diploid cells from haploid algal cells, diploid cells prepared by the method, and a diploid algal cell population obtained by culturing the diploid cells.

[0019] Another object of the present invention is to provide algal cells transformed by self-cloning or multiple self-cloning by using haploid algal cells.

Solution to Problem

[0020] The inventors of the present invention have found that among the algae belonging to Cyanidiophyceae, there are a generation of haploid cells and a generation of diploid cells. Furthermore, the inventors have found that among the algae belonging to Cyanidiophyceae, those having a rigid cell wall are diploid cells, and found a method for inducing cells having no rigid cell wall from the diploid cells having a rigid cell wall. Furthermore, the inventors have found that the cells having no rigid cell wall obtained by the above method are haploid cells. In addition, the inventors have found that the algae belonging to Cyanidiophyceae are rich in nutrients such as amino acids and vitamins. Based on these findings, the inventors of the present invention have accomplished the following invention.

[0021] The present invention includes the following aspects.

[0022] (1) An Alga belonging to Cyanidiophyceae having a diploid cell form and a haploid cell form.

[0023] (1-2) A cell population of the alga according to (1), consisting of a cell population of the diploid cell form, in which the alga belonging to Cyanidiophyceae.

[0024] (1-3) A cell population of the alga according to (1), consisting of a cell population of the haploid cell form, in which the alga belonging to Cyanidiophyceae.

[0025] (1-4) A cell population of the alga according to (1), in which the alga belong to Cyanidiophyceae, and a cell population of the diploid cell form and a cell population of the haploid cell form are mixed together.

[0026] (2) The alga according to (1), in which the haploid cell form is ruptured under a condition of pH 7.

[0027] (3) The alga according to (1) or (2), in which a base sequence of a ribulose 1,5-bisphosphate carboxylase/oxygenase large subunit gene has identity equal to or higher than 90% with a base sequence shown in SEQ ID NO: 1 or 2.

[0028] (4) The alga according to (3), which is selected from the group consisting of *Cyanidium* sp. YFU3 (FERM BP-22334), *Cyanidium* sp. HKN1 (FERM BP-22333), and a mutant of these.

[0029] (5) The alga according to any one of (1) to (4), which is a transformant.

[0030] (6) The alga according to (5), in which the transformant is prepared by self-cloning.

[0031] (7) The alga according to any one of (1) to (6), which is the haploid cell form.

[0032] (8) A method for producing a haploid alga, including (a) step of culturing a diploid cell (alga) of the alga according to any one of (1) to (6), and (b) step of isolating a haploid cell (alga) that are generated during the culturing.

[0033] (9) The method for producing a haploid alga according to (8), in which the culturing in the step (a) is performed under a condition of temperature of 30° C. to 50° C., pH of 1.0 to 5.0, and CO₂ concentration of 1% to 3%.

[0034] (10) An alga culture containing the alga according to any one of (1) to (7), wherein a ratio of the number of haploid cells of the alga to the total number of cells of alga contained in the alga culture is 70% to 100%.

[0035] (11) A drying and swelling treatment product, which is obtained by subjecting the alga according to any one of (1) to (7) to a drying and swelling treatment.

[0036] The present invention also includes the following aspects.

[0037] (12) A method for producing a nutrient, including (a) step of disrupting a cell of the alga according to any one of (1) to (7) so as to obtain a cell disruption product, and (b) step of separating a nutrient from the cell disruption product.

[0038] (13) The method for producing a nutrient according to (12), in which the nutrient is at least one nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber.

[0039] (14) The method for producing a nutrient according to (13), in which the amino acids are at least one kind of amino acid selected from the group consisting of isoleucine, leucine, lysine, methionine, cysteine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, and γ-aminobutyric acid.

[0040] (15) The method for producing a nutrient according to (13), in which the vitamins are at least one kind of vitamin selected from the group consisting of vitamin A, β-carotene, vitamin C, vitamin E, vitamin K₁, and vitamin K₂.

[0041] (16) The method for producing a nutrient according to any one of (12) to (15), in which the disrupting of a cell in the step (a) is performed by at least one kind of treatment selected from the group consisting of a neutralization treatment, a hypotonic treatment, and a freeze-thaw treatment.

[0042] (17) The method for producing a nutrient according to any one of (12) to (16), further including a step of subjecting the alga according to any one of (1) to (7) to a low-temperature treatment at a temperature of 0° C. to 5° C. before the step (a).

[0043] (18) A nutrient composition containing the alga according to any one of (1) to (7) or an extract thereof.

[0044] (19) A food containing the nutrient composition according to (18).

[0045] (20) The food according to (19), which is a functional food or a dietary supplement.

[0046] (21) A feed or a pet food containing the nutrient composition according to (18).

[0047] (22) A cosmetic containing the nutrient composition according to (18).

[0048] (23) A nutritional supplement containing the alga according to any one of (1) to (7) or an extract thereof.

[0049] (24) A food containing the nutritional supplement according to (23).

- [0050] (25) The food according to (24), which is a functional food or a nutritional supplement.
- [0051] (26) A feed or a pet food containing the nutritional supplement according to (23).
- [0052] (27) A cosmetic containing the nutritional supplement according to (23).
- [0053] (28) A composition for supplying a nutrient, containing the nutritional supplement according to (23).
- [0054] (29) The composition for supplying a nutrient according to (28), which is a food.
- [0055] (30) The composition for supplying a nutrient according to (29), which is a functional food or a dietary supplement.
- [0056] (31) The composition for supplying a nutrient according to (28), which is a feed or a pet food.
- [0057] (32) The composition for supplying a nutrient according to (28), which is a cosmetic.
- [0058] (33) A nutrient composition containing the alga culture according to (10) or an extract thereof.
- [0059] (34) A food containing the nutrient composition according to (33).
- [0060] (35) The food according to (34), which is a functional food or a dietary supplement.
- [0061] (36) A feed or a pet food containing the nutrient composition according to (33).
- [0062] (37) A cosmetic containing the nutrient composition according to (33).
- [0063] (38) A nutritional supplement containing the alga culture according to (10) or an extract thereof.
- [0064] (39) A food containing the nutritional supplement according to (38).
- [0065] (40) The food according to (39), which is a functional food or a nutritional supplement.
- [0066] (41) A feed or a pet food containing the nutritional supplement according to (38).
- [0067] (42) A cosmetic containing the nutritional supplement according to (38).
- [0068] (43) A composition for supplying a nutrient, containing the nutritional supplement according to (38).
- [0069] (44) The composition for supplying a nutrient according to (43), which is a food.
- [0070] (45) The composition for supplying a nutrient according to (44), which is a functional food or a dietary supplement.
- [0071] (46) The composition for supplying a nutrient according to (43), which is a feed or a pet food.
- [0072] (47) The composition for supplying a nutrient according to (43), which is a cosmetic.
- [0073] (48) A method for producing a nutrient, including (a) step of collecting an alga from the alga culture according to (10), (b) step of disrupting a cell of the alga so as to obtain a cell disruption product, and (c) step of separating at least one kind of nutrient from the cell disruption product.
- [0074] (49) The method for producing a nutrient according to (48), in which the nutrient is at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber.
- [0075] (50) The method for producing a nutrient according to (49), in which the amino acids are at least one kind of amino acid selected from the group consisting of isoleucine, leucine, lysine, methionine, cysteine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, and γ -aminobutyric acid.
- [0076] (51) The method for producing a nutrient according to (49), in which the vitamins are at least one kind of vitamin selected from the group consisting of vitamin A, β -carotene, vitamin C, vitamin E, vitamin K₁, and vitamin K₂.
- [0077] (52) The method for producing a nutrient according to any one of (48) to (51), in which the disrupting of a cell in the step (a) is performed by at least one kind of treatment selected from the group consisting of a neutralization treatment, a hypotonic treatment, a freeze-thaw treatment, and a drying and swelling treatment.
- [0078] The present invention also includes the following aspects.
- [0079] (53) A nutrient composition containing an alga belonging to Cyanidiophyceae or an extract thereof.
- [0080] (54) The nutrient composition according to (53), in which the alga is a haploid alga.
- [0081] (55) The nutrient composition according to (53) or (54), in which the alga undergoes cell rupture under a condition of pH 7.
- [0082] (56) The nutrient composition according to any one of (53) to (55), in which the alga belongs to *Cyanidioschyzon*.
- [0083] (57) The nutrient composition according to any one of (53) to (56), in which the alga is a transformant having an increased intracellular content of at least one kind of nutrient.
- [0084] (58) The nutrient composition according to (57), in which the transformant is prepared by self-cloning.
- [0085] (59) The nutrient composition according to any one of (53) to (58), containing at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber.
- [0086] (60) A food containing the nutrient composition according to any one of (53) to (59).
- [0087] (61) The food according to (60), which is a functional food or a dietary supplement.
- [0088] (62) A feed or a pet food containing the nutrient composition according to any one of (53) to (59).
- [0089] (63) A cosmetic containing the nutrient composition according to any one of (53) to (59).
- [0090] (64) A method for producing a nutrient, including (a) step of disrupting a cell of an alga belonging to Cyanidiophyceae so as to obtain a cell disruption product, and (b) step of separating at least one kind of nutrient from the cell disruption product.
- [0091] (65) The method for producing a nutrient according to (64), in which the nutrient is at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber.
- [0092] (66) The method for producing a nutrient according to (65), in which the amino acids are at least one kind of amino acid selected from the group consisting of isoleucine, leucine, lysine, methionine, cysteine, phenylalanine, tyrosine, threonine, tryptophan,

valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, and γ -aminobutyric acid.

[0093] (67) The method for producing a nutrient according to (65), in which the vitamins are at least one kind of vitamin selected from the group consisting of vitamin A, β -carotene, vitamin B₁, vitamin B₂, vitamin B₆, vitamin C, vitamin E, vitamin K₁, vitamin K₂, niacin, inositol, folic acid, and biotin.

[0094] (68) The method for producing a nutrient according to any one of (64) to (67), in which the alga is a haploid alga.

[0095] (69) The method for producing a nutrient according to any one of (64) to (68), in which a cell of the alga is ruptured under a condition of pH 7.

[0096] (70) The method for producing a nutrient according to any one of (64) to (69), in which the alga belong to *Cyanidioschyzon*.

[0097] (80) The method for producing a nutrient according to any one of (64) to (70), in which the alga is a transformant having an increased intracellular content of at least one kind of nutrient.

[0098] (81) The method for producing a nutrient according to (80), in which the transformant is prepared by self-cloning.

[0099] (82) The method for producing a nutrient according to any one of (64) to (81), in which the disrupting of a cell in the step (a) is performed by at least one kind of treatment selected from the group consisting of a neutralization treatment, a hypotonic treatment, a freeze-thaw treatment, and a drying and swelling treatment.

[0100] The present invention also includes the following aspects.

[0101] (83) A nutritional supplement containing an alga belonging to *Cyanidiophyceae* or an extract thereof.

[0102] (84) The nutritional supplement according to (83), in which the alga is a haploid alga.

[0103] (85) The nutritional supplement according to (83) or (84), in which a cell of the alga is ruptured under a condition of pH 7.

[0104] (86) The nutritional supplement according to any one of (83) to (85), in which the alga belong to *Cyanidioschyzon*.

[0105] (87) The nutritional supplement according to any one of (83) to (86), in which the alga is a transformant having an increased intracellular content of at least one kind of nutrient.

[0106] (88) The nutritional supplement according to (87), in which the transformant is prepared by self-cloning.

[0107] (89) The nutritional supplement according to any one of (83) to (88), which is for supplying at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber.

[0108] (90) A food containing the nutritional supplement according to any one of (83) to (89).

[0109] (91) The food according to (90), which is a functional food or a dietary supplement.

[0110] (92) A feed or a pet food containing the nutritional supplement according to any one of (83) to (89).

[0111] (93) A cosmetic containing the nutritional supplement according to any one of (83) to (89).

[0112] (94) A composition for supplying a nutrient, containing the nutritional supplement according to any one of (83) to (89).

[0113] (95) The composition for supplying a nutrient according to (94), which is a food.

[0114] (96) The composition for supplying a nutrient according to (95), which is a functional food or a dietary supplement.

[0115] (97) The composition for supplying a nutrient according to (94), which is a feed or a pet food.

[0116] (98) The composition for supplying a nutrient according to (94), which is a cosmetic.

[0117] (99) The composition for supplying a nutrient according to any one of (94) to (98), which is for supplying at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber.

[0118] The present invention also includes the following aspects.

[0119] (100) A method for producing a diploid alga, including (a) step of mixing together two or more kinds of haploid cells of the alga according to any one of (1) to (6) and culturing the mixture, and (b) step of isolating a diploid cell generated during the culturing.

[0120] (101) The method for producing a diploid alga according to (100), in which the alga is selected from the group consisting of *Cyanidium* sp. YFU3 (FERM BP-22334), *Cyanidium* sp. HKN1 (FERM BP-22333), *Cyanidioschyzon merolae*, and mutants of these.

[0121] (102) A diploid cell of *Cyanidium* sp. YFU3 (FERM BP-22334) or a mutant thereof.

[0122] (103) A diploid cell of a *Cyanidium* sp. HKN1 (FERM BP-22333) or a mutant thereof.

[0123] (104) A diploid cell of *Cyanidioschyzon merolae*.

[0124] (105) The method for producing a haploid alga according to (8), in which the alga is selected from the group consisting of alga belonging to *Galdieria* and alga belonging to *Cyanidium*.

[0125] (106) A haploid cell of alga belonging to *Galdieria*.

[0126] (107) The haploid cell according to (106), in which the alga belonging to *Galdieria* are *Galdieria sulphuraria* or *Galdieria partita*.

[0127] (108) A haploid cell of an alga belonging to *Cyanidium*.

[0128] (109) The haploid cell according to (108), in which the alga belonging to *Cyanidium* are *Cyanidium* sp. YFU3 (FERM BP-22334) or *Cyanidium* sp. HKN1 (FERM BP-22333).

Advantageous Effects of Invention

[0129] According to the present invention, there are provided novel microalgae which can be industrially used and a method of using the microalgae. Furthermore, according to the present invention, there are provided a nutrient composition using microalgae rich in nutrients, and a method for producing a nutrient by using the microalgae.

[0130] In addition, according to the present invention, there are provided a method for preparing haploid cells from diploid algal cells, haploid cells prepared by the method, and a haploid algal cell population obtained by culturing the haploid cells.

[0131] Moreover, according to the present invention, there are provided a method for preparing diploid cells from haploid algal cells, diploid cells prepared by the method, and a diploid algal cell population obtained by culturing the diploid cells.

[0132] The present invention also provides algal cells transformed by self-cloning or multiple self-cloning by using haploid algal cells.

BRIEF DESCRIPTION OF DRAWINGS

[0133] FIG. 1A shows the construct of a fragment for transformation (EGFP/URA_{Cm-Cm} fragment) used in Example 2.

[0134] FIG. 1B shows the construct of a fragment for transformation (GAD/URA_{Cm-Cm} fragment) used in Example 2.

[0135] FIG. 1C shows the construct of a fragment for transformation (GAD/URA_{Cm-Gs} fragment) used in Example 2.

[0136] FIG. 2 shows the results of immunoblotting performed using transformants prepared in Example 2. The right figure shows gel stained after SDS-PAGE. The left figure shows the results of immunoblotting using an anti-HA antibody.

[0137] FIG. 3 shows the results of immunoblotting in which a transformant with GAD/URA_{Cm-Cm} fragment prepared in Example 2 is compared with a wild-type strain (WT).

[0138] FIG. 4 shows the results obtained by confirming the number of copies of a glutamate decarboxylase gene in a transformant with GAD/URA_{Cm-Gs} fragment prepared in Example 2.

[0139] FIG. 5 shows the results of SDS-polyacrylamide electrophoresis performed on a centrifugal supernatant and a centrifugal precipitate which are obtained by performing a drying and swelling treatment on *Cyanidioschyzon merolae* 10D and *Cyanidium Caldarium* RK-1 and performing centrifugation on cell suspensions obtained after the swelling treatment.

[0140] FIG. 6A is a picture of microalgae (*Cyanidium* sp. HKN1) isolated from a hot spring in Hakone-machi, Ashigarashimo-gun, Kanagawa prefecture, Japan. The picture shows *Cyanidium* sp. HKN1 in a culture solution that is in the stationary phase after being cultured in MA medium. The arrows indicate *Cyanidioschyzon merolae*-like cells that appeared in the stationary phase.

[0141] FIG. 6B is a picture of microalgae (*Cyanidium* sp. HKN1) isolated from a hot spring in Hakone-machi, Ashigarashimo-gun, Kanagawa prefecture, Japan. The picture shows *Cyanidioschyzon merolae*-like cells isolated from the culture solution in the stationary phase in FIG. 6A.

[0142] FIG. 7 shows the results of sequence analysis on one genomic region of *Cyanidium* sp. HKN1 and the *Cyanidioschyzon merolae*-like cells isolated from the culture solution of *Cyanidium* sp. HKN1 in the stationary phase. It has been confirmed that the *Cyanidioschyzon merolae*-like cells are haploids (HKN1 (haploid)) generated by the meiosis of the *Cyanidium* sp. HKN1 (diploid).

[0143] FIGS. 8A-8E show micrographs of algae belonging to Cyanidiophyceae. (A) is a micrograph of *Galdieria sulphuraria* 074, (B) is a micrograph of *Cyanidium caldarium* RK-1, (C) is a micrograph of *Cyanidioschyzon merolae* 10D,

(D) is a micrograph of YFU3 (haploid), and (E) is a micrograph of HKN1 (haploid). The scale bar represents 5 μm.

[0144] FIG. 9 shows the results of agarose electrophoresis in which YFU3 (haploid), HKN1 (haploid), and *Cyanidioschyzon merolae* are compared with each other in terms of the size of ribosomal DNA ITS1.

[0145] FIG. 10 shows a molecular phylogenetic tree of algae belonging to Cyanidiophyceae based on a chloroplast ribulose 1,5-bisphosphate carboxylase/oxygenase large subunit gene. Near each branch, a local bootstrap value (only a value equal to or greater than 50 is described, left side) determined by the maximum likelihood method and a posterior probability (only a probability equal to or higher than 0.95 is described, right side) determined by the Bayesian method are shown. Known *Cyanidioschyzon merolae* are boxed in a dotted line, and YFU3 and HKN1 are boxed in a solid line.

[0146] FIG. 11 shows the chloramphenicol resistance of chloramphenicol-resistant transformants of YFU3 (haploid) and *Cyanidioschyzon merolae* 10D prepared in Example 6.

[0147] FIG. 12 shows the results of PCR performed to confirm whether a CAT gene has been inserted into the genome of the chloramphenicol-resistant transformants of YFU3 (haploid) and *Cyanidioschyzon merolae* 10D prepared in Example 6.

[0148] FIG. 13 shows the construct of a CAT vector for transformation used in Example 7 (upper figure: PCR product) and the insertion position of the vector in the genome of HKN1 (haploid) (lower figure: Genome). In the lower figure (Genome), the arrows indicate the position of a primer used to check whether an m Venus-CAT gene has been inserted into the genome of the chloramphenicol-resistant transformant.

[0149] FIG. 14 shows the results of PCR and agarose gel electrophoresis performed to confirm whether the m Venus-CAT gene has been inserted into the genome of the chloramphenicol transformant prepared in Example 7.

[0150] FIG. 15 shows fluorescence micrographs of the chloramphenicol transformant prepared in Example 7. The left figure (m Venus) is a fluorescence micrograph in which the fluorescence of m Venus is detected, the middle figure (Chl) is a fluorescence micrograph in which the autofluorescence of the chloroplast is detected, and the right figure (merged) is an image obtained by merging the two fluorescence micrographs.

[0151] FIG. 16 shows the results of SDS-polyacrylamide electrophoresis performed on centrifugal supernatants and centrifugal precipitates which are obtained by performing a drying and swelling treatment on algae having no rigid cell wall and algae having a cell wall and performing centrifugation on cell suspensions obtained after the swelling treatment.

[0152] FIG. 17A is a micrograph of diploid-like cells having a rigid cell wall that are obtained by crossing HKN1 cells (haploids).

[0153] FIG. 17B shows the results of DAPI staining performed on the diploid-like cells having a rigid cell wall and HKN1 (haploid).

[0154] FIG. 18A shows a micrograph (left figure) of normal cells (diploids) of *Galdieria sulphuraria* SAG108. 79 and a micrograph (right figure) of the same strain in the form of cells having no rigid cell wall.

[0155] FIG. 18B shows a micrograph (left figure) of normal cells (diploids) of *Galdieria partita* NBRC102759 which is in the normal cell form (diploid) and a micrograph (right figure) of the same strain in the form of cells having no rigid cell wall.

[0156] FIG. 19 shows the results of sequence analysis on one genomic region of *Galdieria sulphuraria* SAG108.79 in the form of cells having no rigid cell wall. 2N_allele1 (SEQ ID NO: 61) and 2N_allele2 (SEQ ID NO: 62) represent the sequences of alleles of *Galdieria sulphuraria* SAG108.79 in the normal cells (diploids). N_clone1, N_clone2, and N_clone3 represent the sequences of alleles confirmed in three strains in the form of cells having no rigid cell wall.

[0157] FIG. 20 shows the results of SDS-polyacrylamide electrophoresis performed on centrifugal supernatants and centrifugal precipitates which are obtained by performing a drying and swelling treatment on diploid cells and haploid cells of *Galdieria sulphuraria* SAG108.79 and *Galdieria partita* NBRC102759 and performing centrifugation on cell suspensions obtained after the swelling treatment.

[0158] FIG. 21 shows the construct of a fragment for transformation (VE/URA_{Cm-Cm} fragment) used in Example 12.

[0159] FIG. 22 shows the results of immunoblotting performed using a transformant prepared in Example 12. The left figure (TC-HA) shows the results of immunoblotting using an anti-HA antibody, and the right figure (HPT-FLAG) shows the results of immunoblotting using an anti-FLAG antibody.

DESCRIPTION OF EMBODIMENTS

[Algae Belonging to Cyanidiophyceae]

[0160] As shown in Examples which will be described later, it has been found that the algae belonging to Cyanidiophyceae are rich in nutrients such as amino acids, vitamins, proteins, lipids, and dietary fiber. Therefore, the present invention provides a nutrient composition, a nutritional supplement, a food, a feed, a pet food, a pet food, a cosmetic, and the like using the algae belonging to Cyanidiophyceae. Furthermore, the present invention provides a method for producing a nutrient using the algae belonging to Cyanidiophyceae.

[0161] Taxonomically, Cyanidiophyceae is classified into Rhodophyta, Cyanidiophyceae. Currently, three genera of *Cyanidioschyzon*, *Cyanidium*, and *Galdieria* are classified into Cyanidiophyceae. In the nutrient composition according to an embodiment of the present invention and the like may use algae belonging to any of the above genera.

[0162] Whether or not certain algae belong to Cyanidiophyceae can be determined, for example, by phylogenetic analysis using the base sequence of an 18S rRNA gene or a chloroplast rbcL gene. The phylogenetic analysis may be performed by a known method.

[0163] In the present specification, "haploid cell form" means that the cell as an observation target has one set of genetic information, and "diploid cell form" means that the cell as an observation target has two sets of genetic information. Whether a certain cell is haploid or diploid can be determined by measuring the content of DNA in the cell, and also can be determined by a next-generation sequencer which will be described later.

[0164] In the present specification, "having a diploid cell form and a haploid cell form" means that a cell of an alga

belonging to Cyanidiophyceae collected from nature is the haploid cell form in a case, and is the diploid cell form in another case, and also means that the cell of an alga belonging to Cyanidiophyceae which is unknown hitherto and created by the method of the present invention is the haploid cell form in a case and is the diploid cell form in another case.

[0165] In the present specification, "cell population of diploid cell form" and "cell population of haploid cell form" may be a clonal cell population or a non-clonal cell population.

[0166] Among the algae belonging to Cyanidiophyceae, *Cyanidioschyzon merolae* is known to have no rigid cell wall. Most of the algae belonging to Cyanidiophyceae other than *Cyanidioschyzon merolae* have been found as algal cells having a rigid cell wall. As shown in Examples which will be described later, it has been confirmed that these cells having a rigid cell wall are diploids. However, by the method according to an embodiment of the present invention that will be described later, it is possible to create a haploid cell form of algal cells, and many of the created haploid algal cells do not have a rigid cell wall. The haploid cell form of algal cells that do not have a rigid cell wall can be disrupted by a relatively mild treatment such as a neutralization treatment, a hypotonic treatment, or a freeze-thaw treatment. In the present specification, "having no rigid cell wall" means that a cell is ruptured by any of the following cell rupturing treatments (A) to (C).

[0167] (A) Algal cells are suspended in an isotonic solution at pH 7 and left for one week or longer.

[0168] (B) Algal cells are suspended in distilled water and left for 1 minute or longer.

[0169] (C) Algal cells are subjected to a drying treatment and suspended in an isotonic solution at pH 7.

[0170] In the above (A) to (C), in a case where the algal cells are cultured cells, the medium may be removed by centrifugation or the like before each treatment is performed, and the algal cells may be washed with an isotonic solution or the like.

[0171] In the above (A) and (C), examples of the isotonic solution include a pH 7 buffer containing 10% sucrose and 20 mM HEPES.

[0172] In the above (C), examples of the drying treatment include drying in a refrigerator (4° C.), freeze-drying, and the like. For the drying treatment, the precipitate of algal cells collected by centrifugation is used. In a case where the cells are dried in a refrigerator, the drying treatment time depends on the amount of algal cells, and is, for example, 3 days or longer.

[0173] Whether or not the cells rupture has occurred can be determined by performing centrifugation (1,500×g, 3 minutes) on the algal cell suspension after the cell rupturing treatments (A) to (C) described above, and calculating a ratio of the mass of proteins in the centrifugal supernatant to the total mass of proteins in the algal cell suspension. Specifically, in a case where a rupture rate determined by the following equation is equal to or higher than 20%, it can be determined that cell rupture has occurred.

[Equation 1]

Rupture rate =

$$\frac{\text{Protein content in centrifugal supernatant (mass g)}}{\text{Total protein content in algal suspension (mass g)}} \times 100(%)$$

[0174] Alternatively, in a case where the algal cells in the algal cell suspension are observed using an optical microscope (for example, at a magnification of 600 \times), and the ratio of ruptured cells to the total number of the algal cells is found to be about equal to or higher than 10% and preferably about equal to or higher than 20%, it may be determined that cell rupture has occurred.

[0175] In the cell rupturing treatments (A) to (C) described above, an isotonic solution at pH 7 can be used. Accordingly, it can be said that the cells ruptured by any of the cell rupturing treatments (A) to (C) are cells that are ruptured under the condition of pH 7.

[0176] Not only the algae belonging to *Cyanidioschyzon*, but also the algae that undergo cell rupture under the condition of pH 7 are preferable because it is easy to extract nutrients from such algae. In addition, such algae also have an advantage that even though the algal cells are added as themselves to a nutritional supplement or the like which will be described later, the nutrients in the algal cells are easily absorbed after the intake of the nutritional supplement. Therefore, in the nutrient composition according to the present embodiment, the alga, which belongs to Cyanidiophyceae and undergo cell rupture under the condition of pH 7, can be suitably used.

[0177] Whether or not certain alga undergoes cell rupture under the condition of pH 7 can be determined by immersing the algal cell in a buffer at pH 7 or the like, observing the cell for about 10 to 30 minutes, and checking whether or not the algal cells are ruptured.

[0178] In a case where the algal cell has no a rigid cell wall, by the observation using an optical microscope (for example, at a magnification of 600 \times), generally, a cell wall is not observed.

[0179] Generally, *Cyanidioschyzon merolae* does not undergo cell rupture even being subjected to a mild hypotonic treatment under the condition of pH equal to or lower than 6. Therefore, whether or not cell rupture occurs due to the mild hypotonic treatment under the condition of pH equal to or lower than 6 does not exert an influence on determining whether or not the alga has a rigid cell wall.

[0180] Among the algae belonging to Cyanidiophyceae, the algae belonging to *Cyanidioschyzon*, particularly, *Cyanidioschyzon merolae*, is characterized by being a haploid. Furthermore, these algae have no rigid cell wall. Accordingly, the algae belonging to *Cyanidioschyzon*, particularly, *Cyanidioschyzon merolae*, can be relatively easily transformed using gene recombination techniques. For example, as shown in Examples which will be described later, by using a gene recombination technique, it is possible to prepare a transformant having an increased intracellular content of a nutrient.

[0181] Not only the algae belonging to *Cyanidioschyzon*, particularly, *Cyanidioschyzon merolae*, but also haploid algae are preferable because these algae can be relatively easily transformed. For example, the haploid cells of algae, which belong to Cyanidiophyceae having a diploid cell form and a haploid cell form which will be described later, is preferable because these algae can be relatively easily transformed. Therefore, in the nutrient composition and the like according to the embodiment of the present invention, the haploid algae belonging to Cyanidiophyceae can be suitably used.

[0182] Whether certain algae are haploid can be determined by checking the number of copies of the same loci.

That is, in a case where the copy number of the same loci is 1, the algae are determined to be haploid.

[0183] Furthermore, for example, whether certain algae are haploid can also be determined using a next-generation sequencer or the like. For example, whole genome sequence reads are acquired by a next-generation sequencer or the like, the sequence reads are assembled, and then the sequence reads are mapped for the sequence obtained by the assembling. In diploids, the difference in base between alleles are found in various regions on the genome. However, in haploids, because only one allele is present, no such regions are found.

[0184] Furthermore, in a case where certain algae are homodiploid, whether the algae are haploid or diploid can be determined by measuring the DNA content in the cells. The DNA content in haploid cells is one half of the DNA content in diploid cells.

[0185] Examples of the haploid algae, which belong to Cyanidiophyceae and undergo cell rupture under the condition of pH 7, include algae belonging to *Cyanidioschyzon*. Currently, only one species of *Cyanidioschyzon merolae* is classified into *Cyanidioschyzon*. Therefore, *Cyanidioschyzon merolae* is a suitable example of algae used in the nutrient composition or the like according to the embodiment of the present invention.

[0186] In addition, not only the haploid of algae belonging to *Cyanidioschyzon* other than *Cyanidioschyzon merolae* that is obtained by the present invention and is unknown hitherto, but also the haploid of algae belonging to *Galdieria* and the haploid of algae belonging to *Cyanidium* are suitable examples as well.

[0187] The algae belonging to Cyanidiophyceae can grow to high density (about 30 g/L) under high temperature (30° C. to 55° C.) and acidic (pH 0.1 to 3.0) conditions. Under such a high-temperature and acidic environment, other organisms are difficult to grow. Therefore, the algae belonging to Cyanidiophyceae are suitable for being cultured outdoors in large quantities.

[0188] In addition, the algae belonging to Cyanidiophyceae can grow not only under high-temperature and acidic conditions but also under a relatively wide range of environmental conditions such as a medium-temperature environment (15° C. to 30° C.) and a neutral environment (about pH 6). Therefore, the culture conditions can be changed according to the region and the season. In this respect, the algae belonging to Cyanidiophyceae are also suitable for being cultured outdoors in large quantities.

[0189] Furthermore, the algae belonging to Cyanidiophyceae exhibits high salt tolerance and can grow under high salt (300 mM NaCl) conditions. In addition, the range of optical intensity at which the algae can grow is wide. The algae can grow to high density in a range of 5 to 1,500 $\mu\text{mol}/\text{m}^2\text{s}$, and can grow under strong light.

[0190] As will be shown in Examples which will be described later, the algae belonging to Cyanidiophyceae are rich in nutrients such as amino acids, vitamins, proteins, lipids, and dietary fiber. Particularly, it has been confirmed that the algae belonging to Cyanidiophyceae contain amino acids and vitamins at a higher concentration compared to the conventionally used algae (*Chlorella*, *Euglena*, and *Spirulina*). Furthermore, the composition of components of the algae is also specific. Therefore, by using the algae belonging to Cyanidiophyceae, it is possible to manufacture a nutrient composition, a nutritional supplement, and the like.

rich in nutrients selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber, particularly, nutrients such as amino acids or vitamins.

[0191] In the present specification, "amino acids" mean organic compounds having an amino group and a carboxy group.

[0192] Examples of the amino acids contained in the algae belonging to Cyanidiophyceae include isoleucine, leucine, lysine, methionine, cystine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, γ -aminobutyric acid, and the like. One of the characteristics of the algae belonging to Cyanidiophyceae is that the total amino acid content thereof is higher than that of the conventionally used algae. Furthermore, the content of each of the amino acids in the algae belonging to Cyanidiophyceae tends to be high overall.

[0193] For example, the total content of amino acids in the algae belonging to Cyanidiophyceae is 60 to 80 g/100 g based on dry weight. Furthermore, algae belonging to Cyanidiophyceae are also characterized by containing γ -aminobutyric acid. For example, the content of γ -aminobutyric acid in the algae belonging to Cyanidiophyceae is 0.1 to 0.3 g/100 g based on dry weight. In the present specification, "total amino acid content" means a value obtained by adding up the contents of isoleucine, leucine, lysine, methionine, cystine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, and serine.

[0194] In the present specification, "vitamins" mean organic compounds other than carbohydrates, proteins, and lipids among nutrients necessary for the survival of an organism.

[0195] Examples of the vitamins contained in the algae belonging to Cyanidiophyceae include vitamin A, β -carotene, vitamin B₁, vitamin B₂, vitamin B₆, vitamin C, vitamin E, vitamin K₁, vitamin K₂, niacin, inositol, folic acid, biotin, and the like. One of the characteristics of the algae belonging to Cyanidiophyceae is that the content of β -carotene, vitamin C, vitamin E, vitamin K₁, vitamin K₂, and folic acid among the above vitamins is particularly higher in the algae than in the conventionally used algae.

[0196] Regarding the content of each of the vitamins in the algae belonging to Cyanidiophyceae, for example, the content of β -carotene is 200 to 250 mg/100 g based on dry weight, the content of vitamin C is 40 to 70 mg/100 g based on dry weight, the content of vitamin E is 150 to 180 mg/100 g based on dry weight, the content of vitamin K₁ is 3,000 to 5,000 μ g/100 g based on dry weight, the content of vitamin K₂ is 4,000 to 7,000 μ g/100 g based on dry weight, and the content of folic acid is 3,500 to 6,500 μ g/100 g based on dry weight.

[0197] In a case where a human being or a non-human animal ingests algae which have no rigid cell wall among the algae belonging to Cyanidiophyceae, the nutrients described above can be efficiently absorbed into the body of the human being or the non-human animal. Furthermore, in a case where an extract containing the nutrients is prepared from the algae belonging to Cyanidiophyceae, from the algae which have no rigid cell wall, the extract can be prepared by a simple operation such as the cell rupturing treatments (A) to (C) described above.

[0198] The algae belonging to Cyanidiophyceae can preferentially grow in an acidic environment of sulfuric acid

such as an acidic hot spring containing sulfuric acid. Therefore, the algae can be obtained by being isolated from the acidic hot spring containing sulfuric acid and the like. In addition, the algae belonging to Cyanidiophyceae may be obtained from a culture collection and the like. For example, *Cyanidioschyzon merolae* can be obtained from the National Institute for Environmental Studies MICROBIAL CULTURE COLLECTION (16-2 Onogawa, Tsukuba-shi, Ibaraki prefecture, Japan), American Type Culture Collection (ATCC; 10801 University Boulevard Manassas, VA 20110 USA), and the like.

[0199] The algae belonging to Cyanidiophyceae can be cultured using a medium for culturing microalgae. The medium is not particularly limited, and examples thereof include an inorganic salt medium containing a nitrogen source, a phosphorus source, trace elements (such as zinc, boron, cobalt, copper, manganese, molybdenum, and iron), and the like. Examples of the nitrogen source include an ammonium salt, a nitrate, and a nitrite, and the like. Examples of the phosphorus source include a phosphate and the like. Examples of the aforementioned medium include a 2x Allen medium (Allen M B. Arch. Microbiol. 1959 32:270-277.), an M-Allen medium (Minoda A et al. Plant Cell Physiol. 2004 45:667-71)), an MA2 medium (Ohnuma M et al. Plant Cell Physiol. 2008 January; 49 (1): 117-20.), a modified M-Allen medium, and the like.

[0200] As described above, the algae belonging to Cyanidiophyceae can be grown to high density under a relatively wide range of culture conditions. The pH condition can be, for example, pH 1.0 to 6.0, and is preferably pH 1.0 to 5.0. In a case where the algae are cultured outdoors, in order to prevent the growth of other organisms, it is preferable to culture the algae under conditions with a high acidity. The pH of such conditions is, for example, 1.0 to 3.0.

[0201] The temperature condition can be, for example, 15° C. to 50° C., and is preferably 30° C. to 50° C. In a case where the algae are cultured outdoors, in order to prevent the growth of other organisms, it is preferable to culture the algae at a high temperature. The temperature of such condition is, for example, 35° C. to 50° C.

[0202] The light intensity can be, for example, 5 to 2,000 μ mol/m²s, and is preferably 5 to 1,500 μ mol/m²s. In a case where the algae are cultured outdoors, the algae may be cultured under sunlight. In a case where the algae are cultured indoors, the algae may be cultured under continuous light or go through a light-dark cycle (10L:14D or the like).

[0203] By being collected through a known method such as centrifugation or filtration and subjected to washing, drying, and the like as appropriate, the algae belonging to Cyanidiophyceae grown by culturing can be used in the nutritional supplement of the present embodiment.

[0204] The algae belonging to Cyanidiophyceae are not limited to those isolated from the natural world, and may be algae obtained by the mutation of natural algae belonging to Cyanidiophyceae. The mutation may be spontaneous mutation or induced mutation. For example, *Cyanidioschyzon merolae* isolated from the natural world is haploid cells. As the genome size of *Cyanidioschyzon merolae* is small (about 16 Mbp) and the genomic sequence thereof has been completely decoded (Matsuzaki M et al., Nature. 2004 Apr. 8; 428 (6983): 653-7.), *Cyanidioschyzon merolae* can be easily genetically modified. Therefore, a transformant produced by genetically modifying *Cyanidioschyzon merolae* is a suit-

able example of the algae belonging to Cyanidiophyceae. In addition, not only *Cyanidioschyzon merolae*, but also the transformant of the algae belonging to Cyanidiophyceae may be used in the nutrient composition and the like according to the embodiment of the present invention as algae belonging to Cyanidiophyceae, as long as the transformant can be genetically modified. For example, as will be described later, although the algae belonging to *Galdieria* and the algae belonging to *Cyanidium* isolated from nature are diploid cells having a rigid cell wall, these can be made into haploid cells by the method of the present invention. Therefore, these algae have a haploid cell form and a diploid cell form. Cells in any form may be used as a cell population. Among those haploid algal cells, the cells that do not have a rigid cell wall just as *Cyanidioschyzon merolae* can be easily genetically modified. Accordingly, the transformant, which is obtained by genetically modifying haploid cells of the algae belonging to *Galdieria* or *Cyanidium* or cells having no rigid cell wall, is also a suitable example of the algae belonging to Cyanidiophyceae. More specifically, in addition to *Cyanidioschyzon merolae*, it is preferable to use haploids of *Galdieria*, haploids of *Cyanidium*, haploids of *Cyanidioschyzon* other than *Cyanidioschyzon merolae*, and haploids of *Cyanidium*. Furthermore, it is more preferable to use the cells which do not have a rigid cell wall among the above. Moreover, diploids belonging to Cyanidiophyceae which do not have a rigid cell wall are also preferable.

[0205] The transformant of the algae belonging to Cyanidiophyceae is not particularly limited, and examples thereof include a transformant having an increased intracellular content of at least one kind of nutrient. As the nutrient of which the intracellular content is to be increased, for example, at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber is suitable. That is, for example, a transformant of the algae belonging to Cyanidiophyceae is preferable, which has a higher intracellular content of at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber compared to a wild-type strain. In the present specification, "wild-type strain" means an original cell that is a target of transformation but has not yet been transformed.

[0206] Among the above nutrients, at least one kind of nutrient selected from the group consisting of amino acids and vitamins is preferable. Such transformant has a higher intracellular content of a specific nutrient compared to the wild-type strain. Therefore, by using the transformant, it is possible to prepare a nutrient composition, a nutritional supplement, and the like rich in the nutrient. The method for increasing the intracellular content of the nutrient is not particularly limited.

[0207] For example, the algae may be genetically modified such that the expression amount of an enzyme catalyzing any of the reactions in the synthesis pathway of the nutrient of interest is increased. As the genetic modification method for increasing the expression amount of the enzyme, known methods can be used without particular limitation. Examples of such a method include a method of introducing a gene encoding the enzyme (hereinafter, referred to as "enzyme gene"), a method of introducing a gene encoding a factor that promotes the expression of the enzyme gene, and a method of disrupting a gene encoding a factor that inhibits the expression of the enzyme gene, and the like. In a case where the enzyme gene is introduced into an algal

cell, instead of a promoter sequence of the enzyme gene, a promoter of a gene expressed in large amounts in the algal cell into which the gene will be introduced may be used. The enzyme is not particularly limited as long as it is an enzyme that catalyzes any of the reactions in the synthesis pathway of the nutrient of interest. Examples of the enzyme include a synthase of the nutrient of interest, a synthase of a precursor of the nutrient of interest, and the like. "Precursor of the nutrient" may be any of the compounds that are generated at the stage before the nutrient is synthesized in the synthesis pathway of the nutrient of interest.

[0208] The method for increasing the intracellular content of a nutrient may be, for example, genetic modification for reducing the expression amount of a factor (hereinafter, referred to as "inhibitory factor") that inhibits any of the reactions in the synthesis pathway of the nutrient of interest. As the method for reducing the expression amount of the inhibitory factor, known methods can be used without particular limitation. Examples of such a method include a method of disrupting a gene encoding the inhibitory factor, a method of introducing a gene encoding a factor that inhibits the expression of a gene encoding the inhibitory factor, a method of disrupting a gene encoding a factor that promotes the expression of a gene encoding the inhibitory factor, and the like.

[0209] For example, glutamate decarboxylase is a synthase of γ -aminobutyric acid. Therefore, by genetically modifying the algae belonging to Cyanidiophyceae such that the expression amount of the glutamate decarboxylase is increased, it is possible to obtain a transformant having an increased intracellular content of γ -aminobutyric acid. Homogentisate phytoltransferase and tocopherol cyclase are enzymes that catalyze reactions in a vitamin E biosynthesis pathway. That is, the homogentisate phytoltransferase is an enzyme that synthesizes 2-methyl-6-phytylbenzoquinone, and the tocopherol cyclase is an enzyme that synthesizes γ -tocotrienol. Therefore, by genetically modifying the algae belonging to Cyanidiophyceae such that the expression amount of either or both of the enzyme genes described above, it is possible to obtain a transformant having an increased intracellular content of vitamin E. In order to increase the intracellular content of vitamin E, the expression amount of an enzyme that catalyzes another reaction in the vitamin E biosynthesis pathway may be increased. Similarly, for other nutrients, by increasing the expression amount of an enzyme that catalyzes any of the reactions in the biosynthetic pathway of the nutrient of interest (preferably, a rate-limiting reaction in the biosynthetic pathway), the intracellular content of the nutrient can be increased.

[0210] The sequence information of these synthetic enzyme genes can be obtained from a known sequence database such as GenBank. The sequence of the glutamate decarboxylase gene (CMF072C) of *Cyanidioschyzon merolae* is shown in SEQ ID NO: 3, and the amino acid sequence thereof is shown in SEQ ID NO: 4.

[0211] The sequence of the homogentisate phytoltransferase gene (CMN202C) of *Cyanidioschyzon merolae* is shown in SEQ ID NO: 5, and the amino acid sequence thereof is shown in SEQ ID NO: 6. The sequence of the tocopherol cyclase gene (CML326C) of *Cyanidioschyzon merolae* is shown in SEQ ID NO: 7, and the amino acid sequence thereof is shown in SEQ ID NO: 8.

[0212] For example, by introducing a synthetic enzyme gene (for example, the glutamate decarboxylase gene) of the

aforementioned nutrient into the algae belonging to Cyanidiophyceae, it is possible to obtain a transformant having an increased intracellular content of the nutrient. The promoter used for the gene to be introduced may be the promoter of the synthetic enzyme gene or the promoter of another gene. In a case where the promoter of another gene is used, a promoter of a gene that is expressed in large amounts in the algal cell into which the gene is to be introduced is preferable. In the case of *Cyanidioschyzon merolae*, examples of such a promoter include a promoter of APCC (CMO250C) (for example, -600 to -1; “-1” indicates a nucleotide immediately before the start codon), and a promoter of CPCC (CMP166C), a promoter of Catalase (CMI050C), and the like.

[0213] The sequence information of these promoters can be obtained from a known sequence database such as GenBank. The promoter sequence of APCC of *Cyanidioschyzon merolae* is shown in SEQ ID NO: 9, the promoter sequence of CPCC (CMP166C) of *Cyanidioschyzon merolae* is shown in SEQ ID NO: 10, and the promoter sequence of Catalase (CMI050C) of *Cyanidioschyzon merolae* is shown in SEQ ID NO: 11.

[0214] Among the algae belonging to Cyanidiophyceae, *Cyanidioschyzon merolae* is algae capable of self-cloning (Fujiwara et al., PLOS One. 2013 Sep. 5; 8 (9): e73608). “Self-cloning” refers to a gene recombination technique only using, as nucleic acids to be introduced into a cell, (1) a nucleic acid of an organism that taxonomically belongs to the same species as the organism from which the cell is derived, and (2) a nucleic acid of an organism belonging to a species that exchanges nucleic acids with a taxonomic species, to which the organism as an origin of the cell belongs, under the natural conditions. The transformants prepared by self-cloning are excluded from the organisms to be subjected to genetic modification listed in the Cartagena Protocol. Therefore, the transformants can also be cultured outdoors. Accordingly, the transformant of the algae belonging to Cyanidiophyceae that is generated by self-cloning is suitable as algae used in the nutritional supplement of the present embodiment. Particularly, the transformant of *Cyanidioschyzon merolae* that is generated by self-cloning is preferable.

[0215] The method for performing self-cloning in *Cyanidioschyzon merolae* is not particularly limited, and examples thereof include a method using a URA5.3 gene (CMK046C) as a selection marker. *Cyanidioschyzon merolae* includes *Cyanidioschyzon merolae* M4 (Minoda et al., Plant Cell Physiol. 2004 June; 45 (6): 667-71.) which is a uracil auxotrophic mutant. *Cyanidioschyzon merolae* M4 has a mutation in the URA5.3 gene and cannot synthesize uracil. Accordingly, *Cyanidioschyzon merolae* M4 cannot grow in a uracil-free medium. Therefore, self-cloning can be performed using *Cyanidioschyzon merolae* M4 as a parent strain and using the URA5.3 gene of a wild-type strain as a selection marker. More specifically, an arbitrary gene set of *Cyanidioschyzon merolae* is linked to a URA5.3 gene set of a wild-type strain of *Cyanidioschyzon merolae* (for example, a 10D strain), and the resulting gene is introduced into *Cyanidioschyzon merolae* M4. Thereafter, by culturing the cells in a uracil-free medium, it is possible to obtain cells into which an arbitrary gene set has been introduced. “Gene set” described above means a set in which an arbitrary promoter, ORF of a target gene, and arbitrary 3'UTR are linked to each other. The 3'UTR is not particularly limited,

and may be 3'UTR of a target gene or 3'UTR of another gene. As the 3'UTR, for example, 3'UTR of β-tubulin is frequently used. The selection marker is not limited to the URA5.3 gene, and may be a gene related to other types of auxotrophy.

[0216] In a case where the genetic modification is performed using the auxotrophy-related gene as a selection marker as described above, by knocking out the auxotrophy-related gene introduced into the algal cells, genetic modification can be performed again by using the same auxotrophy-related gene as a selection marker. That is, it is possible to perform multiple self-cloning. As the method for knocking out the auxotrophy-related gene introduced as a selection marker, known knockout techniques may be used without particular limitation. Examples of the knockout techniques include homologous recombination, a gene editing technique, and the like.

[0217] For example, as described above, in *Cyanidioschyzon merolae*, self-cloning can be performed using the URA5.3 gene (CMK046C) as a selection marker and *Cyanidioschyzon merolae* M4 as a parent strain. Untransformed cells cannot grow in a uracil-free medium. Therefore, after the transformation, by culturing the cells in a uracil-free medium, it is possible to select transformants. Furthermore, in a case where self-cloning is performed, by a known knockout technique such as homologous recombination, the URA5.3 gene is knocked out. For example, the introduced URA5.3 gene may be totally or partially deleted, or a point mutation may be introduced into the URA5.3 gene. By being cultured in a medium containing uracil and 5-fluoroorotic acid (5-FOA), the URA5.3 gene knockout strain can be selected. This is because in a strain that normally expresses the URA5.3 gene, 5-FOA is converted into toxic 5-fluorouracil by the gene product of the URA5.3 gene. In a case where the URA5.3 knockout strain obtained as above is used as a parent strain, self-cloning can be performed again by using the URA5.3 gene as a selection marker. By repeating the same operation, it is possible to perform self-cloning a desired number of times.

[0218] As the method for introducing an arbitrary nucleic acid into the algae belonging to Cyanidiophyceae, known methods can be used without particular limitation. Examples thereof include a polyethylene glycol method, a lipofection method, a microinjection method, a DEAE dextran method, a gene gun method, an electroporation method, a calcium phosphate method, and the like.

[0219] In a case where the algae belonging to Cyanidiophyceae are subjected to transformation, the nucleic acid to be introduced may be inserted into any of a nuclear genome, a chloroplast genome, and a mitochondrial genome. In a case where the nucleic acid to be introduced is inserted into the genome, the nucleic acid may be inserted into a specific position in the genome or randomly inserted into the genome.

[0220] As the method for inserting the nucleic acid to be introduced into a specific position in the genome, homologous recombination can be used. For example, for *Cyanidioschyzon merolae*, because the entire genome sequence thereof has been completely mapped (Matsuzaki M et al., Nature. 2004 Apr. 8; 428 (6983): 653-7.), the nucleic acid to be introduced can be inserted into a desired position on the genome. There is no particular limitation on the transgene

insertion position in *Cyanidioschyzon merolae*. Examples of the insertion position include, for example, a region between CMD184C and CMD185C.

[0221] Specifically, as mutants of the algae belonging to Cyanidiophyceae, for example, the following (1) to (18) are preferable, but are not limited thereto.

[0222] (1) A transformant of the algae belonging to Cyanidiophyceae, in which the intracellular content of at least one kind of nutrient selected from the group consisting of γ -aminobutyric acid and vitamin E is higher than that in a wild-type strain.

[0223] (2) The transformant of the algae belonging to Cyanidiophyceae according to (1), in which an expression amount of a glutamate decarboxylase gene is larger than in a wild-type strain.

[0224] (3) The transformant of the algae belonging to Cyanidiophyceae according to (2), into which a nucleic acid containing a glutamate decarboxylase gene has been introduced in an expressible form.

[0225] (4) The transformant of the algae belonging to Cyanidiophyceae according to (3), in which the nucleic acid contains a promoter that functions in the cells of the algae belonging to Cyanidiophyceae and a glutamate decarboxylase gene that is operably linked to the promoter.

[0226] (5) The transformant of the algae belonging to Cyanidiophyceae according to (1), in which an expression amount of at least one kind of gene selected from the group consisting of a homogentisate phytoltransferase gene and a tocopherol cyclase gene is larger than in a wild-type strain.

[0227] (6) The transformant of the algae belonging to Cyanidiophyceae according to (5), into which a nucleic acid including at least one kind of gene selected from the group consisting of a homogentisate phytoltransferase gene and a tocopherol cyclase gene has been introduced in an expressible form.

[0228] (7) The transformant of the algae belonging to Cyanidiophyceae according to (6), in which the nucleic acid includes a promoter that functions in the cells of the algae belonging to Cyanidiophyceae and a homogentisate phytoltransferase gene operably linked to the promoter.

[0229] (8) The transformant of the algae belonging to Cyanidiophyceae according to (6) or (7), in which the nucleic acid includes a promoter that functions in the cells of the algae belonging to Cyanidiophyceae and a tocopherol cyclase gene that is operably linked to the promoter.

[0230] (9) The transformant of the algae belonging to Cyanidiophyceae according to any one of (4), (7), and (8), in which the promoter is selected from the group consisting of an APCC promoter, a CPCC promoter, and a Catalase promoter.

[0231] (10) The transformant of the algae belonging to Cyanidiophyceae according to any one of (3), (4), and (6) to (9), in which the nucleic acid does not include a base sequence derived from a cell belonging to another taxonomic species.

[0232] (11) The transformant of the algae belonging to Cyanidiophyceae according to (10), in which the nucleic acid has been introduced into the cells by using an auxotrophy-related gene as a selection marker.

[0233] (12) The transformant of the algae belonging to Cyanidiophyceae according to (10), in which the nucleic acid has been introduced into a knockout cell of the auxotrophy-related gene by using the auxotrophy-related gene as a selection marker.

[0234] (13) The transformant of the algae belonging to Cyanidiophyceae according to any one of (1) to (12), in which the algae belonging to Cyanidiophyceae are algae belonging to *Cyanidioschyzon*.

[0235] (14) The transformant of the algae belonging to Cyanidiophyceae according to (13), in which the algae belonging to Cyanidiophyceae are *Cyanidioschyzon merolae*.

[0236] (15) The transformant of the algae belonging to Cyanidiophyceae according to (14), in which the auxotrophic gene is a URA5.3 gene.

[0237] (16) The transformant of the algae belonging to Cyanidiophyceae according to any one of (1) to (12), in which the algae belonging to Cyanidiophyceae are haploid cells of algae belonging to *Galdieria*.

[0238] (17) The transformant of the algae belonging to Cyanidiophyceae according to (16), in which the algae belonging to *Galdieria* are *Galdieria sulphuraria* or *Galdieria partita*.

[0239] (18) The transformant of the algae belonging to Cyanidiophyceae according to any one of (1) to (12), in which the algae belonging to Cyanidiophyceae are haploid cells of algae belonging to *Cyanidium*.

[0240] In the present specification, "operably linked" means that a first base sequence is disposed sufficiently close to a second base sequence and can exert an influence on the second base sequence or on a region which is under the control of the second base sequence. For example, "gene operably linked to a promoter" means that the gene is linked so as to be expressed under the control of the promoter. "Expressible form" refers to a form where a gene can be transcribed and translated in a cell into which the gene has been introduced.

[Extract of Algae Belonging to Cyanidiophyceae]

[0241] An extract of algae according to an embodiment of the present invention refers to a substance extracted from one kind of algae belonging to Cyanidiophyceae. However, the extract may contain a substance extracted from two or more kinds of algae belonging to Cyanidiophyceae or undisturbed cells of Cyanidiophyceae, and an extract and cells of algae that do not belong to Cyanidiophyceae.

[0242] In the nutrient composition and the like according to the embodiment of the present invention, the extract of the algae belonging to Cyanidiophyceae may be used instead of or together with the algae belonging to Cyanidiophyceae. The extract of the algae belonging to Cyanidiophyceae can also be rich in nutrients such as amino acids, vitamins, proteins, lipids, and dietary fiber, just as the algae belonging to Cyanidiophyceae. Therefore, the present invention also provides a nutrient composition, a nutritional supplement, a food, a feed or a pet food, a cosmetic, and the like using the extract of the algae belonging to Cyanidiophyceae.

[0243] In the present specification, "extract of the algae belonging to Cyanidiophyceae" refers to a substance obtained by extracting intracellular components by performing a physical treatment or a chemical treatment on the cells of the algae belonging to Cyanidiophyceae. For example, the extract of the algae belonging to Cyanidiophyceae may be a

cell disruption product obtained by disrupting the cells of the algae belonging to Cyanidiophyceae by a physical treatment or a chemical treatment. In addition, the extract of algae belonging to Cyanidiophyceae may be a substance obtained by concentrating the cell disruption product, a substance obtained by removing solid contents from the cell disruption product, or a substance obtained by removing some components from the cell disruption product.

[0244] The method of the physical treatment or the chemical treatment for the cells is not particularly limited, and the methods generally used for cell disruption can be used. Examples of the physical treatment include cell disruption performed by means of glass beads, a mortar, an ultrasonic treatment, a French press, a homogenizer, and the like. Examples of the chemical treatment include a neutralization treatment, a hypotonic treatment, a freeze-thaw treatment, a dry and swelling treatment, and the like. More specifically, examples thereof include the cell rupturing treatments (A) to (C) described above.

[0245] In a case where the cell disruption product is concentrated, the concentration method is not particularly limited, and a generally used concentration method may be used. Examples of the concentration method for the cell disruption product include drying, freeze-drying, drying under reduced pressure, and the like.

[0246] In a case where solid contents are removed from the cell disruption product, the solid content-removing method is not particularly limited, and it is possible to use a method that is generally used for removing solid contents and the like. Examples of the solid content-removing method include filtration, centrifugation, and the like.

[0247] In a case where some components are separated from the cell disruption product, the separation method is not particularly limited, and it is possible to use a method generally used for separation, purification, and the like of biochemical substances. Examples of the separation method include salting out, dialysis, solvent extraction, adsorption, column chromatography, ion exchange chromatography, and the like. Each of these methods may be used singly. Alternatively, two or more kinds of treatments may be used in combination. Here, a substance purified into a single component is excluded from "extract of the algae belonging to Cyanidiophyceae". The extract of the algae belonging to Cyanidiophyceae preferably contains 10 or more kinds of cell components of the algae belonging to Cyanidiophyceae, more preferably contains 15 or more kinds of the cell components described above, and even more preferably contains 20 or more kinds of the cell components described above. The extract of the algae belonging to Cyanidiophyceae preferably contains a nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber, and more preferably contains a nutrient selected from the group consisting of amino acids and vitamins. Specific examples of the amino acids include isoleucine, leucine, lysine, methionine, cystine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, γ -aminobutyric acid, and the like. Specific examples of the vitamins include vitamin A, β -carotene, vitamin B₁, vitamin B₂, vitamin B₆, vitamin C, vitamin E, vitamin K₁, vitamin K₂, niacin, inositol, folic acid, biotin, and the like.

[Alga Having a Diploid Cell Form and a Haploid Cell Form]

[0248] In one embodiment, the present invention provides an alga which belongs to Cyanidiophyceae and having a diploid cell form and a haploid cell form.

[0249] One of the characteristics of the alga of the present embodiment is that the alga belong to Cyanidiophyceae and having diploid cell form and a haploid cell form. Hitherto, it has been known that among the algae belonging to Cyanidiophyceae, the algae belonging to *Cyanidioschyzon* include haploid algae. However, among the algae belonging to Cyanidiophyceae, no algae have been known so far which have both the diploid cell form and haploid cell form. That is, from nature, *Cyanidioschyzon merolae* belonging to *Cyanidioschyzon* which belongs to Cyanidiophyceae has been collected only in the form of haploids. Furthermore, from nature, *Galdieria* and *Cyanidium*, which are other algae belonging to Cyanidiophyceae, have been collected only in the form of diploids. The present invention provides algae, which are haploid cells that are created from diploid cells of the algae belonging to Cyanidiophyceae, and algae which are diploid cells generated from two or more haploid cells. In the present specification, the algae described above will be expressed using "having both a diploid cell form and a haploid cell form".

[0250] The algae of the present embodiment having both a diploid cell form and a haploid cell form.

[0251] In the algae of the present embodiment, the haploid cells are generated by the meiosis of the diploid cells. It is considered that by the mating of two haploid cells, diploid cells may be generated. Therefore, in the algae of the present embodiment, by mating haploid cells having desired traits, diploid cells having the traits of the cells can be prepared.

[0252] For example, in a case where the haploid cells are used, it is easier to prepare a transformant by using a gene recombination technique than in a case where the diploid cells are used. Therefore, a method is considered in which a plurality of transformants having any traits are prepared using the haploid cells, and the transformants having any traits are crossing so as to prepare diploids having all the traits of the transformants.

[0253] The algae of the present embodiment can grow in a cell population consisting only of haploid cells.

[0254] The algae of the present embodiment can grow in a cell population consisting only of diploid cells.

[0255] Furthermore, as is evident from Examples which will be described later, it is possible to create haploid cells from diploid cells and to create diploid cells from two or more kinds of haploid cells. In addition, in a case where haploid cells are created from diploid cells or in a case where diploid cells are created from haploid cells, sometimes the haploid cells and the diploid cells are mixed together.

[0256] Whether certain algae are diploids or haploids can be determined by the method exemplified in "[Alga belonging to Cyanidiophyceae]" described above.

[0257] Furthermore, whether certain algae having both the diploid cell form and a haploid cell form can be determined, for example, by the following method. First, the diploid cells are cultured until they reach a stationary phase, and whether cells in the form different from the diploid appear is checked when culturing is continued in the stationary phase. When the cells in the form different from the diploid appear, the cells are collected, and whether they are haploid cells is checked. In a case where the cells are found to be haploid

cells as a result, it is possible to determine that the algae having both a diploid cell form and a haploid cell form.

[0258] In the algae of the present embodiment, it is preferable that at least the cells in either of the cell forms do not have a rigid cell wall. The cells having no rigid cell wall can be disrupted by a relatively mild treatment such as a neutralization treatment, a hypotonic treatment, a freeze-thaw treatment, or the like. Whether or not the algae have a rigid cell wall can be determined by the method described in “[Alga belonging to Cyanidiophyceae]” described above.

[0259] In the algae of the present embodiment, it is preferable that at least either the diploid cells or the haploid cells do not have a rigid cell wall. It is more preferable that the haploid cells do not have a rigid cell wall. That is, it is preferable that the haploid cells are ruptured by any of the cell rupturing treatments (A) to (C) described above.

[0260] In a case where a cell does not have a rigid cell wall, generally, the cell wall is not observed by the observation with an optical microscope (for example, at a magnification of 600 \times). In the algae of the present embodiment, usually, the diploid cells have a rigid cell wall while the haploid cells do not have a rigid cell wall.

[0261] As shown in Examples which will be described later, the algae of the present embodiment are rich in nutrients such as amino acids, vitamins, proteins, lipids, and dietary fiber. Particularly, it has been confirmed that the algae belonging to Cyanidiophyceae contain amino acids and vitamins at a higher concentration compared to the conventionally used algae (*Chlorella*, *Euglena*, and *Spirulina*). Therefore, the algae of the present embodiment can be used for a method for producing a nutritional supplement or a nutrient which will be described later.

[0262] Examples of the amino acids contained in the algae of the present embodiment include isoleucine, leucine, lysine, methionine, cystine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, γ -aminobutyric acid, and the like. One of the characteristics of the algae of the present embodiment is that the total amino acid content thereof is higher than that of the conventionally used algae. Furthermore, the content of each of the amino acids in the algae of the present embodiment tends to be high in general.

[0263] For example, the total amino acid content in the algae of the present embodiment is preferably equal to or greater than 50 g/100 g based on dry weight. The range of the total amino acid content is, for example, 50 g to 70 g/100 g based on dry weight. In the present specification, “total amino acid content” means a value obtained by adding up the contents of isoleucine, leucine, lysine, methionine, cystine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, and serine.

[0264] The content of isoleucine in the algae of the present embodiment is preferably equal to or greater than 1.5 g/100 g based on dry weight. The range of the content of isoleucine is, for example, 1.5 g to 5 g/100 g based on dry weight.

[0265] The content of leucine in the algae of the present embodiment is preferably equal to or greater than 4.5 g/100 g based on dry weight. The range of the content of isoleucine is, for example, 4.5 g to 8 g/100 g based on dry weight.

[0266] The content of lysine in the algae of the present embodiment is preferably equal to or greater than 2.5 g/100

g based on dry weight. The range of the content of lysine is, for example, 2.5 g to 6 g/100 g based on dry weight.

[0267] The content of methionine in the algae of the present embodiment is preferably equal to or greater than 0.5 g/100 g based on dry weight. The range of the content of methionine is, for example, 0.5 g to 4 g/100 g based on dry weight.

[0268] The content of cystine in the algae of the present embodiment is preferably equal to or greater than 0.5 g/100 g based on dry weight. The range of the content of cystine is, for example, 0.5 g to 3 g/100 g based on dry weight.

[0269] The content of phenylalanine in the algae of the present embodiment is preferably equal to or greater than 1.5 g/100 g based on dry weight. The range of the content of phenylalanine is, for example, 1.5 g to 5 g/100 g based on dry weight.

[0270] The content of tyrosine in the algae of the present embodiment is preferably equal to or greater than 2.0 g/100 g based on dry weight. The range of the content of tyrosine is, for example, 2.0 g to 5 g/100 g based on dry weight.

[0271] The content of threonine in the algae of the present embodiment is preferably equal to or greater than 2.0 g/100 g based on dry weight. The range of the content of threonine is, for example, 2.0 g to 5 g/100 g based on dry weight.

[0272] The content of tryptophan in the algae of the present embodiment is preferably equal to or greater than 0.5 g/100 g based on dry weight. The range of the content of tryptophan is, for example, 0.5 g to 3 g/100 g based on dry weight.

[0273] The content of valine in the algae of the present embodiment is preferably equal to or greater than 2.5 g/100 g based on dry weight. The range of the content of valine is, for example, 2.5 g to 6 g/100 g based on dry weight.

[0274] The content of arginine in the algae of the present embodiment is preferably equal to or greater than 4.5 g/100 g based on dry weight. The range of the content of phenylalanine is, for example, 4.5 g to 8 g/100 g based on dry weight.

[0275] The content of histidine in the algae of the present embodiment is preferably equal to or greater than 0.5 g/100 g based on dry weight. The range of the content of histidine is, for example, 0.5 g to 3 g/100 g based on dry weight.

[0276] The content of alanine in the algae of the present embodiment is preferably equal to or greater than 3.5 g/100 g based on dry weight. The range of the content of alanine is, for example, 3.5 g to 7 g/100 g based on dry weight.

[0277] The content of aspartic acid in the algae of the present embodiment is preferably equal to or greater than 4.5 g/100 g based on dry weight. The range of the content of aspartic acid is, for example, 4.5 g to 8 g/100 g based on dry weight.

[0278] The content of glutamic acid in the algae of present embodiment is preferably equal to or greater than 5.5 g/100 g based on dry weight. The range of the content of glutamic acid is, for example, 5.5 g to 9 g/100 g based on dry weight.

[0279] The content of glycine in the algae of the present embodiment is preferably equal to or greater than 2.0 g/100 g based on dry weight. The range of the content of glycine is, for example, 2.0 g to 5.5 g/100 g based on dry weight.

[0280] The content of proline in the algae of the present embodiment is preferably equal to or greater than 1.5 g/100 g based on dry weight. The range of the content of proline is, for example, 1.5 g to 5 g/100 g based on dry weight.

[0281] The content of serine in the algae of present embodiment is preferably equal to or greater than 2.0 g/100 g based on dry weight. The range of the content of serine is, for example, 2.0 g to 5 g/100 g based on dry weight.

[0282] The content of γ -aminobutyric acid in the algae of the present embodiment is preferably equal to or greater than 0.05 g/100 g based on dry weight. The range of the content of γ -aminobutyric acid is, for example, 0.05 g to 0.3 g/100 g based on dry weight.

[0283] Examples of the vitamins contained in the algae of the present embodiment include vitamin A, β -carotene, vitamin C, vitamin E, vitamin K₁, vitamin K₂, folic acid, and the like.

[0284] One of the characteristics of the algae of the present embodiment is that the content of β -carotene, vitamin C, vitamin E, vitamin K₁, vitamin K₂, and folic acid is particularly higher in the algae of the present embodiment than in the conventionally used algae.

[0285] The content of vitamin A in the algae of the present embodiment is preferably equal to or greater than 8 mg/100 g based on dry weight. The range of the content of vitamin A is, for example, 8 mg to 20 mg/100 g based on dry weight.

[0286] The content of β -carotene in the algae of the present embodiment is preferably equal to or greater than 100 mg/100 g based on dry weight. The range of the content of β -carotene is, for example, 100 mg to 200 mg/100 g based on dry weight.

[0287] The content of vitamin C in the algae of the present embodiment is preferably equal to or greater than 20 mg/100 g based on dry weight. The range of the content of vitamin C is, for example, 20 mg to 50 mg/100 g based on dry weight.

[0288] The content of vitamin E in the algae of the present embodiment is preferably equal to or greater than 80 mg/100 g based on dry weight. The range of the content of vitamin E is, for example, 80 mg to 150 mg/100 g dry weight.

[0289] The content of vitamin K₁ in the algae of the present embodiment is preferably equal to or greater than 4,000 μ g/100 g based on dry weight. The range of the content of vitamin K₁ is, for example, 4,000 μ g to 8,000 μ g/100 g based on dry weight.

[0290] The content of vitamin K₂ in the algae of the present embodiment is preferably equal to or greater than 1,000 μ g/100 g based on dry weight. The range of the content of vitamin K₂ is, for example, 1,000 μ g to 3,000 μ g/100 g based on dry weight.

[0291] The content of folic acid in the algae of the present embodiment is preferably equal to or greater than 1,500 μ g/100 g based on dry weight. The range of the content of folic acid is, for example, 1,500 μ g to 4,000 μ g/100 g based on dry weight.

[0292] The algae of the present embodiment can be cultured using a medium for culturing microalgae. The culturing can be performed in the same manner as in the method described above in “[Alga belonging to Cyanidiophyceae]”.

[0293] The algae of the present embodiment can also be cultured in a medium using acidic hot spring drainage. “Acidic hot spring drainage” means acidic drainage discharged from hot spring facilities. The acidic hot spring drainage is not particularly limited, and the pH thereof is preferably 1.0 to 4.0 and more preferably 1.0 to 3.0. “Medium using acidic hot spring drainage” means a medium prepared by adding a nitrogen source, a phosphorus source, trace elements, and the like to acidic hot spring drainage. As

the medium using acidic hot spring drainage, a medium obtained by adding a nitrogen source to acidic hot spring drainage is preferable, and a medium obtained by adding a nitrogen source and a phosphorus source to acidic hot spring drainage is more preferable (for example, see Hirooka S and Miyagishima S. Y. (2016) Cultivation of Acidophilic Algae *Galdieria sulphuraria* and *Pseudochlorella* sp. YKT1 in Media Derived from Acidic Hot Springs. Front Microbiol. December 20; 7:2022.). Examples of the nitrogen source include ammonium salts (such as ammonium sulfate), urea, nitrates (such as sodium nitrate), and the like. Among these, ammonium salts and urea are preferable, and ammonium salts are more preferable. The amount of the nitrogen source to be added is, for example, 1 to 50 mM as the amount of nitrogen to be added. The amount of the nitrogen source to be added is preferably 5 to 40 mM and more preferably 10 to 30 mM as the amount of nitrogen to be added. Examples of the phosphorus source include phosphates (such as potassium dihydrogen phosphate). The amount of the phosphorus source to be added is, for example, 0.1 to 10 mM as the amount of phosphorus to be added. The amount of the phosphorus source to be added is preferably 0.5 to 5 mM and more preferably 1 to 3 mM as the amount of phosphorus to be added. The algae of the present embodiment can be cultured in a medium using acidic hot spring drainage. Therefore, the algae of the present embodiment make it possible to effectively use acidic hot spring drainage and can be cultured at low costs.

[0294] In a case where the algae of the present embodiment are algae belonging to *Galdieria*, as the aforementioned nitrogen source, an ammonium salt or urea is preferable, and an ammonium salt is more preferable. In a case where the algae of the present embodiment are algae belonging to *Cyanidium*, as the aforementioned nitrogen source, an ammonium salt or a nitrate is preferable, and an ammonium salt is more preferable.

(YFU3 Strain and HKN1 Strain)

[0295] Specific examples of the algae of the present embodiment include a *Cyanidium* sp. YFU3 (FERM BP-22334) (hereinafter, referred to as “YFU3 strain”), a *Cyanidium* sp. HKN1 (FERM BP-22333) (hereinafter referred to as “HKN1 strain”), and allied species, mutants, and progenies of these.

[0296] Both the YFU3 strain and the HKN1 strain have a diploid cell form and a haploid cell form. Hereinafter, when the YFU3 strain is described by distinguishing between diploid cells and haploid cells, the diploid cells will be described as “YFU3 strain (diploid)”, and the haploid cells will be described as “YFU3 strain (haploid)”. Likewise, when the HKN1 strain is described by distinguishing between diploid cells and haploid cells, the diploid cells will be described as “HKN1 strain (diploid)”, and the haploid cells will be described as “HKN1 strain (haploid)”. When the strains are simply referred to as “YFU3 strain” or “HKN1 strain”, each of the strains includes both the diploid cells and the haploid cells.

[0297] The YFU3 strain (haploid) is unicellular red algae isolated from hot acidic water of a hot spring in Yufu-shi, Oita prefecture, Japan. The YFU3 strain was deposited in the Patent Microorganism Depository Center of National Institute of Technology and Evaluation (2-5-8 Kazusa Kamatari, Kisarazu-shi, Chiba prefecture, Japan) on Jun. 28, 2017 under the accession No. FERM P-22334, and transferred to

the International Depository on May 23, 2018 under the accession No. FERM BP-22334.

[0298] The HKN1 strain is unicellular red algae isolated from hot acidic water of a hot spring in Hakone-machi, Ashigarashimo-gun, Kanagawa prefecture, Japan. The HKN1 strain (haploid) was deposited in the Patent Micro-organism Depository Center of National Institute of Technology and Evaluation on Jun. 28, 2017 under the accession No. FERM P-22333, and transferred to the International Depository on May 23, 2018 under the accession No. FERM BP-22333.

[0299] Both the YFU3 strain and the HKN1 strain have a blue pigment, phycocyanin, in addition to chlorophyll a. Therefore, these strains appear blue-green. Both the YFU3 strain and the HKN1 strain grow suitably in a high-temperature acidic environment, at an optimum temperature of about 42° C. and an optimum pH of around 2.

[0300] Both the YFU3 strain (diploid) and the HKN1 strain (diploid) have a cell size of about 4 μm . In addition, both the YFU3 strain (haploid) and the HKN1 strain (haploid) have a cell size of about 2 μm . Furthermore, both the YFU3 strain (diploid) and the HKN1 strain (diploid) have a rigid cell wall. That is, these strains do not undergo cell rupture under the condition of pH 7. In contrast, neither the YFU3 strain (haploid) nor the HKN1 strain (haploid) has a rigid cell wall. That is, these strains undergo cell rupture under the condition of pH 7.

[0301] The cells of the YFU3 strain (haploid) and the HKN1 strain (haploid) are similar to the cells of *Cyanidioschyzon merolae* (see FIG. 8). Among the algae belonging to Cyanidiophyceae, *Cyanidioschyzon merolae* is the only known species of haploid algae having no rigid cell wall. In the present specification, the cells similar to the cells of *Cyanidioschyzon merolae* will be described as “*Cyanidioschyzon merolae*-like cells” in some cases.

[0302] When the YFU3 strain (diploid) is cultured until the strain enters the stationary phase, and the culturing is continued in the stationary phase, several YFU3 strains (diploid) undergo meiosis. As a result, from one cell of the YFU3 strain (diploid), four cells of the YFU3 strain (haploid) are generated. Likewise, when the HKN1 strain (diploid) is cultured until the strain enters the stationary phase, and the culturing is continued in the stationary phase, several HKN1 strains (diploid) undergo meiosis. As a result, from one cell of the HKN1 strain (diploid), four cells of HKN1 strain (haploid) are generated. Through two-cell division, both the YFU3 strain (haploid) and the HKN1 strain (haploid) can grow while maintaining the form of haploid cells.

[0303] As the algae of the present embodiment, for example, in addition to the YFU3 strain and the HKN1 strain, algae, which are allied species of the YFU3 strain or the HKN1 strain and having a diploid cell form and a haploid cell form, are suitable. Examples of the algae as allied species of the YFU3 strain and the HKN1 strain include algae in which the base sequence of the rbcL gene shares identity equal to or higher than 90% with the base sequence of the rbcL gene of the YFU3 strain or the HKN1 strain. The base sequence of the rbcL gene of the YFU3 strain is shown in SEQ ID NO: 1. The base sequence of the rbcL gene of the HKN1 strain is shown in SEQ ID NO: 2. Therefore, the algae, in which the base sequence of the rbcL gene shares identity equal to or higher than 90% with the base sequence shown in SEQ ID NO: 1 or 2, are also a suitable example of the algae of the present embodiment.

The identity between the base sequence of the rbcL gene of the algae and the base sequence shown in SEQ ID NO: 1 or 2 is preferably equal to or higher than 95%, more preferably equal to or higher than 97%, even more preferably equal to or higher than 98%, and particularly preferably equal to or higher than 99%.

[0304] The base sequence of the rbcL gene of algae can be obtained by a known method. For example, by extracting DNA from algal cells of interest by a known method, amplifying DNA fragments of the rbcL gene by PCR or the like, and analyzing the base sequence of the amplified DNA fragments by using a DNA sequencer, the base sequence of the rbcL gene of the algae of interest can be obtained. Examples of primers for amplifying the rbcL gene include the primers used in Examples of the present specification and the like.

[0305] As the algae of the present embodiment, for example, algae, which are a mutant of the YFU3 strain or the HKN1 strain and have a diploid cell form and a haploid cell form, are also suitable. In the present specification, “mutant” means an algal strain generated when spontaneous or induced mutation occurs in the genome of the original algal strain (including a nuclear genome, a chloroplast genome, and a mitochondrial genome; the same is true of the following description). The technique for inducing mutation in the genome is not particularly limited, and examples thereof include ultraviolet irradiation, radiation irradiation, a chemical treatment using nitrous acid; genetic engineering techniques such as gene introduction and genome editing, and the like. In the present specification, “mutant of the YFU3 strain” refers to an algal strain which is generated by the mutation of the genome of the YFU3 strain and has a diploid cell form and a haploid cell form. Furthermore, “mutant of the HKN1 strain” refers to an algal strain which is generated by the mutation of the genome of the HKN1 strain and has a diploid cell form and a haploid cell form.

[0306] In the mutant of the YFU3 strain, the proportion of mutations in the whole genome of the YFU3 strain is preferably equal to or lower than 10%, more preferably equal to or lower than 5%, even more preferably equal to or lower than 3%, and particularly preferably equal to or lower than 2% or 1%.

[0307] In the mutant of the HKN1 strain, the proportion of mutations in the whole genome of the HKN1 strain is preferably equal to or lower than 10%, more preferably equal to or lower than 5%, even more preferably equal to or lower than 3%, and particularly preferably equal to or lower than 2% or 1%.

[0308] In the above description, the proportion of mutations in the whole genome is calculated by comparing the whole genomes of the diploid cells or comparing the whole genomes of the haploid cells.

[0309] Among the above, as the allied species or the mutant of the YFU3 strain or the HKN1 strain, those having a nutritional composition similar to the nutritional composition of the YFU3 strain or the HKN1 strain are preferable.

[0310] For example, one of the characteristics of the YFU3 strain and the HKN1 strain is that these strains are rich in amino acids and vitamins. Therefore, it is preferable that the content of amino acids or vitamins in the allied species or the mutant is similar to those in the YFU3 strain and the HKN1 strain. Examples of the content of amino acids and vitamins include the contents exemplified above.

[0311] Specific examples of the mutant of the YFU3 strain or the HKN1 strain include transformants obtained by genetic modification using genetic engineering techniques. The genetic modification may be performed on the diploid cells or the haploid cells. However, it is easier to genetically modify the haploid cells. The type of genetic modification is not particularly limited, and any type of modification may be adopted.

[0312] Examples of the transformant of the YFU3 strain or the HKN1 strain include a transformant having an increased intracellular content of at least one kind of nutrient. As the nutrient of which the intracellular content is to be increased, for example, at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber is suitable. That is, a transformant of the YFU3 strain or the HKN1 strain is suitable, in which the intracellular content of at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber is higher than that in a wild-type strain.

[0313] Among the above nutrients, at least one kind of nutrient selected from the group consisting of amino acids and vitamins is preferable. Such a transformant has a higher intracellular amount of a specific nutrient compared to a wild-type strain. Therefore, by using this transformant, it is possible to prepare a nutritional supplement rich in the nutrient.

[0314] The method for increasing the intracellular content of a nutrient is not particularly limited, and any method can be used. Examples thereof include a method similar to the method exemplified in “[Alga belonging to Cyanidiophyceae]” described above. For example, by modifying the YFU3 strain or the HKN1 strain such that the expression amount of any enzyme gene involved in the synthesis of nutrients such as glutamate decarboxylase, homogenitise phytoltransferase, and tocopherol cyclase is increased, the intracellular content of any nutrient can be increased.

[0315] The sequence information of these synthetic enzyme genes can be obtained by cloning the gene of interest and analyzing the sequence of the cloned gene of interest by a known method.

[0316] For example, primers may be designed based on the sequence information of known homologous genes of the algae belonging to Cyanidiophyceae, and amplified fragments of the gene of interest may be obtained by PCR or the like by using the genome of the YFU3 strain or the HKN1 strain as a template. Alternatively, a probe may be designed based on the sequence information of a known homologous gene of the algae belonging to Cyanidiophyceae, and a cDNA library of the YFU3 strain or the HKN1 strain may be screened out.

[0317] Furthermore, as a transgene, a gene of an organism of another species may be used as long as the gene functions in the transfected cell. For example, a known gene of the algae belonging to Cyanidiophyceae may be used. The sequence information of a known gene can be obtained from a known sequence database such as GenBank. For example, it is possible to use a glutamate decarboxylase gene (CMF072C), a homogenitise phytoltransferase gene (CMN202C), a tocopherol cyclase gene (CML326C), of *Cyanidioschyzon merolae*, and the like.

[0318] For example, by introducing a synthetic enzyme gene (for example, a glutamate decarboxylase gene) of the nutrient described above into the YFU3 strain or the HKN1

strain, it is possible to obtain a transformant having an increased intracellular content of the nutrient. The promoter used for the gene to be introduced may be the promoter of the synthetic enzyme gene or the promoter of another gene. In a case where the promoter of another gene is used, a promoter of a gene that is expressed in large amounts in the algal cell into which the gene is to be introduced is preferable. Examples of such a promoter include an APCC (CMO250C) promoter, a CPCC (CMP166C) promoter, a Catalase (CMI050C) promoter, and the like. The promoter may be derived from an organism of the same species as the cell to be transfected, or may be derived from an organism of another species, as long as the promoter functions in the cell to be transfected.

[0319] In a case where a promoter of a known gene is used, the sequence information thereof can be obtained from a known sequence database such as GenBank.

[0320] The transformant of the YFU3 strain or the HKN1 strain can be prepared by the introduction of any nucleic acid. However, the transformant is more preferably prepared by self-cloning. For example, among the algae belonging to Cyanidiophyceae, for *Cyanidioschyzon merolae*, a self-cloning method has been established (Fujiwara et al., PLOS One. 2013 Sep. 5; 8 (9): e73608). Therefore, self-cloning can also be performed on the YFU3 strain or the HKN1 strain by the same method as the method used for *Cyanidioschyzon merolae*.

[0321] The self-cloning method is not particularly limited, and examples thereof include a method using an auxotrophic mutant. For example, for *Cyanidioschyzon merolae*, a method using the URA5.3 gene (CMK046C) as a selection marker has been suggested (see “[Algae belonging to Cyanidiophyceae]” described above).

[0322] For the YFU3 strain and the HKN1 strain, by knocking out an auxotrophy-related gene by a known gene knockout technique or the like so as to prepare an auxotrophic mutant, self-cloning can be performed using the gene to be knocked out as a selection marker. As the knockout technique, a known method may be used without particular limitation. Examples of the knockout technique include homologous recombination, a gene editing technique, and the like. The gene to be knocked out is not particularly limited as long as the auxotrophy is changed by the knockout of the gene. Examples of the gene to be knocked out include orotidine-5'-decarboxylase and the like.

[0323] In addition, in a case where genetic modification is performed using the auxotrophy-related gene as a selection marker as described above, by knocking out the auxotrophy-related gene introduced into algal cells, the genetic modification can be performed again by using the same auxotrophy-related gene as a selection marker. That is, it is possible to perform multiple self-cloning. As the method for knocking out the auxotrophy-related gene introduced as a selection marker, known knockout techniques may be used without particular limitation. Examples of the knockout technique include homologous recombination, a gene editing technique, and the like.

[0324] For example, as described above, in *Cyanidioschyzon merolae*, self-cloning can be performed using the URA5.3 gene (CMK046C) as a selection marker and *Cyanidioschyzon merolae* M4 as a parent strain. In a case where a URA5.3 gene knockout strains are prepared in the YFU3 strain and the HKN1 strain, the knockout strains cannot grow in a uracil-free medium. In a case where the afore-

mentioned strains are used as parent strains, the URA5.3 gene is introduced into the strains so as to cause transformation, and the transformed cells are cultured in uracil-free medium, transformants can be selected. Furthermore, in a case where self-cloning is performed, the URA5.3 gene is knocked out again by a known knockout technique such as homologous recombination. By culturing the cells in a medium containing uracil and 5-fluoroorotic acid, the re-knockout strain of the URA5.3 gene can be selected. This is because in a strain that normally expresses the URA5.3 gene, 5-FOA is converted into toxic 5-fluorouracil by the gene product of the URA5.3 gene. In a case where the re-knockout strain of URA5.3 obtained as above is used as a parent strain, self-cloning can be performed again by using the URA5.3 gene as a selection marker. By repeating the same operation, it is possible to perform self-cloning a desired number of times. By the knockout of the URA5.3 gene, the URA5.3 gene may be totally or partially deleted, or a point mutation may be introduced into the URA5.3 gene.

[0325] The method for introducing an arbitrary nucleic acid into the YFU3 strain or the HKN1 strain is not particularly limited, and a known method can be used. Examples thereof include a polyethylene glycol method, a lipofection method, a microinjection method, a DEAE dextran method, a gene gun method, an electroporation method, a calcium phosphate method, and the like.

[0326] In a case where the YFU3 strain or the HKN1 strain is transformed, the nucleic acid to be introduced may be inserted into any of a nuclear genome, a chloroplast genome, and a mitochondrial genome. In a case where the nucleic acid to be introduced is inserted into the genome, the nucleic acid may be inserted into a specific position in the genome or randomly inserted into the genome. As the method for inserting the nucleic acid to be introduced into a specific position in the genome, homologous recombination can be used.

[0327] An example of the transformation method for the YFU3 strain or the HKN1 strain will be described below, but the present invention is not limited thereto.

[0328] First, the YFU3 strain or the HKN1 strain is diluted with an appropriate medium such as MA2U medium (see Example 7) such that the concentration thereof becomes appropriate. Then, the cells are cultured for about 40 to 80 hours with aeration under appropriate culture conditions (for example, light-dark cycle: 12L:12D, light: 50 $\mu\text{mol}/\text{m}^2\text{s}$, temperature: 42° C.). Thereafter, Tween-20 is added to the culture solution such that the final concentration becomes 0.002%, and then the cells are collected by centrifugation and suspended in an appropriate medium such as MA2U medium. The culture solution is dissolved in 450 μL of MA2U medium containing PEG4000 (95° C., 10 minutes), thereby preparing a 60% (w/v) PEG4000 solution. Subsequently, the PEG 4000 solution was kept at 42° C. on a heat block until used.

[0329] The cell suspension of the YFU3 strain or the HKN1 strain is added to a mixture for transformation containing a vector for transformation including an appropriate selection marker and polyethylene glycol (such as PEG4000), and the mixture is stirred. The cells are moved to an appropriate medium such as MA2U medium and cultured in a static state for about 40 to 60 hours under appropriate culture conditions (for example, continuous light: 20 $\mu\text{mol}/\text{m}^2\text{s}$, temperature: 42° C.). Then, the cells are

collected by centrifugation, and suspended in an appropriate medium such as Tsukahara mineral spring medium (Hirooka et al. 2016 Front in Microbiology) or modified MA medium (see Example 7). The cell suspension is added to a selective medium compatible with the used selection marker (for example, a medium obtained by adding a specific substance to the Tsukahara mineral spring medium, the modified MA medium, or the like, or a medium obtained by removing a specific substance from the medium described above), and the cells are cultured in a static state for about 5 to 10 days under appropriate culture conditions (for example, continuous light: 20 $\mu\text{mol}/\text{m}^2\text{s}$, temperature: 42° C., 3% CO₂). The culture solution of a green portion, which has become darker, is added to the newly collected selective medium, the cells are further cultured in a static state for about 5 to 10 days, and transformants are selected. For example, under an inverted microscope, the transformants are isolated one by one by using a Pasteur pipette with a sharp tip, and cultured in a static state in an appropriate medium such as the Tsukahara mineral spring medium or the modified MA medium. In this way, the transformants can be obtained.

[0330] In the above section, an example of transformation by a polyethylene glycol method was described. However, other transformation methods such as a lipofection method, a microinjection method, a DEAE dextran method, a gene gun method, an electroporation method, and a calcium phosphate method may also be used.

[0331] Specific examples of strains preferred as the mutants of the YFU3 strain or the HKN1 strain include, but are not limited to, the following (1) to (18).

[0332] (1) A transformant of the YFU3 strain or the HKN1 strain, which has a higher intracellular content of at least one kind of nutrient selected from the group consisting of γ -aminobutyric acid and vitamin E as compared to a wild-type strain.

[0333] (2) The transformant of the YFU3 strain or the HKN1 strain according to (1), in which an expression amount of a glutamate decarboxylase gene is larger than in a wild-type strain.

[0334] (3) The transformant of the YFU3 strain or the HKN1 strain according to (2), into which a nucleic acid containing the glutamate decarboxylase gene has been introduced in an expressible form.

[0335] (4) The transformant of the YFU3 strain or the HKN1 strain according to (3), in which the nucleic acid contains a promoter that functions in cells of algae belonging to Cyanidiophyceae and a glutamate decarboxylase gene operably linked to the promoter.

[0336] (5) The transformant of the YFU3 strain or the HKN1 strain according to (1), in which an expression amount of at least one kind of gene selected from the group consisting of a homogentisate phytoltransferase gene and a tocopherol cyclase gene is larger than in the wild-type strain.

[0337] (6) The transformant of the YFU3 strain or the HKN1 strain according to (5), into which a nucleic acid containing at least one kind of gene selected from the group consisting of a homogentisate phytoltransferase gene and a tocopherol cyclase gene has been introduced in an expressible form.

[0338] (7) The transformant of the YFU3 strain or the HKN1 strain according to (6), in which the nucleic acid contains a promoter that functions in cells of algae

belonging to Cyanidiophyceae and a homogentisate phytoltransferase gene operably linked to the promoter.

[0339] (8) The transformant of the YFU3 strain or the HKN1 strain according to (6) or (7), in which the nucleic acid contains a promoter that functions in cells of algae belonging to Cyanidiophyceae and a tocopherol cyclase gene operably linked to the promoter.

[0340] (9) The transformant of the YFU3 strain or the HKN1 strain according to any one of (4), (7), and (8), in which the promoter is selected from the group consisting of an APCC promoter, a CPCC promoter, and a Catalase promoter.

[0341] (10) The transformant of the YFU3 strain or the HKN1 strain according to any one of (3), (4), and (6) to (9), in which the nucleic acid does not include a base sequence derived from a cell belonging to another taxonomic species.

[0342] (11) The transformant of the algae belonging to Cyanidiophyceae according to (10), in which the nucleic acid has been introduced into the cells by using an auxotrophy-related gene as a selection marker.

[0343] (12) The transformant of the YFU3 strain or the HKN1 strain according to (10), in which the nucleic acid has been introduced into a knockout cell of the auxotrophy-related gene by using the auxotrophy-related gene as a selection marker.

[0344] (13) The transformant of the YFU3 strain or the HKN1 strain according to any one of (3), (4), and (9), in which the glutamate decarboxylase gene is a glutamate decarboxylase gene of *Cyanidioschyzon merolae*.

[0345] (14) The transformant of the YFU3 strain or the HKN1 strain according to any one of (6), (7), and (9), in which the homogentisate phytoltransferase gene is a homogentisate phytoltransferase gene of *Cyanidioschyzon merolae*.

[0346] (15) The transformant of the YFU3 strain or the HKN1 strain according to any one of (6), (8), and (9), in which the tocopherol cyclase gene is a tocopherol cyclase gene of *Cyanidioschyzon merolae*.

[0347] (16) The transformant of the YFU3 strain or the HKN1 strain according to any one of (4) and (7) to (9), in which the promoter is a promoter derived from *Cyanidioschyzon merolae*.

[0348] (17) The transformant of the YFU3 strain or the HKN1 strain according to any one of (1) to (16), which is a haploid cell.

[0349] (18) The transformant of the YFU3 strain or the HKN1 strain according to any one of (1) to (16), which is a diploid cell.

[0350] In addition, as shown in Examples which will be described later, it has been confirmed that the algae belonging to *Galdieria* also have a haploid cell form and a diploid cell form. Therefore, the algae of the present embodiment may be algae belonging to *Galdieria*. Examples of the algae belonging to *Galdieria* include *Galdieria sulphuraria* (for example, an SAG108.79 strain), *Galdieria partita* (for example, an NBRC 102759 strain), and the like. Furthermore, the algae of the present embodiment may be algae belonging to *Cyanidium* other than the YFU3 strain and the HKN1 strain.

[Method for Producing Haploid Algae]

[0351] The present invention provides a method for producing a haploid cell of alga, including (a) step of culturing

a diploid cell of an alga belonging to Cyanidiophyceae having a diploid cell form and a haploid cell form and (b) step of isolating a haploid cell generated during the culturing.

[0352] The algae which belong to Cyanidiophyceae and have a diploid cell form and a haploid cell form (hereinafter, referred to as "the present algae" in some cases) are the same as the algae described above in "[Alga having a diploid cell form and a haploid cell form]". As the present algae, for example, the YFU3 strain, the HKN1 strain, mutants or analogues of these, and the like.

[0353] In addition, as shown in Examples which will be described later, it has been confirmed that the algae belonging to *Galdieria* (*Galdieria sulphuraria* SAG108.79 and *Galdieria partita* NBRC 102759) also have a diploid cell form and a haploid cell form. This result implies that in Cyanidiophyceae (for example, *Galdieria* and *Cyanidium*), probably, there may be a wide variety of algae having a diploid cell form and a haploid cell form. Therefore, the algae to which the producing method of the present embodiment can be applied may be algae belonging to *Galdieria* or *Cyanidium*, and are not limited to those exemplified above. Specific examples of the algae belonging to *Galdieria* include *Galdieria sulphuraria* and *Galdieria partita*.

[0354] According to the producing method of the present embodiment, haploid cells of algae can be prepared from diploid cells of the algae. Haploid algae can be transformed more easily compared to diploid algae.

(Step (a))

[0355] The step (a) is a step of culturing a diploid cell of an alga (present algae) which belongs to Cyanidiophyceae and has a diploid cell form and a haploid cell form.

[0356] The present algae are not particularly limited, and diploid cells of algae, which belong to Cyanidiophyceae and have been found to have a diploid cell form and a haploid cell form, may be cultured. Specific examples thereof include the YFU3 strain (diploid), the HKN1 strain (diploid), and allied species and mutants of these, and the like. In addition, examples thereof also include diploid cells of algae belonging to *Galdieria* (such as *Galdieria sulphuraria* and *Galdieria partita*), diploid cells of algae belonging to *Cyanidium*, and the like.

[0357] Examples of the culture conditions include the culture conditions exemplified in "[Alga belonging to Cyanidiophyceae]" or "[Alga having a diploid cell form and a haploid cell form]". Specifically, the pH condition is pH 1.0 to 6.0 for example, and is preferably pH 1.0 to 5.0. Furthermore, the temperature condition is 15° C. to 50° C. for example, and is preferably 30° C. to 50° C. Examples of the medium include those exemplified above in "[Alga belonging to Cyanidiophyceae]" or "[Alga having a diploid cell form and a haploid cell form]". Among these, a medium using acidic hot spring drainage is preferable. Specific examples of the medium using acidic hot spring drainage is include the Tsukahara mineral spring medium (Hirooka et al. 2016 Front in Microbiology).

[0358] The algae of the above embodiment are preferably cultured until the algae enter the stationary phase, and more preferably continuously cultured for an arbitrary period of time in the stationary phase. The duration of culturing is, for example, 2 to 60 days, 3 to 40 days, or 5 to 35 days. The period from the start of culturing to the stationary phase varies with the type of algae. Therefore, the duration of

culturing can be set according to the type of algae. Furthermore, cells may be collected from the culture solution in the stationary phase, subcultured, and further cultured for about 1 to 5 days.

(Step (b))

[0359] The step (b) is a step of isolating a haploid cell generated during the culturing.

[0360] By the culturing in the step (a), diploid cells of the present algae undergo meiosis, and as a result, haploid cells of the present algae are generated. In a case where the haploid cells are observed with an optical microscope, the shape of the haploid cells looks different from the shape of the diploid cells (for example, the haploid cells are found to have a size smaller than that of the diploid cells and found to have no rigid cell wall). Therefore, based on the difference in the cell shape, it is possible to discern the haploid cells. By collecting and isolating the haploid cells by using a Pasteur pipette or the like, it is possible to manufacture haploid cells of the algae.

[0361] The isolated haploid cells of the algae may be cultured using the medium exemplified in “[Alga belonging to Cyanidiophyceae]” or “[Alga having a diploid cell form and a haploid cell form]” described above, and the like. By isolating one by one the haploid cells that have appeared in the culture solution of the diploid cells and allowing the isolated cells to grow, it is possible to obtain monoclonal haploid cells of the algae.

[0362] In another aspect, the present invention provides a haploid cell of an alga (cell) (present alga) that belongs to Cyanidiophyceae and having a diploid cell form and a haploid cell form. The algae can be obtained by the method for producing haploid algae described above. The present invention also provides a haploid alga (cell) of the YFU3 strain, a haploid alga (cell) of the HKN1 strain, a haploid alga (cells) of algae belonging to *Cyanidium*, and a haploid alga (cell) of algae belonging to *Galdieria*. Specific examples of the algae belonging to *Galdieria* include *Galdieria sulphuraria* and *Galdieria partita*.

[Method for Producing a Diploid Alga]

[0363] The present invention provides a method for producing a diploid alga of an alga having a diploid cell form and a haploid cell form, the method including (a) step of mixing together two or more kinds of haploid cells and culturing the cells, and (b) step of isolating a diploid cell generated during the culturing.

[0364] As shown in Examples which will be described later, by mixing together and culturing haploid cells, diploid cells can be obtained.

(Step (a))

[0365] Step (a) is a step of mixing together and culturing two or more kinds of haploid cells of an alga having a diploid cell form and a haploid cell form.

[0366] The two or more kinds of haploid cells, which are mixed together in this step, are preferably algal cells of the same species. It is more preferable to use haploid cells derived from the diploid cells of the same strain.

[0367] Examples of the culture conditions include the culture conditions exemplified in “[Alga belonging to Cyanidiophyceae]” or “[Alga having a diploid cell form and a haploid cell form]” described above. Specifically, the pH

condition is pH 1.0 to 6.0 for example, and is preferably pH 1.0 to 5.0. Furthermore, the temperature condition is 15° C. to 50° C. for example, and is preferably 30° C. to 50° C. Examples of the medium include the media described in “[Alga belonging to Cyanidiophyceae]” or “[Alga having a diploid cell form and a haploid cell form]” described above.

[0368] The cells are preferably cultured, for example, for about 1 to 4 weeks, and may be further subcultured and then cultured for about 3 to 10 days.

(Step (b))

[0369] The step (b) is a step of isolating a diploid cell generated during the culturing.

[0370] By the culturing in the step (a), haploid cells are mated to each other, and as a result, diploid cells are generated. By collecting and isolating the diploid cell by using a Pasteur pipette or the like, a diploid alga can be prepared.

[0371] The isolated diploid algae may be cultured using the medium exemplified in “[Alga belonging to Cyanidiophyceae]” or “[Alga having a diploid cell form and a haploid cell form]” described above (preferably an artificial synthetic medium such as M-Allen medium), and the like.

[0372] In another aspect, the present invention provides a diploid alga (cell) of an alga (present alga) which belongs to Cyanidiophyceae and has a diploid cell form and a haploid cell form. The alga can be obtained by the method for producing a diploid alga described above. The present invention also provides a diploid alga (cell) of the YFU3 strain, a diploid alga (cell) of the HKN1 strain, and a diploid alga (cell) of an alga belonging to *Cyanidioschyzon*. Specific examples of the algae belonging to *Cyanidioschyzon* include *Cyanidioschyzon merolae*.

[Culture of the Present Algae]

[0373] The present invention provides an alga culture containing algae (present algae) which belong to Cyanidiophyceae and have a diploid cell form and a haploid cell form. In the alga culture of the present embodiment, the ratio of the number of haploid algal cells to the total number of algal cells is 70% to 100%.

[0374] Presumably, in the natural world, the present algae may exist as both the diploid cells and haploid cells. However, all of the known algae belonging to *Galdieria* and *Cyanidium* are discovered as cells having a rigid cell wall. Therefore, it is considered that most of the present algae may exist as diploid cells in nature.

[0375] On the other hand, according to the method for producing a haploid alga of the above embodiment, when the diploid cells of the present algae are cultured until they enter the stationary phase and then continuously cultured as they are, the diploid cells of the present algae undergo meiosis, and as a result, haploid cells of the present algae appear. The haploid cells of the present algae grow by two-cell division, and the proportion of the haploid cells of the present algae increases as the culture is continued. Then, by further continuing the culture or isolating the haploid cells and culturing them, it is possible to obtain an alga culture in which the proportion of the haploid cells of the present algae is 70% to 100%.

[0376] The present algae are not particularly limited, and diploid cells of algae which belong to Cyanidiophyceae and are confirmed to have a diploid cell form and a haploid cell

form may be cultured. Specific examples thereof include the YFU3 strain (diploid), the HKN1 strain (diploid), allied species and mutants of these, and the like. In addition, examples thereof also include diploid cells of algae belonging to *Galdieria* (such as *Galdieria sulphuraria* and *Galdieria partita*), diploid cells of algae belonging to *Cyanidium*, and the like.

[0377] Examples of the culture conditions for the present algae in the form of diploid cells are the same as the culture conditions exemplified in “[Alga belonging to Cyanidiophyceae]” or “[Alga having a diploid cell form and a haploid cell form]” described above.

[0378] In the alga culture of the present embodiment, the proportion of the haploid cells is 70 to 100%, which is significantly different from the proportion of haploid cells in nature.

[0379] In another aspect, the present invention provides a cell population including diploid cells of algae which belong to Cyanidiophyceae and have a diploid cell form and a haploid cell form. As the algae belonging to Cyanidiophyceae, for example, the YFU3 strain, the HKN1 strain, mutants of these, and algae belonging to *Cyanidioschyzon*. Examples of the algae belonging to *Cyanidioschyzon* include *Cyanidioschyzon merolae*.

[0380] In another aspect, the present invention provides a cell population including haploid cells of algae which belong to Cyanidiophyceae and have a diploid cell form and a haploid cell form. As the algae belonging to Cyanidiophyceae, for example, the YFU3 strain, the HKN1 strain, mutants of these, algae belonging to *Cyanidium* other than the above, and algae belonging to *Galdieria*. Examples of the algae belonging to *Galdieria* include *Galdieria sulphuraria* and *Galdieria partita*.

[0381] In another aspect, the present invention provides a cell population of algae that belong to Cyanidiophyceae and have a diploid cell form and a haploid cell form, in which the diploid cells and the haploid cells are mixed together.

[Drying and Swelling Treatment Product of the Present Algae]

[0382] The present invention provides a dry and swollen product obtained by performing a drying and swelling treatment on algae (present algae) which belong to Cyanidiophyceae and have a diploid cell form and a haploid cell form.

[0383] By performing a drying and swelling treatment on the present algae, it is possible to kill the algal cells. Therefore, even a transformant obtained not using self-cloning can be handled in an open system. “Drying and swelling treatment” refers to a treatment of drying algal cells and then adding an aqueous medium or the like to the dried product so as to wet again the dried product. Specific examples of the drying and swelling treatment include the cell disruption treatment described above in (3).

[0384] The present algae subjected to the drying and swelling treatment may be diploid cells, haploid cells, or a mixture of diploid cells and haploid cells. For example, the present algae may be cells collected from the alga culture of the above embodiment. The ploidy-variable algae subjected to the drying and swelling treatment are preferably haploid cells.

[Extract of the Present Algae]

[0385] As described above, the present algae are rich in nutrients such as amino acids, vitamins, proteins, lipids, and dietary fiber. Accordingly, the present algae can be used as a nutrient composition or a nutritional supplement which will be described below. In addition, in a case where the present algae are blended with foods, feeds, pet foods, cosmetics, and the like, it is possible to provide nutrient-enriched products. Furthermore, in the products described above, an extract of the present algae may be used instead of or together with the present algae. Therefore, the present invention also provides an extract of the present algae.

[0386] In the present specification, “extract of the present algae” refers to a substance obtained by extracting intracellular components from cells of the present algae by performing a physical treatment or a chemical treatment. For example, the extract of the present algae may be a cell disruption product obtained by disrupting the cells of the present algae by means of a physical treatment or a chemical treatment. In addition, the extract of the present algae may be a substance obtained by concentrating the cell disruption product, a substance obtained by removing solid contents from the cell disruption product, or a substance obtained by separating some components from the cell disruption product.

[0387] The method of the physical treatment or the chemical treatment for the cells is not particularly limited, and the methods generally used for cell disruption can be used. Examples of the physical treatment include cell disruption performed by means of glass beads, a mortar, an ultrasonic treatment, a French press, a homogenizer, and the like.

[0388] Examples of the chemical treatment include a neutralization treatment, a hypotonic treatment, a freeze-thaw treatment, a drying and swelling treatment, and the like. More specifically, examples thereof include the cell rupturing treatments (A) to (C) described above.

[0389] In a case where the cell disruption product is concentrated, the concentration method is not particularly limited, and a generally used concentration method may be used. Examples of the concentration method for the cell disruption product include drying, freeze-drying, drying under reduced pressure, and the like.

[0390] Furthermore, in a case where solid contents are removed from the cell disruption product, the solid content-removing method is not particularly limited, and it is possible to use a method that is generally used for removing solid contents. Examples of the solid content-removing method include filtration, centrifugation, and the like.

[0391] In a case where some components are separated from the cell disruption product, the separation method is not particularly limited, and it is possible to use a method generally used for separation, purification, and the like for biochemical substances. Examples of the separation method include salting out, dialysis, solvent extraction, adsorption, column chromatography, ion exchange chromatography, and the like. Each of these methods may be used singly. Alternatively, two or more kinds of treatments may be used in combination. Here, a substance purified into a single component is excluded from “extract of the present algae”. The extract of the present algae preferably contains 10 or more kinds of cell components of the present algae, more preferably contains 15 or more kinds of cell components of the present algae, and even more preferably contains 20 or more kinds of cell components of the present algae.

[0392] The extract of the present algae preferably contains a nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber, and more preferably contains a nutrient selected from the group consisting of amino acids and vitamins. Specific examples of the amino acids include isoleucine, leucine, lysine, methionine, cysteine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, γ -aminobutyric acid, and the like. Specific examples of the vitamins include vitamin A, β -carotene, vitamin C, vitamin E, vitamin K₁, vitamin K₂, folic acid, and the like.

[Nutrient Composition]

[0393] In one embodiment, the present invention provides a nutrient composition containing the algae belonging to Cyanidiophyceae or an extract thereof.

[0394] The nutrient composition of the present embodiment contains the algae belonging to Cyanidiophyceae or an extract thereof. Accordingly, the nutrient composition is rich in nutrients abundantly contained in the algae. For example, the nutrient composition can be rich in nutrients selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber. Particularly, the nutrient composition can be rich in nutrients selected from the group consisting of amino acids and vitamins. Therefore, the nutrient composition can be used by being blended with a nutritional supplement, a food, a feed, a pet food, a cosmetic, or the like which will be described later.

[0395] The algae belonging to Cyanidiophyceae are not particularly limited. As the algae, for example, the present algae or algae belonging to *Cyanidioschyzon* are preferable.

[0396] The present algae may be diploid cells, haploid cells, or a mixture of diploid cells and haploid cells. For example, the present algae may be cells collected from the alga culture of the above embodiment. Examples of the present algae contained in the nutrient composition of the present embodiment include those exemplified in “[Alga having a diploid cell form and a haploid cell form]” described above. Specific examples of the present algae include the YFU3 strain, the HKN1 strain, allied species and mutants of these, and the like. Furthermore, examples of the present algae also include algae belonging to *Cyanidium* other than the above and algae belonging to *Galdieria*. Among these, the YFU3 strain, the HKN1 strain, mutants of these, and the algae belonging to *Galdieria* are preferable, and the YFU3 strain, the HKN1 strain, and mutants of these are more preferable.

[0397] In a case where either the diploid cells or the haploid cells having no rigid cell wall, it is preferable to use the cells having no rigid cell wall. Using such cells is advantageous because nutrients can be easily extracted. Furthermore, using such cells is advantageous because even in a case where the algal cells are directly blended with the nutrient composition, the nutrients in the algal cells are easily absorbed after the intake of the nutrient composition. For example, in a case where the YFU3 strain, the HKN1 strain, or mutants of these are used, it is preferable to use the YFU3 strain (haploid), the HKN1 strain (haploid), or the mutants of these. In addition, for example, haploid cells of the algae belonging to *Galdieria* (for example, *Galdieria sulphuraria*, *Galdieria partita*, and the like) and haploid cells of the algae belonging to *Cyanidium* are also preferable. Among these, the YFU3 strain (haploid), the HKN1

strain (haploid), mutants of these, and haploid cells of the algae belonging to *Galdieria* are preferable.

[0398] The algae belonging to Cyanidiophyceae may be algae belonging to Cyanidiophyceae other than the present algae. Among these, the algae belonging to *Cyanidioschyzon* have no rigid cell wall. Therefore, their cells can be disrupted by a relatively mild treatment such as a neutralization treatment, a hypotonic treatment, a freeze-thaw treatment, or the like. Accordingly, the algae belonging to *Cyanidioschyzon* (for example, *Cyanidioschyzon merolae*) are a preferred example of the algae belonging to Cyanidiophyceae.

[0399] By being cultured and allowed to grow using an appropriate medium, collected through a known method such as centrifugation or filtration, and subjected to washing, drying, and the like as appropriate, the algae belonging to Cyanidiophyceae can be used in the nutritional supplement of the present embodiment. Alternatively, by being collected from the alga culture of the above embodiment and subjected to washing, drying, and the like as appropriate, the algal cells may be used in the nutrient composition of the present embodiment.

[0400] Furthermore, the nutrient composition of the present embodiment may contain an extract of the algae belonging to Cyanidiophyceae instead of or together with the algae belonging to Cyanidiophyceae.

[0401] The nutrient composition of the present embodiment may contain other components as appropriate in addition to the algae belonging to Cyanidiophyceae or an extract thereof. Those other components are not particularly limited, and examples thereof include a pharmaceutically acceptable carrier and the like. “Pharmaceutically acceptable carrier” means a carrier which does not inhibit the function of nutrients contained in the algae belonging to Cyanidiophyceae and substantially does not exhibit toxicity to an administration subject. “Substantially does not exhibit toxicity” means that the component does not exhibit toxicity to the administration subject at the usual dose thereof. Examples of the pharmaceutically acceptable carrier include, but are not limited to, excipients, binders, disintegrants, lubricants, emulsifiers, stabilizers, diluents, oleaginous bases, thickeners, antioxidants, reducing agents, oxidants, chelating agents, solvents, and the like. One kind of pharmaceutically acceptable carrier may be used singly, or two or more kinds of pharmaceutically acceptable carriers may be used in combination.

[0402] The content of the algae belonging to Cyanidiophyceae or an extract thereof in the nutrient composition of the present embodiment is not particularly limited, and can be appropriately selected, for example, in a range of 1% to 100% by mass. The content of the algae belonging to Cyanidiophyceae or an extract thereof in the nutritional supplement of the present embodiment is preferably 50% to 100% by mass, more preferably 60% to 100% by mass, and even more preferably 70% to 100% by mass.

[0403] The algae belonging to Cyanidiophyceae or an extract thereof can be appropriately mixed with other components and made into a form such as dry powder, granules, tablets, jelly, a liquid, or capsules according to a common method.

[Nutritional Supplement]

[0404] In one embodiment, the present invention provides a nutritional supplement containing the algae belonging to Cyanidiophyceae or an extract thereof.

[0405] Nutritional supplements are used by human beings or non-human animals for supplying nutrients, and there is no particular limitation on the dosage form, the degree of purification, and the like thereof. As described above, the algae belonging to Cyanidiophyceae are rich in nutrients such as amino acids and vitamins. Therefore, by using the algae belonging to Cyanidiophyceae or an extract thereof, it is possible to obtain a nutritional supplement rich in nutrients such as amino acids and vitamins.

[0406] The nutritional supplement of the present embodiment is rich in nutrients such as amino acids and vitamins. Therefore, the nutritional supplement of the present embodiment can be used for human beings or non-human animals as a nutritional supplement for supplying the nutrients. Examples of the nutrients supplied by the nutritional supplement of the present embodiment include amino acids, vitamins, proteins, lipids, dietary fiber, and the like. Particularly, the nutritional supplement of the present embodiment can be suitably used for supplying amino acids (such as isoleucine, leucine, lysine, methionine, cystine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, and γ -aminobutyric acid) and vitamins (such as vitamin A, β -carotene, vitamin B1, vitamin B2, vitamin B6, vitamin C, vitamin E, vitamin K₁, vitamin K₂, niacin, inositol, folic acid, and biotin). In a case where the present algae are algae belonging to Cyanidiophyceae, the nutritional supplement of the present embodiment can be particularly suitably used for supplying amino acids (such as isoleucine, leucine, lysine, methionine, cystine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, and γ -aminobutyric acid) and vitamins (such as vitamin A, β -carotene, vitamin B1, vitamin B2, vitamin B6, vitamin C, vitamin E, vitamin K₁, vitamin K₂, niacin, inositol, folic acid, and biotin).

[0407] The nutritional supplement of the present embodiment can be particularly suitably used for supplying γ -aminobutyric acid, β -carotene, vitamin C, vitamin E, vitamin K₁, vitamin K₂, and folic acid.

[0408] The nutritional supplement of the present embodiment may contain other components as appropriate in addition to the algae belonging to Cyanidiophyceae and an extract thereof. Those other components are not particularly limited, and examples thereof include a pharmaceutically acceptable carrier and the like. Examples of the pharmaceutically acceptable carrier are the same as those exemplified in "Nutrient composition]" described above. One kind of pharmaceutically acceptable carrier may be used singly, or two or more kinds of pharmaceutically acceptable carriers may be used in combination.

[0409] The content of the algae belonging to Cyanidiophyceae or an extract thereof in the nutritional supplement of the present embodiment is not particularly limited, and can be appropriately selected, for example, in a range of 1% to 100% by mass. The content of the algae belonging to Cyanidiophyceae or an extract thereof in the nutritional supplement of the present embodiment is preferably 50% to 100% by mass, more preferably 60% to 99% by mass, and even more preferably 70% to 99% by mass.

[0410] The algae belonging to Cyanidiophyceae or an extract thereof can be appropriately mixed with other components and made into a form such as dry powder, granules, tablets, jelly, a liquid, or capsules according to a common method.

[0411] Furthermore, the nutritional supplement of the present embodiment may be used as it is for human beings or non-human organisms, or may be used by being blended with a composition for supplying nutrients such as a food, a feed, a pet food, or a cosmetic which will be described later. By blending the nutritional supplement of the present embodiment, it is possible to prepare a composition suitable for supplying nutrients such as amino acids and vitamins.

[0412] In another aspect, the present invention provides a method for producing a nutritional supplement, including a step of culturing algae belonging to Cyanidiophyceae, a step of collecting the cultured algae belonging to Cyanidiophyceae, and a step of formulating the collected algae belonging to Cyanidiophyceae.

[0413] In another aspect, the present invention provides a method for producing a nutritional supplement, including a step of culturing algae belonging to Cyanidiophyceae, a step of collecting the cultured algae belonging to Cyanidiophyceae, a step of obtaining an extract of the collected algae belonging to Cyanidiophyceae, and a step of formulating the extract of the algae belonging to Cyanidiophyceae.

[Composition for Supplying Nutrients]

[0414] In one embodiment, the present invention provides a composition for supplying nutrients containing the nutritional supplement described above.

[0415] In the present specification, "composition for supplying nutrients" refers to a composition used by human beings or non-human animals for the intake of nutrients into the body. The intake of nutrients into the body may be oral or parenteral.

[0416] The composition for supplying nutrients of the present embodiment is rich in amino acids, vitamins, proteins, lipids, dietary fiber, and the like. These components are nutrients abundantly contained in the aforementioned nutritional supplement.

[0417] One of the characteristics of the composition for supplying nutrients of the present embodiment is that the composition is rich in amino acids and vitamins among the above nutrients.

[0418] The composition is characterized by being particularly rich in γ -aminobutyric acid among amino acids and in β -carotene, vitamin C, vitamin E, vitamin K₁, vitamin K₂, and folic acid among vitamins. For example, γ -aminobutyric acid is known to have a brain function improving effect and a blood pressure lowering effect. Furthermore, β -carotene, vitamin C and vitamin E are known to have an antioxidant activity. In addition, vitamin K₁ and vitamin K₂ are known to have an osteoporosis improving effect. Moreover, folic acid is known to be necessary for the development of the fetus during pregnancy, and is reported to have a cardiovascular disease improving effect.

[0419] Therefore, the intake of the composition for supplying nutrients of the present embodiment can bring about the physical condition improving effects that the aforementioned nutrients have. Therefore, the composition for supplying nutrients of the present embodiment is suitably used for supplying at least one kind of nutrient selected from amino acids, vitamins, proteins, lipids, and dietary fiber. Furthermore, the composition for supplying nutrients of the present embodiment is more suitably used for supplying at least one kind of nutrient selected from the group consisting of amino acids and vitamins.

[0420] The composition for supplying nutrients of the present embodiment is not particularly limited as long as the composition is used by human beings or non-human animals for the intake of nutrients into the body. Examples of the composition for supplying nutrients of the present embodiment include foods, feeds, pet foods, cosmetics, and the like.

(Food)

[0421] The composition for supplying nutrients of the present embodiment may be a food. Therefore, the present invention also provides a food containing the nutrient composition or the nutritional supplement of the above embodiment. In addition, the present invention also provides a food containing the algae belonging to Cyanidiophyceae or an extract thereof.

[0422] In a case where the composition for supplying nutrients of the present embodiment is a food, the nutrient composition or the nutritional supplement described above may be added to the food as a food additive. By adding the aforementioned nutrient composition or nutritional supplement to a food, it is possible to prepare a food enriched with nutrients contained in the aforementioned nutrient composition or nutritional supplement. Therefore, in another aspect, the present invention provides a food additive containing the algae belonging to Cyanidiophyceae or an extract thereof. By adding the aforementioned nutrient composition or nutritional supplement to a raw material of a food and adding other food additives thereto as appropriate, the food of the present embodiment can be manufactured by a known method according to the type of the food.

[0423] In the food of the present embodiment, the type of the food is not particularly limited. Examples of the food include, but are not limited to, various noodles such as

[0424] Soba, Udon, Harusame, Chinese noodles, instant noodles, and cup noodles; carbohydrates such as bread, flour, rice flour, pancake, and mashed potatoes; beverages such as green vegetable juice, soft drinks, carbonated drinks, nutritious drinks, fruit drinks, vegetable drinks, lactic acid drinks, milk-based drinks, sports drinks, tea, and coffee; bean products such as tofu, Okara, and Natto; various soups such as curry roux, stew, and instant soups; frozen desserts such as ice cream, ice sorbet, and shaved ice; confectioneries such as cookies, candies, gums, chocolates, tablet-shaped sweets, snacks, biscuits, jellies, jams, creams, and other baked cakes; processed marine and livestock products such as Kamaboko, Hanpen, hams, and sausages; dairy products such as processed milk, fermented milk, butter, cheese, and yogurt, oil and fat as well as oil and fat processed foods such as salad oil, oil for frying, margarine, mayonnaise, shortening, whipping cream, and dressing; flavors such as sauces, dressings, Miso, soy sauce, and seasonings; other processed foods such as various retort foods, Furikake, and pickles; and the like.

[0425] In the foods described above, the content of the aforementioned nutrient composition or the nutritional supplement is not particularly limited, and may be appropriately set according to the type of the food. Considering the flavor of the food and the like, the content of the aforementioned nutrient composition or nutritional supplement in the food can be, for example, 0.01% to 30% by mass as the content of the algae belonging to Cyanidiophyceae or an extract thereof. From the viewpoint of flavor of the food and the like, the content of the nutrient composition or the nutritional supplement is preferably 0.05% to 20% by mass,

more preferably 0.1% to 15% by mass, even more preferably 0.1% to 10% by mass, and particularly preferably 0.1% to 5% by mass.

[0426] The food may be a functional food or a dietary supplement. The functional food or the dietary supplement may be in the form of a general food as described above, or in the form of dry powder, granules, tablets, jellies, drinks, and the like. In this case, by mixing the aforementioned nutrient composition or nutritional supplement with other components as appropriate, foods in the form of dry powders, granules, tablets, jellies, drinks, and the like can be prepared according to a common method. Those other components are not particularly limited, and examples thereof include a pharmaceutically acceptable carrier and the like. Examples of the pharmaceutically acceptable carrier are the same as those exemplified in “[Nutrient composition]” described above. Furthermore, in order to improve the flavor and the like, sweeteners, flavoring agents, various flavors, spices, oils and fats, other food additives, and the like may be used as those other components. One kind of each of those other components may be used singly, or two or more kinds of those other components may be used in combination.

[0427] In the functional food or the dietary supplement described above, the content of the aforementioned nutrient composition or nutritional supplement is not particularly limited, and may be appropriately set according to the type of the functional food or the dietary supplement. In a case where the functional food or the dietary supplement is in the form of dry powder, granules, tablets, or the like, in the functional food or the dietary supplement, the content of the nutrient composition or the nutritional supplement described above is, for example, 0.1% to 99% by mass as the content of the algae belonging to Cyanidiophyceae or an extract thereof.

[0428] From the viewpoint of flavor and efficient supplement of nutrients, the content of the nutrient composition or the nutritional supplement described above is preferably 1% to 90% by mass, more preferably 10% to 85% by mass, even more preferably 20% to 85% by mass, and particularly preferably 25% to 85% by mass. Furthermore, in a case where the functional food or the dietary supplement is in the form of a jelly, a drink, or the like, in the functional food or the dietary supplement, the content of the nutritional supplement described above is, for example, 0.05% to 80% by mass as the content of the algae belonging to Cyanidiophyceae or an extract thereof. From the viewpoint of flavor and efficient supplement of nutrients, the content of the nutritional supplement described above is preferably 1% to 75% by mass, more preferably 10% to 70% by mass, even more preferably 15% to 70% by mass, and particularly preferably 20% to 70% by mass.

[0429] The food of the present embodiment can be ingested for efficiently supplying the aforementioned nutrients contained in the algae belonging to Cyanidiophyceae. Particularly, the food of the present embodiment can be ingested for efficiently supplying nutrients selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber. The food of the present embodiment is effective for the intake of nutrients selected from the group consisting of amino acids and vitamins among the nutrients described above.

(Feed and Pet Food)

[0430] The composition for supplying nutrients according to the present embodiment may be a feed or a pet food. Accordingly, the present invention also provides a feed or a pet food containing the nutrient composition or the nutritional supplement of the above embodiment. In addition, the present invention also provides a feed or a pet food which contains the algae belonging to Cyanidiophyceae or an extract thereof.

[0431] In a case where the composition for supplying nutrients of the present embodiment is a feed or a pet food, the nutrient composition or the nutritional supplement described above may be added to the feed or the pet food as a feed additive or a pet food additive. By adding the aforementioned nutrient composition or nutritional supplement to a feed or a pet food, it is possible to prepare a feed or a pet food enriched with nutrients contained in the nutrient composition or the nutritional supplement described above. Therefore, in another aspect, the present invention provides a feed additive or a pet food additive containing the algae belonging to Cyanidiophyceae or an extract thereof.

[0432] By adding the aforementioned nutrient composition or nutritional supplement to raw materials of a feed or raw materials of a pet food and adding other feed additives or pet food additives as appropriate, the feed or the pet food of the present embodiment can be manufactured by a known method according to the type of the raw materials of the feed or the pet food.

[0433] The type of animal to which the feed or the pet food of the present embodiment is provided is not particularly limited. Examples of thereof include, but are not limited to, livestock (such as cows, pigs, chickens, horses, sheep, and goats), fish, shellfish, pets (such as dogs, cats, hamsters, rabbits, parakeets, tropical fish, reptiles, amphibians, and insects), and the like.

[0434] In the feed or the pet food of the present embodiment, the content of the aforementioned nutrient composition or nutritional supplement is not particularly limited, and may be appropriately set according to the type of the feed or the pet food. In the feed or the pet food, the content of the aforementioned nutrient composition or nutritional supplement is, for example, 0.01% to 90% by mass as the content of the algae belonging to Cyanidiophyceae or an extract thereof. The content of the aforementioned nutrient composition or nutritional supplement is preferably 0.1% to 80% by mass, more preferably 1% to 70% by mass, and particularly preferably 1% to 60% by mass.

[0435] The feed or the pet food of the present embodiment can be used for efficiently supplying the aforementioned nutrients, which are contained in the algae belonging to Cyanidiophyceae, to the animal. The feed or the pet food of the present embodiment can be used particularly for efficiently supplying nutrients, which are selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber, to the animal. The feed or the pet food of the present embodiment is effective for the intake of nutrients selected from the group consisting of amino acids and vitamins among the above nutrients.

(Cosmetic)

[0436] The composition for supplying nutrients according to the present embodiment may be a cosmetic. Therefore, the present invention also provides a cosmetic containing the

nutrient composition or the nutritional supplement of the above embodiment. In addition, the present invention also provides a cosmetic containing the algae belonging to Cyanidiophyceae or an extract thereof.

[0437] In a case where the composition for supplying nutrients of the present embodiment is a cosmetic, the nutrient composition or the nutritional supplement described above may be added to the cosmetic as a cosmetic additive. By adding the aforementioned nutrient composition or nutritional supplement to the cosmetic, it is possible to prepare a cosmetic containing the nutrients contained in the nutrient composition or the nutritional supplement described above. Therefore, in another aspect, the present invention provides a cosmetic additive containing the algae belonging to Cyanidiophyceae or an extract thereof.

[0438] By mixing the aforementioned nutrient composition or nutritional supplement with other components as appropriate, the cosmetic of the present embodiment can be manufactured by a known method according to the type of the cosmetic. Those other components are not particularly limited, and examples thereof include a pharmaceutically acceptable carrier and the like. Examples of the pharmaceutically acceptable carrier are the same as those exemplified in “[Nutrient composition]” described above. In addition, materials known as cosmetic additives may be used as those other components. One kind of each of those other components may be used singly, or two or more kinds of those other components may be used in combination.

[0439] In the cosmetic of the present embodiment, the type of the cosmetic is not particularly limited. Examples of the cosmetic include, but are not limited to, skin care cosmetics such as toners, emulsions, lotions, creams, gels, sunscreens, facial packs, masks, and serums; makeup products such as foundations, makeup bases, lipsticks, lip glosses, and blushers; cleansers such as facial cleansers, body washes, and cleansing agents; hair care cosmetics such as shampoos, rinses, hair conditioners, treatments, and hair styling agents; body care cosmetics such as body powders and body lotions; and the like.

[0440] In the cosmetic of the present embodiment, the content of the aforementioned nutrient composition or nutritional supplement is not particularly limited, and may be appropriately set according to the type of the cosmetic. Considering the texture of the cosmetic and the like, the content of the aforementioned nutrient composition or nutritional supplement in the cosmetic can be, for example, 0.01% to 30% by mass as the content of the algae belonging to Cyanidiophyceae or an extract thereof. From the viewpoint of texture of the cosmetic and the like, the content of the nutrient composition or the nutritional supplement is preferably 0.1% to 20% by mass, more preferably 0.1% to 15% by mass, even more preferably 0.1% to 10% by mass, and particularly preferably 0.1% to 5% by mass.

[0441] The cosmetic of the present embodiment can be used by being applied to the skin or hair so as to supply the aforementioned nutrients, which are contained in the algae belonging to Cyanidiophyceae, to the skin or hair. Particularly, the cosmetic of the present embodiment can be used by being applied to the skin or hair so as to supply nutrients, which are selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber, to the skin or hair. The cosmetic of the present embodiment is effective for supplying nutrients selected from the group consisting of amino acids and vitamins among the above nutrients.

[0442] In another aspect, the present invention provides a method for producing a food, including a step of blending algae belonging to Cyanidiophyceae or an extract thereof with a food.

[0443] In another aspect, the present invention provides a method for producing a functional food or a dietary supplement, including a step of mixing algae belonging to Cyanidiophyceae or an extract thereof with a pharmaceutically acceptable carrier.

[0444] In another aspect, the present invention provides a method for producing a feed or a pet food, including a step of blending algae belonging to Cyanidiophyceae or an extract thereof with a feed or a pet food.

[0445] In another aspect, the present invention provides a method for producing a cosmetic, including a step of blending algae belonging to Cyanidiophyceae or an extract thereof with a cosmetic.

[0446] In another aspect, the present invention provides a method for producing a food additive, a feed additive, a pet food additive, or a cosmetic additive, including a step of culturing algae belonging to Cyanidiophyceae, a step of collecting the cultured algae belonging to Cyanidiophyceae, and a step of formulating the collected algae belonging to Cyanidiophyceae.

[0447] In another aspect, the present invention provides a method for producing a food additive, a feed additive, a pet food additive, or a cosmetic additive, including a step of culturing algae belonging to Cyanidiophyceae, a step of collecting the cultured algae belonging to Cyanidiophyceae, a step of obtaining an extract of the collected algae belonging to Cyanidiophyceae, and a step of formulating the extract of the algae belonging to Cyanidiophyceae.

[Method for Producing Nutrient]

[0448] In one embodiment, the present invention provides a method for producing a nutrient. The method for producing a nutrient of the present embodiment includes (a) step of disrupting cells of algae belonging to Cyanidiophyceae so as to obtain a cell disruption product and (b) step of separating at least one kind of nutrient from the cell disruption product.

[0449] Hereinafter, each of the steps of the producing method of the present embodiment will be described.

(Step (a))

[0450] The step (a) is a step of disrupting cells of algae belonging to Cyanidiophyceae so as to obtain a cell disruption product.

[0451] As the algae belonging to Cyanidiophyceae used in this step, for example, the present algae described above and algae belonging to *Cyanidioschyzon* (for example, *Cyanidioschyzon merolae*) are suitable. As the present algae, for example, the same algae as those listed in “[Alga having a diploid cell form and a haploid cell form]” described above are suitable. Specific examples thereof include the YFU3 strain, the HKN1 strain, allied species and mutants of these, and the like. Furthermore, examples thereof also include algae belonging to *Cyanidium* other than the above and algae belonging to *Galdieria*. Among these, the YFU3 strain, the HKN1 strain, mutants of these, and the algae belonging to *Galdieria* are preferable, and the YFU3 strain, the HKN1 strain, and mutants of these are more preferable.

[0452] The present algae may be diploid cells, haploid cells, or a mixture of diploid cells and haploid cells.

[0453] For example, the present algae may be cells collected from the alga culture of the above embodiment. Therefore, the producing method of the present embodiment also provides a method for producing a nutrient, including (a) step of collecting algae from the alga culture of the above embodiment, (b) step of disrupting cells of the algae so as to obtain a cell disruption product, and (c) step of separating at least one kind of nutrient from the cell disruption product.

[0454] In a case where either the diploid cells or the haploid cells have no rigid cell wall, it is preferable to use the cells having no rigid cell wall. The cells having no rigid cell wall can be disrupted by a relatively mild treatment. For example, in a case where the YFU3 strain, the HKN1 strain, or mutants of these are used, it is preferable to use the YFU3 strain (haploid), the HKN1 strain (haploid), or mutants of these. In addition, for example, haploid cells of the algae belonging to *Galdieria* (such as *Galdieria sulphuraria*, and *Galdieria partita*) and haploid cells of the algae belonging to *Cyanidium* are also preferable. Among these, the YFU3 strain (haploid), the HKN1 strain (haploid), mutants of these, and haploid cells of the algae belonging to *Galdieria* are preferable.

[0455] The method for disrupting the cells of the algae belonging to Cyanidiophyceae is not particularly limited, and a known method can be used. Examples of the method for disrupting the cells include physical treatments by means of glass beads, a mortar, an ultrasonic treatment, a French press, a homogenizer, and the like; chemical treatments such as a neutralization treatment, a hypotonic treatment, a freeze-thaw treatment, and a drying and swelling treatment; and the like. One kind of each of these treatments may be performed singly, or two or more kinds of these treatments may be performed in combination.

[0456] In a case where the algae belonging to Cyanidiophyceae have no rigid cell wall, it is possible to disrupt the cells by relatively mild treatment such as a neutralization treatment, a hypotonic treatment, a freeze-thaw treatment, or a drying and swelling treatment. These treatments bear energy costs lower than that of physical treatments and can be performed in a simple manner. Therefore, in a case where the algae belonging to Cyanidiophyceae that do not have a rigid cell wall are used, as the cell disruption method, a neutralization treatment, a hypotonic treatment, a freeze-thaw treatment, and a drying and swelling treatment are preferable. Furthermore, any of the cell rupturing treatments (1) to (3) described above is also an example suitable as the cell disruption method.

[0457] Examples of the neutralization treatment method include a method of immersing the algal cells, which belong to Cyanidiophyceae and have no rigid cell wall, in a neutralizing solution at a pH of about 7 to 10. The algae belonging to Cyanidiophyceae are adapted to the acidic pH range. Therefore, in a case where the algae have no rigid cell wall, the cells are disrupted by being immersed in the neutralizing solution which is neutral or basic. The composition of the neutralizing solution is not particularly limited, and for example, a buffer such as a phosphate buffer and a Tris buffer can be used. The time for which the cells are immersed in the neutralizing solution may be set such that the cells are disrupted. For example, the cells are immersed in the neutralizing solution for about one week. For example, the cell rupturing treatment described above in (1) is suitable.

[0458] Examples of the hypotonic treatment method include a method of immersing the algal cells belonging to Cyanidiophyceae in a hypotonic solution such as water. Among the algae belonging to Cyanidiophyceae, those having no rigid cell wall undergo cell rupture by being immersed in a hypotonic solution such as water. The composition of the hypotonic solution is not particularly limited as long as it is a hypotonic liquid in which the algal cells belonging to Cyanidiophyceae are ruptured. Examples of the hypotonic solution include water, a buffer with a low salt concentration, and the like. The time for which the cells are immersed in the hypotonic solution may be set such that the cells are ruptured. For example, the cells are immersed in the hypotonic solution for about 1 to 30 minutes. Furthermore, after the immersion in the hypotonic solution, a process of collecting the algal cells by centrifugation or the like and resuspending the cells in the hypotonic solution may be repeated. The number of times of resuspension is not particularly limited, but is, for example, 1 to 5. For example, the cell rupturing treatment described above in (2) is suitable.

[0459] Examples of the freeze-thaw treatment method include a method in which a cycle of freezing and thawing is performed once or more on the algal cells belonging to Cyanidiophyceae. The number of cycles of freezing and thawing is not particularly limited, and may be set such that the algal cells belonging to Cyanidiophyceae having no rigid cell wall are disrupted. The number of cycles of freezing and thawing is, for example, about 1 to 5. Each of the freezing time and the thawing time is not particularly limited, but is, for example, about 10 to 30 minutes.

[0460] Examples of the drying and swelling treatment method include a method in which a cycle of drying and resuspension in a buffer is performed once or more on the algal cells. The number of cycles of drying and resuspension is not particularly limited, and may be set such that the algal cells belonging to Cyanidiophyceae having no rigid cell wall are disrupted. The number of cycles of drying and resuspension is, for example, about 1 to 5. For example, the cell rupturing treatment described above in (3) is suitable.

[0461] By disrupting the algal cells belonging to Cyanidiophyceae by the methods described above, a cell disruption product of the algae belonging to Cyanidiophyceae can be obtained.

(Step (b))

[0462] The step (b) is a step of separating at least one kind of nutrient from the cell disruption product of the algae belonging to Cyanidiophyceae.

[0463] In this step, the nutrient to be separated is not particularly limited as long as it is a nutrient of the algae belonging to Cyanidiophyceae. As described above in “[Alga having a diploid cell form and a haploid cell form]”, the algae belonging to Cyanidiophyceae are rich in nutrients such as amino acids, vitamins, proteins, lipids, and dietary fiber. Therefore, examples of the nutrient to be separated in this step include at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber.

[0464] In this step, it is preferable to separate at least one kind of nutrient selected from the group consisting of amino acids and vitamins among the nutrients described above. Specific examples of the amino acids include at least one kind of amino acid selected from the group consisting of isoleucine, leucine, lysine, methionine, cystine, phenylala-

nine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, and γ -aminobutyric acid. As the amino acid, for example, γ -aminobutyric acid is preferable.

[0465] Specific examples of the vitamins include at least one kind of vitamin selected from the group consisting of vitamin A, β -carotene, vitamin B₁, vitamin B₂, vitamin B₆, vitamin C, vitamin E, vitamin K₁, vitamin K₂, niacin, inositol, folic acid, and biotin. Particularly, in a case where the algae belonging to Cyanidiophyceae are the present algae, specific examples of the vitamins include at least one kind of vitamin selected from the group consisting of vitamin A, β -carotene, vitamin C, vitamin E, vitamin K₁, vitamin K₂, and folic acid. As the vitamins, for example, at least one kind of vitamin selected from the group consisting of β -carotene, vitamin C, vitamin E, vitamin K₁, vitamin K₂, and folic acid is preferable.

[0466] In this step, one kind of nutrient or two or more kinds of nutrients may be separated. Furthermore, amino acids, fat-soluble vitamins (such as vitamin A, β -carotene, vitamin E, vitamin K₁, and vitamin K₂), water-soluble vitamins (such as vitamin B₁, vitamin B₂, vitamin B₆, vitamin C, niacin, inositol, folic acid, and biotin), and the like may be separated by type.

[0467] The method for separating the nutrient from the cell disruption product is not particularly limited, and may be appropriately selected according to the type of the nutrient. As the nutrient separation method, the methods generally used for the separation, purification, and the like for biochemical substances can be used in combination as appropriate. Examples of the separation method include, but are not limited to, centrifugation, washing, salting out, dialysis, recrystallization, reprecipitation, solvent extraction, adsorption, concentration, filtration, gel filtration, ultrafiltration, various types of chromatography (such as thin layer chromatography, column chromatography, ion exchange chromatography, high performance liquid chromatography, and adsorption chromatography), and the like.

(Optional Step)

[0468] The producing method of the present embodiment may include other steps in addition to the steps (a) and (b). Examples of those other steps include a step of culturing algae belonging to Cyanidiophyceae (culturing step), a step of collecting the algae belonging to Cyanidiophyceae from a culture solution (collecting step), a step of washing cells of the algae belonging to Cyanidiophyceae (washing step), a step of treating the algae belonging to Cyanidiophyceae at a low temperature (low-temperature treatment step), a step of drying the algae belonging to Cyanidiophyceae (drying step), a step of freezing the algae belonging to Cyanidiophyceae (freezing step), and the like. These steps can be performed before the step (a).

[0469] The culturing step can be performed by the method described above in “[Alga belonging to Cyanidiophyceae]” or “[Alga having a diploid cell form and a haploid cell form]”. Furthermore, the collecting step can be performed by a known method such as filtration or centrifugation. The washing step can be performed by suspending the cells in a washing solution (such as a buffer) at a pH of 1.0 to 6.0, and then collecting the cells from the washing solution by a method such as filtration or centrifugation.

[0470] The low-temperature treatment step can be performed by treating the algae belonging to Cyanidiophyceae at a temperature of 0° C. to 5° C.

[0471] For example, in a case where the algal cells belonging to Cyanidiophyceae collected through the culturing step and the collecting step are placed in an environment at a temperature of 0° C. to 5° C., the algal cells can be killed. Therefore, even a transformant obtained not using self-cloning can be handled in an open system. The time of the low-temperature treatment is not particularly limited, and is, for example, 8 days or longer, and preferably 10 days or longer.

[0472] The drying step can be performed by drying the algae belonging to Cyanidiophyceae with a dryer such as a room temperature dryer, a low temperature dryer, or a freeze dryer. For example, in a case where the algal cells belonging to Cyanidiophyceae collected through the culturing step and the collecting step are dried with the dryer, the algal cells can be killed. Therefore, even a transformant obtained not using self-cloning can be handled in an open system.

[0473] The freezing step can be performed by freezing the algae belonging to Cyanidiophyceae in liquid nitrogen, a freezer, or the like. For example, in a case where the algal cells belonging to Cyanidiophyceae collected through the culturing step and the collecting step are placed in an environment with a temperature equal to or lower than -4° C. and preferably equal to or lower than -20° C., the algal cells can be killed. Therefore, even a transformant obtained not using self-cloning can be handled in an open system.

[0474] The time of the freezing treatment is not particularly limited, and is, for example, 10 minutes or longer and preferably 30 minutes or longer.

[0475] The nutrient produced by the producing method of the present embodiment can be used for various uses such as nutritional supplements, foods, feeds, pet foods, cosmetics, pharmaceutical products, and reagents.

[Other Aspects]

[0476] The algae belonging to Cyanidiophyceae are considered to be resistant to gastric acid because they are resistant to acid. On the other hand, among the algae belonging to Cyanidiophyceae, those having no rigid cell wall (for example, the YFU3 strain (haploid), the HKN1 strain (haploid), algae belonging to *Cyanidioschyzon* (for example, *Cyanidioschyzon merolae*), haploids of algae belonging to *Galdieria*, haploids of algae belonging to *Cyanidium*, and the like) undergo cell rupture in a neutral environment (pH of about 7 to 10). Therefore, these algae are considered to undergo cell rupture in the intestinal tract. Furthermore, for these algae, a transformation system has been established, and accordingly, any gene can be introduced into the algae. Therefore, by preparing a transformant into which a gene for generating an arbitrary drug has been introduced, the transformant can be used as a capsule of a gastric acid-resistant drug. Consequently, the present invention also provides a pharmaceutical composition containing algae belonging to Cyanidiophyceae into which a drug-generating gene has been introduced.

[0477] "Drug-generating gene" means a gene expressed in the cells of algae belonging to Cyanidiophyceae, in which a translation product of the gene generates a drug for treating or preventing diseases in human beings or non-human animals (such as mammals). The translation product of the

drug-generating gene may be a drug or may be an enzyme that catalyzes a drug synthesis reaction.

[0478] The drug is not particularly limited, and examples thereof include antigenic proteins of pathogenic microorganisms or pathogenic viruses. The pathogenic viruses are not particularly limited, and examples thereof include a rabies virus, a porcine circovirus, a bovine rotavirus, an influenza virus, an AIDS virus, and the like. The algae belonging to Cyanidiophyceae into which antigenic proteins of these pathogenic microorganisms or pathogenic viruses have been introduced can be used as vaccines against infectious diseases caused by the microorganisms or the viruses.

EXAMPLE

[0479] Hereinafter, the present invention will be described based on examples, but the present invention is not limited to the following examples.

[Example 1] Analysis of Components of Algae Belonging to Cyanidiophyceae

(Culturing of Algae Belonging to Cyanidiophyceae)

[0480] *Cyanidioschyzon merolae* 10D was cultured with aeration by using 5 L of a 2× Allen medium. The cells were cultured for 2 weeks at a culturing temperature of about 35° C. under continuous white light (500 μmol/m²s).

[0481] Table 1 shows the composition of the 2× Allen medium.

TABLE 1

2x Allen medium	
(NH ₄) ₂ SO ₄	2.6 g
KH ₂ PO ₄	0.54 g
MgSO ₄ •7H ₂ O	0.5 g
CaCl ₂ •2H ₂ O	0.15 g
FeCl ₃ •6H ₂ O	0.784 mg
MnCl ₂ •4H ₂ O	0.144 mg
ZnSO ₄ •7H ₂ O	0.0888 mg
CoCl ₂ •6H ₂ O	0.016 mg
Na ₂ MoO ₄ •2H ₂ O	0.01 mg
Na ₂ EDTA•2H ₂ O	4 mg
H ₂ SO ₄	1 ml
Distilled water	999 ml

(Analysis of Components of Algae Belonging to Cyanidiophyceae)

[Amino Acids, Vitamins, and the Like]

[0482] The cells of *Cyanidioschyzon merolae* 10D cultured as described above were collected by centrifugation, and the component analysis thereof was deputed to Japan Food Research Laboratories. The analytical value per wet weight obtained by the component analysis was divided by 0.248 so as to be converted into a value per dry weight.

[0483] Tables 2 and 3 show the analysis method for each component.

TABLE 2

Component	Analysis method	Note
Isoleucine	Automatic amino acid analysis method	
Leucine	Automatic amino acid analysis method	
Lysine	Automatic amino acid analysis method	
Methionine	Automatic amino acid analysis method	Performic acid oxidation treatment, followed by hydrochloric acid hydrolysis and measurement
Cystine	Automatic amino acid analysis method	Performic acid oxidation treatment, followed by hydrochloric acid hydrolysis and measurement
Phenylalanine	Automatic amino acid analysis method	
Tyrosine	Automatic amino acid analysis method	
Threonine	Automatic amino acid analysis method	
Tryptophan	High performance liquid chromatography method	
Valine	Automatic amino acid analysis method	
Arginine	Automatic amino acid analysis method	
Histidine	Automatic amino acid analysis method	
Alanine	Automatic amino acid analysis method	
Aspartic acid	Automatic amino acid analysis method	
Glutamic acid	Automatic amino acid analysis method	
Glycine	Automatic amino acid analysis method	
Proline	Automatic amino acid analysis method	
Serine	Automatic amino acid analysis method	
γ-Aminobutyric acid	Automatic amino acid analysis method	

TABLE 3

Component	Analysis method	Note
Vitamin A (retinol activity equivalent)		Each of α-carotene (24 µg) and β-carotene (12 µg) was regarded as retinol activity equivalent (1 µg).
α-carotene	High performance liquid chromatography	
β-carotene	High performance liquid chromatography	
Vitamin B ₁	High performance liquid chromatography	
Vitamin B ₂	High performance liquid chromatography	
Vitamin B ₆	microbiological assay	Used strain: <i>Saccharomyces cerevisiae</i> (<i>S. uvarum</i>) ATCC 9080
Vitamin B ₁₂	microbiological assay	Used strain: <i>Lactobacillus delbrueckii</i> subsp. <i>lactis</i> (<i>L. leichmannii</i>) ATCC 7830
Vitamin C	High performance liquid chromatography	Made into derivative by using hydrazine, followed by measurement
Vitamin D	High performance liquid chromatography	
Vitamin E	High performance liquid chromatography	
Vitamin K ₁	High performance liquid chromatography	
Vitamin K ₂	High performance liquid chromatography	
Niacin equivalent		The total amount of niacin (nicotinic acid equivalent) and 1/60 tryptophan were regarded as niacin equivalent.

TABLE 3-continued

Component	Analysis method	Note
Niacin (nicotinic acid equivalent) tryptophan	microbiological assay High performance liquid chromatography	Used strain: <i>Lactobacillus plantarum</i> ATCC 8014
Inositol	microbiological assay	Used strain: <i>Saccharomyces cerevisiae</i> (<i>S. uvarum</i>) ATCC 9080
Folic acid	microbiological assay	Used strain: <i>Lactobacillus rhamnosus</i> (<i>L. casei</i>) ATCC 7469
Pantothenic acid	microbiological assay	Used strain: <i>Lactobacillus plantarum</i> ATCC 8014
Biotin	microbiological assay	Used strain: <i>Lactobacillus plantarum</i> ATCC 8014
Choline	Reinecke salt precipitation	
Taurine	Automatic amino acid analysis method	
Protein	Kjeldahl method	
Lipid	Acidolysis	
Dietary fiber	Enzyme-gravimetric method	

[0484] Tables 4 to 6 show the results of the component analysis. Table 4 shows the analysis results of amino acids, Table 5 shows the analysis results of vitamins, and Table 6 shows the analysis results of other nutrients. In Tables 4 and 5, analytical values published for *Chlorella*, *Euglena*, and *Spirulina* are also shown for reference. In Tables 4 to 6, *Cyanidioschyzon merolae* 10D is described as "Schyzon".

TABLE 4

Amino acids	Unit (per dry weight of 100 g)	Chlorella ¹⁾ Euglena ²⁾ Spirulina ³⁾ Spirulina ⁴⁾ Schyzon				
Isoleucine	g	2.06	1.57	3.17	3.209	3.34
Leucine	g	4.45	3.3	5.02	4.947	6.4
Lysine	g	3.07	2.83	2.7	3.025	3.66
Methionine	g	1.24	0.87	2.19	1.149	2.01
Cystine	g	0.63	0.66		0.662	0.89
Phenylalanine	g	2.59	1.79	5	2.777	3.5
Tyrosine	g	1.79	1.73		2.584	3.94
Threonine	g	2.53	2	2.78	2.97	3.58
Tryptophan	g	1.09	0.7	0.84	0.929	1.21
Valine	g	3.11	2.57	3.48	3.512	4.19
Arginine	g	3.51	4.19	3.6	4.147	6.04
Histidine	g	1.05	1.06	1.09	1.085	1.25
Alanine	g	4.27	3.03	4.11	4.515	5.96
Aspartic acid	g	4.82	3.83	5.47	5.793	7.04
Glutamic acid	g	6.01	4.9	8.02	8.386	7.77
Glycine	g	3.01	2.03	2.85	3.099	3.38
Proline	g	2.48	2.45	2.04	2.382	2.82
Serine	g	2.15	1.71	2.74	2.998	3.54
Total amount ⁵⁾	g	49.86	41.22	55.1	58.169	70.52
γ-Aminobutyric acid	g	—	—	—	—	0.197

¹⁾Information posted on the website of SUN CHLORELLA CORP.²⁾Physiology and Biochemistry of Euglena, Seizaburo Kitaoka, Gakkai Shuppan Center, 1989³⁾Information posted on the website of DIC LIFETEC Co., Ltd.⁴⁾USDA Food Composition Databases⁵⁾Total amount of 18 kinds of amino acids of isoleucine to serine

TABLE 5

Vitamins		Unit (per dry weight of 100 g)	Chlorella ¹⁾	Euglena ²⁾	Spirulina ³⁾	Spirulina ⁴⁾	Schyzon
Vitamin A	mg	—	—	16.2	—	17.75	
α-carotene	mg	17.4	—	—	—	0	
β-carotene	mg	—	7.03	—	—	212.88	
Vitamin B ₁	mg	1.55	18.1	3.78	2.38	3.98	
Vitamin B ₂	mg	4.73	6.06	3.99	3.67	4.06	
Vitamin B ₆	mg	1.58	1.36	0.84	0.364	1.52	
Vitamin B ₁₂	μg	310	190	—	0	—	
Vitamin C	mg	8	5	—	10.1	56.34	
Vitamin D	μg	794	—	—	0	0	
Vitamin E	mg	3.7	23.8	9.1	5	162.98	
Vitamin K ₁	μg	980	—	1080	25.5	4708	
Vitamin K ₂	μg	0	—	80	—	5674	
Niacin equivalent	mg	40.2	—	38.5	12.82	40.64	
Inositol	mg	274	—	105	—	237	
Folic acid	μg	1200	990	170	94	5232	
Pantothenic acid	mg	3.04	—	1.3	3.48	0	
Biotin	μg	213	—	30.4	—	49.5	
Choline	mg	274	—	—	66	0	

¹⁾⁻⁴⁾Same as in Table 4

TABLE 6

Nutrient	Unit (per dry weight of 100 g)	Schyzon
Taurine	g	0
Protein	g	76.056
Lipid	g	12.5
Dietary fiber	g	2.01

[0485] As shown in Table 4, it has been revealed that the total amino acid content is higher in the *Cyanidioschyzon merolae* 10D than in other algae conventionally used in foods and the like. Furthermore, the content of each of the amino acids tended to be higher in Schyzon than in other algae.

[0486] In addition, it has been confirmed that the *Cyanidioschyzon merolae* contains γ-aminobutyric acid. γ-Aminobutyric acid is a substance that functions as an inhibitory neurotransmitter, and has been found to have a brain function improving effect, a blood pressure lowering action, a sedative action, and the like. Tomato is known as a food rich in γ-aminobutyric acid, and has been reported to contain γ-aminobutyric acid in an amount of 0.062 g/100 g based on wet weight (Overview on Wide-spreading Dietary Supplements for the Appropriate Use in Dialysis Patients [paper] 8 GABA, Clinical Dialysis Vol. 24. No. 13, December 2008). It can be said that the γ-aminobutyric acid content in Schyzon is by no means lower than that in tomatoes.

[0487] As shown in Table 5, the *Cyanidioschyzon merolae* tended to be richer in vitamins compared to other algae. Particularly, the content of β-carotene, vitamin C, vitamin E, vitamin K₁, vitamin K₂, and folic acid was higher in the *Cyanidioschyzon merolae* than in other algae.

[0488] Almond is known as a food rich in vitamin E, and has been reported to contain vitamin E in an amount of 30.3 mg/100 g based on wet weight (Standard Tables of Food Composition in Japanese, 2015). Furthermore, Natto is known as a food rich in vitamin K, and has been reported to contain vitamin K in an amount of 600 μg/100 g based on wet weight (Standard Tables of Food Composition in Japanese, 2015). It can be said that the content of vitamin E and vitamin K is higher in Schyzon than in these foods.

[0489] From the above results, it has been confirmed that the content of nutrients such as amino acids and vitamins is higher in the *Cyanidioschyzon merolae* than in other algae conventionally used in foods and the like.

[Example 2] Preparation of γ-Aminobutyric Acid-Producing Ability-Enhanced Strain

(Preparation of Fragment for Transformation)

[0490] Fragments for transformation were prepared by the method described in Fujiwara et al. (PLOS One. 2013 Sep. 5; 8 (9): e73608). Table 7 shows the primers used for preparing the fragments for transformation. Hereinafter, the primers used will be described using the Primer No. described in Table 7.

TABLE 7

No.	Primer	Sequence (5'→3')	SEQ ID NO:
1	URA5_3 (-897) F	GAACTGAGGGCGAACGCA	33
2	URA5_3 (+471) R	CCCTAGCAGCTGACTGTATC	34
3	D184 (773) pqeF	<u>CACCATCACCATCAC</u> GCGTGAGTCAGTTCACTTGAC	35
4	D184 (+1884) pqeR	<u>AAGCTCAGCTAATTACAGCT</u> TGCTGACCTTACCC	36
5	pQE_R	GTGATGGTGATGGTGATGGG	37
6	pQE_F	TAATTAGCTGAGCTTGGACT CCTG	38
7	D184 (+25) o250R	<u>TATACTTCTCGTCGCGTCA</u> CCCTCGGGACTTG	39
8	D184 (+28) btF	<u>GCAGGCAAAAGTGT</u> GAAACCGCTCAGCGACCA	40

TABLE 7-continued

No.	Primer	Sequence (5'→3')	SEQ ID NO:
9	O250 (-600) pqeF	<u>AAGCTCAGCTAATTACGACG</u> AGAACGTATAAGGAGTG	41
10	O250 (-1) gfpR	<u>GCCCTTGCTCACCATGGTCA</u> ACGAACGAAAGAAACACA	42
11	GFP(1)F	ATGGTGAGCAAGGGCGAG	43
12	GFP(717)R	CTTGTACAGCTCGTCCATGC	44
13	bt (+1) gfpF	<u>GACGAGCTGTACAAGTAAC</u> TAGCTATTTATCTGGTACAT ATCATTC	45
14	bt (+200) pqeR	<u>CACCATCACCATCACACACT</u> TTTTGCCCTGCACAAGT	46
15	O250 (-600) F	CGACGAGAACGTATAAGGAG TG	47
16	bt (+200) R	ACACTTTTGCCTGCACAAAG T	48
17	bt (+200) uraR	<u>TTCGCCCCCTCAGTTCACACT</u> TTTTGCCCTGCACAAGT	49
18	D184 (+28) uraF	<u>AGTCAGCTGTAGGGAAAC</u> CGCTCAGCGACCA	50
19	D184 (1270) F	ACACGAATCACACGGTGCTG	51
20	D184 (+1448) R	TTGCCGATAACGCAGAAGAG A	52
21	URA5.3 (1412) R	AGGTCGCTGATGCGGAA	53
22	URA5.3_ (+4) F	<u>GAATACGTTGAATGATT CCT</u> AATGGGCAGAAGCAAG	54
23	GsURA5.3F	<u>CCGCATCAGCGACCTTGT CG</u> GAACACTCCGCC	55
24	GsURA5.3R	TCATTCAACGTATTCTCAA GTCGTTG	56

<Preparation of EGFP/URA_{Cm-Cm} Fragment>

[0491] In order to prepare a URA_{Cm-Cm} selection marker, a DNA fragment containing the URA5.3 gene (including an 897 bp upstream sequence and a 471 downstream sequence) was amplified by PCR by using the genomic DNA of the *Cyanidioschyzon merolae* 10D as a template and using primer set No. 1/2. A CMD184C fragment (including the last 1,961 bp and 1.9 kb downstream sequences of CDCl84C ORF) and a pQE80 vector (QIAGEN) were amplified by PCR by using primer set No. 3/4, and primer set No. 5/6 respectively. The CMD184C fragment was subcloned into the pQE80 vector by using an In-Fusion HD Cloning Kit (Clontech Laboratories, Inc.). The resulting pD184 vector was amplified by PCR by using primer set No. 7/8. The upstream sequence of CMO250C (-600 to -1; SEQ ID NO: 9), EGFP ORF (Takara Bio Inc.), and the downstream sequence of β-tubulin (+1 to +200) were amplified by PCR by using primer set No. 9/10, primer set no. 11/12, and

primer set No. 13/14 respectively, and subcloned into pQE by using the In-Fusion HD Cloning Kit. By using the resulting pQ250-EGFP vector as a template DNA, an assembly fragment (upstream region of CMO250C [-600 to -1], EGFP ORF, and downstream region of β-tubulin [+1 to +200]) was amplified by PCR by using primer set No. 15/16, and subcloned into the amplified pD184 vector. The resulting pD184-O250-EGFP vector was amplified by PCR by using primer set No. 17/18. The URA_{Cm-Cm} selection marker was subcloned into the amplified pD184-O250-EGFP vector. The resulting pD184-O250-EGFP-URA_{Cm-Cm} vector was used as a template for amplifying an EGFP/URA_{Cm-Cm} fragment by using primer set No. 19/20. The amplified DNA fragment (EGFP/URA_{Cm-Cm} fragment) was used for transformation. The EGFP/URA_{Cm-Cm} fragment was used as a negative control. FIG. 1(A) shows the construct of the EGFP/URA_{Cm-Cm} fragment.

<Preparation of GAD/URA_{Cm-Cm} Fragment>

[0492] A pD184-O250-GAD-URA_{Cm-Cm} vector was prepared by the same method as the method used in “<Preparation of EGFP/URA_{Cm-Cm} Fragment>” described above, except that instead of EGFP ORF, a DNA fragment was used which was obtained by adding a 3×HA tag to the 3' terminal of CMF072C (glutamate decarboxylase: GAD: SEQ ID NO: 3) of *Cyanidioschyzon merolae*. By using a pD184-O250-GAD-URA_{Cm-Cm} vector as a template and using primer set No. 19/20, a GAD/URA_{Cm-Cm} fragment was amplified by PCR. The amplified DNA fragment (GAD/URA_{Cm-Cm} fragment) was used for transformation. FIG. 1(B) shows the construct of the GAD/URA_{Cm-Cm} fragment.

<Preparation of GAD/URA_{Cm-Gs} Fragment>

[0493] In order to remove a sequence including an OMP-decarboxylase domain of URA_{Cm-Cm} from the pD184-O250-GAD-URA_{Cm-Cm} vector, the vector was amplified by PCR by using primer set No. 21/22. A sequence including an OMP-decarboxylase domain of URA5.3 of *Galdieria sulphuraria* was amplified by PCR by using primer set No. 23/24 and using a URA_{Cm-Gs} selection marker as template DNA, and the sequence was subcloned into the pD184-O250-GAD-URA_{Cm-Cm} vector. The resulting pD184-O250-EGAD-URA_{Cm-Gs} vector was used as a template for amplifying a GAD/URA_{Cm-Gs} fragment by using primer set No. 19/20 and PrimeSTAR (registered trademark) MAX (Takara Bio Inc.). The amplified DNA fragment (GAD/URA_{Cm-Gs} fragment) was used for transformation. FIG. 1(C) shows the construct of the GAD/URA_{Cm-Gs} fragment.

(Transformation)

[0494] As a parent strain for transformation, a uracil auxotrophic mutant, *Cyanidioschyzon merolae* M4 (Minoda et al., Plant Cell Physiol. 2004 June; 45 (6): 667-71.) was used. The cells were allowed to grow in MA2 medium containing uracil (0.5 mg/mL) and 5-fluoroorotic acid (0.8 mg/mL) (Ohnuma M et al. Plant Cell Physiol. 2008 January; 49 (1): 117-20.). The cells were cultured by being shaken at 130 rpm under the conditions of continuous white light (80 μmol/m²s) and 40° C. The MA2 medium was coagulated using 0.5% (w/v) gellan gum (Wako Pure Chemical Industries, Ltd.). In order to adjust the pH to 2.3, sulfuric acid was added thereto such that the final concentration became 0.05% (v/v). The cells were transformed by the method

described in Ohnuma M et al (Plant Cell Physiol. 2008 January; 49 (1): 117-20.) and Imamura S et al (Plant Cell Physiol. 2010 May; 51 (5): 707-17). For transformation, 4 µg of each of the fragments for transformation prepared as above was used. The cells were cultured at 40° C. under continuous white light until colonies were formed. Thereafter, the colonies were moved to starch on MA2 solid medium containing uracil (0.5 mg/mL).

(Immunoblotting)

[0495] A cell lysate (4 µg as protein amount) of each of the transformants obtained using each of the fragments for transformation was separated by SDS-PAGE (gel having a thickness of 1 mm, 10% acrylamide, 50 minutes at 25 mA). Then, the protein in the gel was transferred to a PVDF membrane (300 V, 400 mA, 75 minutes). Blocking was performed using 5% skim milk, and as a primary antibody, an anti-HA antibody (16B12, 1:2,000) was reacted for 1 hour. Thereafter, as a secondary antibody, an anti-mouse HRP-labeled antibody (1:25,000) was reacted for 1 hour. Chemiluminescent signals were detected using Immobilin Western Chemiluminescent HRP Substrate (Millipore) and LAS-3000 (FUJIFILM Corporation).

[0496] FIG. 2 shows the results. In FIG. 2, the right figure shows the gel stained after SDS-PAGE. The left figure shows the results of immunoblotting. Chemiluminescent signals were detected from the GAD/URA_{Cm-Cm} fragment transformant and the GAD/URA_{Cm-Gs} fragment transformant. On the other hand, no signal was detected from those transformed using the EGFP/URA_{Cm-Cm} fragment. This result shows that HA-tagged GAD is expressed in the GAD/URA_{Cm-Cm} fragment transformant and the GAD/URA_{Cm-Gs} fragment transformant. The expression amount of the HA-tagged GAD was larger in the GAD/URA_{Cm-Gs} fragment transformant than in the GAD/URA_{Cm-Cm} fragment transformant.

[0497] FIG. 3 shows the results of comparison between the GAD/URA_{Cm-Cm} fragment transformant and a wild-type strain (WT). As well as FIG. 2, the expression of the HA-tagged GAD was confirmed in the GAD/URA_{Cm-Cm} fragment transformant, but no signal was detected in the wild-type strain.

(Checking Number of GAD-Introduced Copies in Transformant)

[0498] Real-time PCR was performed using the primers shown in Table 8. It has been confirmed that a wild-type strain of *Cyanidioschyzon merolae* has one copy of GAD (CMF072C). Therefore, in the EGFP/URA_{Cm-Cm} fragment transformant used as a negative control, a value determined by number of copies amplified using the primer set in A of Table 8 (A value)/number of copies amplified using the primer set in B of Table 8 (B value) is regarded as 1, and the number of copies of GAD (CMF072C) in other transformants was checked.

[0499] As a result, it has been confirmed that one copy of GAD is introduced into the GAD/URA_{Cm-Cm} fragment transformant by transformation and the transformant has two copies of GAD including endogenous GAD. On the other hand, it has been confirmed that 10 copies of GAD is introduced into the GAD/URA_{Cm-Gs} fragment transformant by transformation and the transformant has 11 copies of GAD including the endogenous GAD. FIG. 4 shows the

results of checking the number of copies in the GAD/URA_{Cm-Gs} fragment transformant.

TABLE 8

Type of primer	Primer	Sequence (5'→3')	SEQ ID NO:
A	Primers for amplifying a portion of GAD gene	TGGTGCGGGTCGGTTTA CCGAATATCCTGGACCAGCAT (1359)R	57 58
B	Primers for amplifying a genomic DNA sequence irrelevant to GDA gene (internal standard)	CMD184 (452)F CMD184 (513)R	CGCTGATCAACCTGGGACTT GTCAAAGCCAAGCTCGATGAG 59 60

(Measurement of Intracellular γ-Aminobutyric Acid Content in Transformant)

[0500] The EGFP/URA_{Cm-Cm} fragment transformant, the GAD/URA_{Cm-Cm} fragment transformant, and the GAD/URA_{Cm-Gs} fragment transformant were allowed to grow in MA2 medium. The cells were cultured under the conditions of continuous white light and 40° C.

[0501] After being cultured, the cells were collected by centrifugation, and a soluble matter was extracted from the collected cells by using 75% ethanol. γ-Aminobutyric acid in the extracted soluble matter was quantified by OPA-post column derivatization by using HPLC.

[0502] The results are shown in Table 9. Table 9 shows a relative concentration determined in a case where the concentration of γ-aminobutyric acid (per dry weight of cells) in the EGFP/URA_{Cm-Cm} fragment transformant is regarded as 1.0.

TABLE 9

Transformant	Concentration of γ-aminobutyric acid (relative concentration)
EGFP/URA _{Cm-Cm} fragment transformant	1.0
GAD/URA _{Cm-Cm} fragment transformant	4.04
GAD/URA _{Cm-Gs} fragment transformant	12.29

[0503] As shown in Table 9, in the GAD/URA_{Cm-Cm} fragment transformant and the GAD/URA_{Cm-Gs} fragment transformant, the concentration of γ-aminobutyric acid was higher than in the EGFP/URA_{Cm-Cm} fragment transformant as a negative control. Furthermore, in the GAD/URA_{Cm-Gs} fragment transformant, the concentration of γ-aminobutyric acid was higher than in the GAD/URA_{Cm-Cm} fragment transformant. This result shows that the intracellular γ-aminobutyric acid concentration increases according to the number of introduced GAD copies.

[0504] From these results, it has been confirmed that in the *Cyanidioschyzon merolae*, it is possible to increase the intracellular concentration of a specific nutrient by transformation.

[Example 3] Cell Rupturing Treatment for Algae
Belonging to *Cyanidioschyzon*

(Drying and Swelling Treatment for Algal Cells)

[0505] The *Cyanidioschyzon merolae* 10D was cultured in the same manner as in Example 1, and 1 mL of the culture solution was subjected to centrifugation (1,500×g, 2 minutes). The supernatant obtained after the centrifugation was discarded, the precipitate of the algal cells was suspended in an isotonic solution (10% sucrose, 20 mM HEPES, pH 7.0) and subjected to centrifugation (1,500×g, 3 minutes), and the algal cells were washed. The supernatant obtained after the centrifugation was discarded, and the precipitate of the algal cells was left in a refrigerator (4° C.) for 3 days so as to dry the algal cells.

[0506] Three days later, the algal cells were suspended in 45 µL of an isotonic solution (10% sucrose, 20 mM HEPES, pH 7.0), and subjected to centrifugation (1,500×g, 3 minutes), and centrifugal supernatant and precipitate were collected.

[0507] In addition, as algae belonging to Cyanidiophyceae having a rigid cell wall, *Cyanidium caldarium* RK-1 was cultured and subjected to the same drying and swelling treatment as that described above, and centrifugal supernatant and precipitate were obtained.

(SDS-Polyacrylamide Electrophoresis (SDS-PAGE))

[0508] In order to prepare a sample for SDS-PAGE, 15 µL of a 4×SDS-PAGE sample buffer was added to the centrifugal supernatant. Furthermore, 60 µL of 1×SDS-PAGE sample buffer was added to the centrifugal precipitate. SDS-PAGE was performed on the supernatant sample and the precipitate sample prepared as above. The gel after SDS-PAGE was stained with coomassie brilliant blue, and proteins in the centrifugal supernatant and precipitate.

[0509] FIG. 5 shows the results. For the *Cyanidioschyzon merolae* 10D, a plurality of kinds of proteins were detected in both the supernatant and the precipitate obtained after the drying and swelling treatment. From this result, it has been confirmed that the cells of *Cyanidioschyzon merolae* 10D are ruptured by the drying and swelling treatment. On the other hand, for the *Cyanidium Caldarium* RK-1, no protein was detected in the supernatant obtained after the drying and swelling treatment. This result shows that the cells of *Cyanidium Caldarium* RK-1 are not ruptured by the drying and swelling treatment.

[Example 4] Isolation of Novel Microalgae

(Isolation of Novel Microalgae)

[0510] Hot acidic water was collected from a hot spring in Yufu-shi, Oita prefecture, Japan. Likewise, hot acidic water was collected from a hot spring in Hakone-machi, Ashigashimo-gun, Kanagawa prefecture, Japan. The collected hot acidic water was cultured in M-Allen medium (40° C., pH 2.0, 100 µmol/m²s), and microalgae were isolated. The YFU3 strain was isolated from the hot acidic water collected in Yufu-shi, and the HKN1 strain was isolated from the hot acidic water collected in Hakone-machi.

[0511] Table 10 shows the composition of M-Allen medium (hereinafter, referred to as "MA medium" as well).

TABLE 10

M-Allen Medium	
(NH ₄) ₂ SO ₄	2.64 g
KH ₂ PO ₄	0.54 g
MgSO ₄ •7H ₂ O	0.5 g
CaCl ₂ •2H ₂ O	0.14 g
A2 trace element	2 mL
Distilled water	994 mL
Adjusted to pH 2.0 by using sulfuric acid	
Autoclaving followed by addition of 4 mL of A2 Fe stock	
A2 trace element	
H ₃ BO ₄	2.85 g
MnCl ₂ •4H ₂ O	1.8 g
ZnCl ₂	0.105 g
Na ₂ MoO ₄ •2H ₂ O	0.39 g
CoCl ₂ •6H ₂ O	40 mg
CuCl ₂ •2H ₂ O	43 mg
Distilled water	1000 mL
A2 Fe stock	
EDTA•2Na	7 g
FeCl ₃ •6H ₂ O	4 g
Distilled water	1000 mL
Sterilization using filter	

(Isolation of Haploid Algae)

[0512] The HKN1 strain isolated from the hot acidic water collected in Hakone-machi was cultured in M-Allen medium in a static state for about one month (40° C., 2% CO₂). After about one month of the culturing, the growth phase ended, and the cells entered a stationary phase. While the cells were being in the stationary phase, the culture solutions of the HKN1 strain were observed. As a result, small *Cyanidioschyzon merolae*-like cells were observed in all of the culture solutions. FIG. 6(A) is a picture (magnification: 600×) of a culture solution of the HKN1 strain in the stationary phase. In FIG. 6(A), the arrows indicate *Cyanidioschyzon merolae*-like cells.

[0513] Under an inverted microscope (CKX41; Olympus Corporation), by using a Pasteur pipette with a sharp tip, the *Cyanidioschyzon merolae*-like cells were isolated one by one, and the cells were cultured in a static state in 1 mL of a hot spring medium or a modified MA medium (see Example 7). As the hot spring medium, a medium was used which was obtained by adding 10 mM (NH₄)₂SO₄ to hot spring water collected from Tsukahara hot spring (pH 1.15) or Tamagawa hot spring (pH 1.14) (Hirooka S and Miyagishima S Y., Front Microbiol. 2016 Dec. 20; 7:2022.). After growing in the hot spring medium, the cells having entered the stationary phase were maintained in M-Allen medium or MA2 medium (Ohnuma M et al. Plant Cell Physiol. 2008 January; 49 (1): 117-20.). FIG. 6(B) is a photograph (magnification: 600×) of *Cyanidioschyzon merolae*-like cells isolated from the culture solution of the HKN1 strain in the stationary phase. The *Cyanidioschyzon merolae*-like cells maintained their shape even though the cells were continuously cultured thereafter.

[0514] In order to confirm the relationship between the *Cyanidioschyzon merolae*-like cell isolated as above and the HKN1 strain, the sequence of one genomic region of these cells was analyzed using MiSeq (Illumina, Inc.). As a result, it has been confirmed that all of the *Cyanidioschyzon merolae*-like cells isolated from the culture solution of the HKN1 strain in the stationary phase (Schyzon-like cells

derived from HKN1 strain) have only one allele sequence for the analyzed region and therefore the *Cyanidioschyzon merolae*-like cells are haploid cells. On the other hand, it has been confirmed that the HKN1 strain has two allele sequences for the analyzed region and therefore the HKN1 strain is diploid cells. Similarly, it has been confirmed that the allele sequence of the Schyzon-like cells derived from the HKN1 strain is a sequence in which two allele sequences of HKN1 are crossed. FIG. 7 shows two allele sequences of the HKN1 strain (HKN1_allele 1: SEQ ID NO: 12, HKN1_allele 2: SEQ ID NO: 13) and an allele sequence of the Schyzon-like cell derived from the HKN1 (HKN1_Schyzon-like_allele 1: SEQ ID NO: 14). In FIG. 7, the arrowheads indicate bases in which polymorphism is observed between alleles.

[0515] From these results, it has been confirmed that the Schyzon-like cells derived from the HKN1 strain are haploid cells generated by the meiosis of the diploid HKN1 strain.

[0516] In addition, the YFU3 strain isolated from the hot acidic water collected in Yufu-shi was a *Cyanidioschyzon*-like cell, and a haploid cell.

[0517] Hereinafter, diploid cells of the YFU3 strain will be described as "YFU strain (diploid)", haploid Schyzon-like cells derived from the YFU3 strain will be described as "YFU3 strain (haploid)", and the YFU strain (diploid) and the YFU3 strain (haploid) will be collectively described as "YFU3 strain" in some cases. Furthermore, diploid cells of the HKN1 strain will be described as "HKN1 strain (diploid)", haploid Schyzon-like cells derived from the HKN1 strain will be described as "HKN1 strain (haploid)", and the HKN1 strain (diploid) and the HKN1 strain (haploid) will be collectively described as "HKN1 strain" in some cases.

[0518] FIG. 8 shows pictures (magnification: 600 \times) of the YFU3 strain (haploid), the HKN1 strain (haploid), and other cells of *Cyanidium* observed using a microscope (BX51; Olympus Corporation). FIG. 8(A) is a micrograph of *Galdieria sulphuraria* 074, FIG. 8(B) is a micrograph of *Cyanidium caldarium* RK-1, FIG. 8(C) is a micrograph of *Cyanidioschyzon merolae* 10D, FIG. 8(D) is a micrograph of the YFU3 strain (haploid), and FIG. 8(E) is a micrograph of the HKN1 strain (haploid). The scale bar in FIG. 3 represents 5 μ m. The *Galdieria sulphuraria* 074 (FIG. 8A) and the *Cyanidium caldarium* RK-1 (FIG. 8B) have a rigid cell wall and grow by forming endospores within the mother cell wall. In contrast, it has been confirmed that the YFU3 strain (haploid) (FIG. 8D) and the HKN1 strain (haploid) (FIG. 8E) do not have a rigid cell wall just as the *Cyanidioschyzon merolae* 10D (FIG. 8C) and grow by two-cell division. Furthermore, both the YFU3 strain (haploid) and the HKN1 strain (haploid) had a cell size of about 2 μ m.

(Comparison of Size of Ribosomal DNA ITS1)

[0519] The YFU3 strain (haploid), the HKN1 strain (haploid), and the *Cyanidioschyzon merolae* 10D were compared with each other in terms of the size of the ribosomal DNA ITS1.

[0520] DNA was extracted from the cells of the YFU3 strain, the HKN1 strain, and the *Cyanidioschyzon merolae* 10D, and the ribosomal DNA ITS1 was amplified by PCR. The amplified ITS1 fragment was subjected to agarose gel electrophoresis, and the sizes of ITS1 of the microalgae were compared with each other. The primers used for the amplification of ITS1 are as follows.

Forward primer:

(SEQ ID NO: 15)

TAGAGGAAGGAGAACGCTAA

Reverse primer:

(SEQ ID NO: 16)

TTGGCGTCAAAGACTCGATGATT

[0521] FIG. 9 shows the results of the agarose gel electrophoresis. As shown in FIG. 9, both the YFU3 strain (haploid) and the HKN1 strain (haploid) had an ITS1 size of about 1.5 kb. Furthermore, both the YFU3 strain (haploid) and the HKN1 strain (haploid), had an ITS1 size larger than that of the *Cyanidioschyzon merolae* 10D.

(Phylogenetic Analysis Based on rbcL Gene Sequence)

[0522] By using DNA extracted from cells of the YFU3 strain (haploid) and the HKN1 strain (haploid) as templates, the rbcL gene was amplified by PCR, and the sequence thereof was analyzed. The base sequences of the rbcL genes of the YFU3 strain (haploid) and the HKN1 strain (haploid) are shown in SEQ ID NOS: 1 and 2 respectively.

[0523] The sequences of the primers used for the amplification of rbcL are as follows.

Forward primer:

(SEQ ID NO: 17)

aaaactttccaaggrrccwgc

Reverse primer:

(SEQ ID NO: 18)

gcwggttgtgttchacwaaatc

[0524] Based on the base sequence of the rbcL gene of the YFU3 strain (haploid) and the HKN1 strain (haploid), molecular phylogenetic analysis was performed by the maximum likelihood method. FIG. 10 shows a molecular phylogenetic tree based on the base sequence of the rbcL gene.

[0525] From the results of the molecular phylogenetic analysis, it has been determined that the YFU3 strain and the HKN1 strain belong to *Cyanidium*.

(Culturing of YFU3 Strain (Haploid) and HKN1 Strain (Haploid) in Acidic Hot Spring Drainage Medium)

[0526] Whether the YFU3 strain (haploid) and the HKN1 strain (haploid) can be cultured in an acidic hot spring drainage medium was checked. As the acidic hot spring drainage, water from the Tsukahara hot spring or the Tamagawa hot spring was used. By adding 10 mM (NH₄)₂SO₄ as a nitrogen source to the acidic hot spring drainage from the Tsukahara hot spring or the Tamagawa hot spring, an acidic hot spring drainage medium was prepared. Alternatively, by adding 10 mM (NH₄)₂SO₄ as a nitrogen source and 2 mM KH₂PO₄ as a phosphorus source to the acidic hot spring drainage, an acidic hot spring drainage medium was prepared. The YFU3 strain (haploid) or the HKN1 strain (haploid) was cultured with aeration by using 50 mL of the acidic hot spring drainage medium. The cells were cultured for 2 weeks at a culturing temperature of 40° C. under continuous white light (100 μ mol/m²s).

[0527] As a result, the YFU3 strain (haploid) and the HKN1 strain (haploid) could be cultured in any of the acidic hot spring drainage media.

[Example 5] Analysis of Nutrient of Novel Microalgae

[0528] By using 5 L of M-Allen medium, the YFU3 strain (haploid) was cultured with aeration. The cells were cultured for 2 weeks at a culturing temperature of 35° C. under continuous white light (500 $\mu\text{mol}/\text{m}^2\text{s}$).

[0529] The cells of the YFU3 strain (haploid) cultured as described above were collected by centrifugation, and the component analysis thereof was deputed to Japan Food Research Laboratories. The analytical value per wet weight obtained by the component analysis was divided by 0.248 so as to be converted into a value per dry weight.

[0530] Each of the components was analyzed by the method shown in Tables 2 and 3 described above.

[0531] Tables 11 and 12 show the results of the component analysis. Table 11 shows the analysis results for amino acids, and Table 12 shows the analysis results for vitamins. In Tables 11 and 12, analytical values published for *Chlorella*, *Euglena*, and *Spirulina* are also shown for reference.

and the like. Furthermore, the content of each of the amino acids tended to be higher in the YFU3 strain (haploid) than in other algae.

[0533] In addition, it has been confirmed that the YFU3 strain (haploid) contains γ -aminobutyric acid. γ -Aminobutyric acid is a substance that functions as an inhibitory neurotransmitter, and has been found to have a brain function improving effect, a blood pressure lowering action, a sedative action, and the like. Tomato is known as a food rich in γ -aminobutyric acid, and has been reported to contain γ -aminobutyric acid in an amount of 0.062 g/100 g based on wet weight (Overview on Wide-spreading Dietary Supplements for the Appropriate Use in Dialysis Patients [paper] 8 GABA, Clinical Dialysis Vol. 24. No. 13, December 2008). It can be said that the γ -aminobutyric acid content in the YFU strain (haploid) is by no means lower than that in tomatoes.

[0534] As shown in Table 12, the YFU3 strain (haploid) tended to be richer in vitamins compared to other algae.

TABLE 11

Type of amino acid	Unit (per dry weight of 100 g)	Chlorella ¹⁾	Euglena ²⁾	Spirulina ³⁾	Spirulina ⁴⁾	YFU3 strain
Isoleucine	g	2.06	1.57	3.17	3.209	2.58
Leucine	g	4.45	3.3	5.02	4.947	5.40
Lysine	g	3.07	2.83	2.7	3.025	3.47
Methionine	g	1.24	0.87	2.19	1.149	1.65
Cystine	g	0.63	0.66		0.662	0.81
Phenylalanine	g	2.59	1.79	5	2.777	2.78
Tyrosine	g	1.79	1.73		2.584	3.06
Threonine	g	2.53	2	2.78	2.97	3.02
Tryptophan	g	1.09	0.7	0.84	0.929	1.05
Valine	g	3.11	2.57	3.48	3.512	3.59
Arginine	g	3.51	4.19	3.6	4.147	5.48
Histidine	g	1.05	1.06	1.09	1.085	1.17
Alanine	g	4.27	3.03	4.11	4.515	4.80
Aspartic acid	g	4.82	3.83	5.47	5.793	5.81
Glutamic acid	g	6.01	4.9	8.02	8.386	6.73
Glycine	g	3.01	2.03	2.85	3.099	3.02
Proline	g	2.48	2.45	2.04	2.382	2.50
Serine	g	2.15	1.71	2.74	2.998	3.15
Total amount ⁵⁾	g	49.86	41.22	55.1	58.169	60.08
γ -Aminobutyric acid	g	—	—	—	—	0.14

¹⁾Information posted on the website of SUN CHLORELLA CORP.

²⁾Physiology and Biochemistry of Euglena, Seizaburo Kitao, Gakkai Shuppan Center, 1989

³⁾Information posted on the website of DIC LIFETEC Co., Ltd.

⁴⁾USDA Food Composition Databases

⁵⁾Total amount of 18 kinds of amino acids of isoleucine to serine

TABLE 12

Vitamins	Unit (per dry weight of 100 g)	Chlorella ¹⁾	Euglena ²⁾	Spirulina ³⁾	Spirulina ⁴⁾	Schyzon
Vitamin A	mg	—	—	16.2	—	13.1
α -carotene	mg	17.4	—	—	—	0
β -carotene	mg	—	7.03	—	—	156.8
Vitamin C	mg	8	5	—	10.1	36
Vitamin E	mg	3.7	23.8	9.1	5	110
Vitamin K ₁	μg	980	—	1080	25.5	6350
Vitamin K ₂	μg	0	—	80	—	1407
Folic acid	μg	1200	990	170	94	2338

¹⁾⁻⁴⁾Same as in Table 4

[0532] As shown in Table 11, it has been revealed that the total amino acid content is higher in the YFU3 strain (haploid) than in other algae conventionally used in foods

Particularly, the content of β -carotene, vitamin C, vitamin E, vitamin K₁, vitamin K₂, and folic acid was higher in the YFU3 strain (haploid) than in other algae.

[0535] Almond is known as a food rich in vitamin E, and has been reported to contain vitamin E in an amount of 30.3 mg/100 g based on wet weight (Standard Tables of Food Composition in Japanese, 2015). Furthermore, Natto is known as a food rich in vitamin K, and has been reported to contain vitamin K in an amount of 600 µg/100 g based on wet weight (Standard Tables of Food Composition in Japanese, 2015). It can be said that the content of vitamin E and vitamin K in the YFU strain is higher than those in the foods described above.

[0536] From the above results, it has been confirmed that the content of nutrients such as amino acids or vitamins is higher in the YFU3 strain (haploid) than in other algae conventionally used in foods and the like.

[Example 6] Transformation of Novel Microalgae (YFU3 Strain)

(Preparation of Fragment for Transformation)

[0537] Fragments for transformation were prepared by the method described in Fujiwara et al. (Front Plant Sci. 2017 Mar. 14; 8:343). Table 13 shows the primers used for preparing the fragments for transformation. Hereinafter, the primers used will be described using Primer No. described in Table 13.

TABLE 13

No.	Primer	Sequence (5'→3')	SEQ ID NO:
1	APCC(1) Fapc	cttcgttgcgttggaccATGTTCG TTCAGACCACTTCTTT	19
2	APCC(180) R	ATCATTGCAACGCCAGATG	20
3	B-TUB (+1)	TAAACTAGCTATTTATCTGGTA CATATCATTATCAT	21
4	B-TUB (+200) Rd184	tgcgtgagcggtttcACACCTTTT GCCTGCACAAGTTTCGTAC	22
5	CAT(1) Fapcc	ggcggttgcgaatgtatGAACCTTT AATAAAATTGATTTAGACAATTGG	23
6	CAT(650) taaRbt	taaatagctagttaTAAAAGCCA GTCATTAGGCCTATC	24
7	APCC(-1) R	GGTCAACGAAACGAAAGAACACAGA GAACAAAGATAT	25
8	D184(+28) F	GAAACCGCTCAGCGACCAAGCGAC	26
9	D184(1270) F	ACACGAATCACACGGTGCTG	27
10	D184(+1448) R	TTGCCGATAACGCAGAAGAGA	28
11	APCC(-600) F	CGACGAGAACGTATAAGGAGTGCG CAGC	29
12	B-TUB (+200) R	ACACTTTTGCTGCACAAGTTT CGTAGC	30

The lower cases represent an adapter sequence of In-Fusion reaction.

<Preparation of CAT Vector for Transforming of *Cyanidioschyzon merolae*>

[0538] By using the genomic DNA of the *Cyanidioschyzon merolae* 10D as a template and using primer set No. 1/2,

1 to 180 nucleotides encoding a chloroplast transition sequence (60 amino acids) in APCC ORF (CMO250C) of *Cyanidioschyzon merolae* were amplified by PCR. Furthermore, by using the genomic DNA of the *Cyanidioschyzon merolae* 10D as a template and using primer set No. 3/4, a downstream nucleotide (200 bp, β13') of β-tubulin ORF of the *Cyanidioschyzon merolae* was amplified by PCR. By using a chloramphenicol acetyltransferase gene of pC194 (pC194, Gene ID: 4594904; *Staphylococcus aureus*) as a template and using primer set No. 5/6, CAT ORF was amplified by PCR. By using an InFusion Cloning Kit (Takara Bio Inc.), the sequence encoding a chloroplast transition sequence of APCC, CAT ORF, and the downstream sequence of β-tubulin were cloned into pD184-O250-EGFP-URA_{Cm-Cm} (Fujiwara et al., PLOS One. 2013 Sep. 5; 8 (9): e73608) amplified using primer set No. 7/8, thereby constructing a plasmid pD184-CAT. By using a plasmid pD18 as a template and using primer set No 9/10, a DNA fragment was amplified by PCR, thereby preparing a CAT vector for transforming *Cyanidioschyzon merolae*.

<Preparation of CAT Vector for Transforming Novel Microalgae>

[0539] By using the plasmid pD18 prepared as above as a template and using primer set No. 11/12, a DNA fragment was amplified by PCR, thereby preparing a CAT vector for transforming novel microalgae.

(Transformation)

[0540] The cells were transformed by a method obtained by modifying the method described in Ohnuma M et al (Plant Cell Physiol. 2008 January; 49 (1): 117-20.). The *Cyanidioschyzon merolae* 10D and the YFU3 strain (haploid) were used as parent strains for transformation. The cells of each of the parent strains were diluted with 50 mL of an MA2U medium (MA2 medium containing 0.5 mg/mL uracil) so as to yield a concentration of OD750=0.3. The cells were cultured for 19 hours with aeration (600 mL/ambient air/min) under continuous light (100 µmol/m²s). Then, Tween-20 was added to the culture solution such that the final concentration became 0.002%, and the cells were collected by centrifugation (2,000 g, 5 minutes). The collected cells were suspended in 270 µL of an MA2U medium. Transformation was performed according to the protocol using polyethylene glycol (PEG).

[0541] PEG4000 (Sigma-Aldrich Co. LLC., #81240, 0.6 g) was dissolved in 450 µL of an MA2U medium (95°C., 10 minutes), thereby preparing a 60% (w/v) PEG4000 solution. Subsequently, the PEG 4000 solution was kept at 42°C. on a heat block until used.

[0542] Each of the CAT vectors (4 µg) for transformation prepared as above was dissolved in 90 µL of water. The vector solution (90 µL), 10 µL of a 10×TF solution (400 mM (NH₄)₂SO₄, 40 mM MgSO₄, 0.3% H₂SO₄), and 100 µL of the PEG 4000 solution were mixed together in a 1.5 mL tube by means of pipetting. Then, 25 µL of the cell suspension was added to 200 µL of a TF-CAT vector-PEG4000 mixed solution, stirred by being flipped upside down three to four times, and immediately moved to 40 mL of an MA2U medium. The cells were cultured for 1 day with aeration (300 mL ambient air/min) under continuous light (100 µmol/m²s). Thereafter, the cells were collected by centrifugation (1,500 g, 5 minutes) and suspended in 2 mL of an

MA2U medium. The cells were cultured for 2 to 3 days in a 24-well plate (TPP Techno Plastic Products AG) under the conditions of continuous light, 42°C., and 5% CO₂. Subsequently, chloramphenicol (CP) was added to the culture solution, the cells were cultured for 10 days, and CP-resistant transformants were selected. The CP resistant transformants were diluted stepwise by being washed with a CP-free MA2U medium, and spotted on a starch bed on an MA2U gellan gum plate. The starch bed and the MA2U gellan gum plate were prepared by a method obtained by adding slight modifications (Fujiwara et al., PLOS One. 2013 Sep. 5; 8 (9): e73608) to the method described in Imamura S et al (Plant Cell Physiol. 2010 May; 51 (5): 707-17). The plate was cultured for 2 weeks in a 5% CO₂ incubator until colonies appeared. According to the method described in Fujiwara et al. (PLOS One. 2013 Sep. 5; 8 (9): e73608), the colonies were moved to a starch bed on a new MA2U medium plate.

(Evaluation of Chloramphenicol (CP) Resistance)

[0543] After being transformed as described above, the cells were cultured in an MA2 liquid medium supplemented with CP so as to select CP-resistant transformants of *Cyanidioschyzon merolae* 10D and the YFU3 strain (haploid). In order to evaluate CP-resistant concentration of each of the CP-resistant transformants, a wild-type strain (WT) and the transformant (TF) were cultured for 10 days in an MA2 liquid media containing CP at various concentrations (0.50, 100, 150, 200, and 250 µg/mL).

[0544] The results are shown in FIG. 11. In FIG. 11, 10D_TF and 10D_WT represent the CP-resistant transformant of the *Cyanidioschyzon merolae* 10D and the wild-type strain respectively. YFU3_TF and YFU3_WT represent the CP-resistant transformant of the YFU3 strain (haploid) and the wild-type strain respectively. As shown in FIG. 11, in both the *Cyanidioschyzon merolae* 10D and the YFU3 strain (haploid), the CP-resistant transformant exhibited higher CP concentration resistance compared to the wild-type strain.

[0545] In addition, in order to check whether the CAT gene has been introduced into the cells cultured as shown in FIG. 11, primers were designed for the 5' terminal sequence of the chloroplast transition sequence of APCC and for the 3' terminal sequence of CAT ORF. The sequences of the primers are as follows.

APCC (1) forward primer:	(SEQ ID NO: 31)
ATGTTCGTTCAGACCAGTTCTTT	
CAT (650) reverse primer:	(SEQ ID NO: 32)
TAAAAGCCAGTCATTAGGCCTA	

[0546] In FIG. 11, CP-resistant transformant cells (TF_0, TF_150) of the *Cyanidioschyzon merolae* 10D cultured in CP at 0 µg/mL or 150 µg/mL and the CP-resistant transformant cells (TF_0, TF_100) of the YFU3 strain (haploid) cultured in CP at 0 µg/mL or 100 µg/mL were collected and subjected to PCR by using the above primer set. Then, agarose gel electrophoresis was performed to check where or not amplified fragments generated by PCR are present.

[0547] The results are shown in FIG. 12. In FIG. 12, "w/o TF" represents a cell that has not been subjected to transformation (wild-type strain). As shown in FIG. 12, in the CP-resistant transformant of the YFU3 strain (haploid), even

though the cells were cultured in CP at a concentration of 0 µg/mL and at a concentration of 100 µg/mL, the amplified fragment of the CAT gene was checked. In the *Cyanidioschyzon merolae*, the amplified fragment of the CAT gene was checked in the cells cultured in CP at a concentration of 150 µg/mL, but no amplified fragment of the CAT gene was checked in the cells cultured in CP at a concentration of 0 µg/mL. It is considered that in a case where the *Cyanidioschyzon merolae* is cultured in the absence of CP, the introduced CAT gene may be eliminated.

[0548] The above results show that the YFU3 strain (haploid) can be transformed just as the *Cyanidioschyzon merolae*.

[Example 7] Transformation of Novel Microalgae (HKN1 Strain)

(Preparation of Fragment for Transformation)

[0549] Table 14 shows the primers used for preparing fragments for transformation. Hereinafter, the primers used will be described using Primer No. shown in Table 14.

TABLE 14

No.	Primer	Sequence (5' → 3')	SEQ ID NO:
1	URAS.3 (-2500) F	CGGTACCCGGGGATCGTCCTCTCGAAAAA TGATTAC	63
2	URAS.3 (-501) R	CGACTCTAGAGGATCTGTGAAGACTATGTCG GTGTT	64
3	APCC promoter	CGGTACCCGGGGATCCATTCCCCTATTCAT FGAATG	65
4	APCC promoter R	GACTCTTCGGCGTTATTCGCG	66
5	mVenus EcoRV F	AACGCCGAAAGAGTCgatatcATGGTTAGCA AGGGCGAAG	67
6	mVenus NotI R	CCATGAAATGCAAGTgcggccgcTCACTTAT ACAGTTCATCCATACCC	68
7	Btub 3'UTR F	ACTTGCATTTCATGGCGAGGC	69
8	Btub 3'UTR R	CGACTCTAGAGGATCTCAAGCTCGGAAGAGA AGCTTC	70
9	CPCC promoter	CGGTACCCGGGGATCTCACAGCATATGCAA FGAGCCTGC	71
10	CPCC promoter R	GTTTGCAGTGTTCGATCAGAGTTAG	72
11	Tp of POP F	GGAAACACTGCAAACATGCGGTTAGCGTAGG GTC	73
12	Tp of POP R	CTTGTAAAGTTCATAGCCGAACGGCGGGCG	74
13	CAT F	ATGAACCTTAACAAGATTGACCTGG	75
14	CAT R	AGCGCACATACAAACTCACAGGAGCCAGTCGT TGG	76
15	UBQ 3'UTR F	GTTTGTATGTGCGCTTCGTAAAGC	77

TABLE 14-continued

No.	Primer	Sequence (5'→3')	SEQ ID NO:
16	UBQ 3'UTR R	CGACTCTAGAGGATCTTGACCTAAGACCACA CACCTAG	78
17	pmVenus F	TGATTCAAGAAAGAACATTCCCTATTCATG	79
18	pmVenus R	TGACATATGCTGTGATCAAGCTCGGAAGAGAA GCTTC	80
19	pCAT F	TCACAGCATATGTCAAGAGCCTGC	81
20	pCAT R	ACTGTATTCGAATGTCCTGACCTAAGACCACA CACCTAG	82
21	URA5.3 (-1501) F	CATTCGAAATACAGTGTTCGTG	83
22	URA5.3 (-1502) R	TTTCCTTCTGAATCAAACGAGACG	84
23	M13 F	GTAAAACGACGGCCAGT	85
24	M13 R	CAGGAAACAGCTATGAC	86

<Preparation of CAT Vector for Transforming Novel Microalgae>

[0550] The construct for introducing genes into the upstream of the URA5.3 gene was prepared as below. By using the genomic DNA of the HKN1 strain as a template and using primer set No. 1/2, a DNA fragment of the upstream sequence (-2500 to -501) of the URA5.3 gene was amplified by PCR. Then, by using the In-Fusion HD Cloning Kit, the fragment amplified by PCR was subcloned into pUC19. The resulting vector will be referred to as pURA5.3up vector.

[0551] Subsequently, by using the genomic DNA of the HKN1 strain as a template and using primer set No. 3/4, a DNA fragment of the upstream sequence (-500 to -1) of APCC was amplified by PCR. Furthermore, by using primer set No. 5/6, m Venus ORF (synthesized by Integrated DNA Technologies) was amplified by PCR. In addition, by using the genomic DNA of the HKN1 strain as a template and using primer set No. 7/8, a DNA fragment of the downstream sequence (+1 to +250) of β -tubulin was amplified by PCR. By using the In-Fusion HD Cloning Kit, the three fragments amplified by PCR were subcloned into pUC19. The resulting vector will be referred to as pm Venus vector.

[0552] Thereafter, by using the genomic DNA of the HKN1 strain as a template and using primer set No. 9/10, a DNA fragment of the upstream sequence (-500 to -1) of CPCC was amplified by PCR. Furthermore, by using the genomic DNA of the *Cyanidioschyzon merolae* 10D as a template and using primer set No. 11/12, a DNA fragment of the organelle transition sequence of POP was amplified by PCR. In addition, by using primer set No. 13/14, CAT ORF (synthesized by Integrated DNA Technologies) was amplification by PCR. Moreover, by using the genomic DNA of the HKN1 strain as a template and using primer set No. 15/16, a DNA fragment of the downstream sequence (+1 to +250) of ubiquitin was amplified by PCR. By using the In-Fusion HD Cloning Kit, the four fragments amplified by

PCR were subcloned into pUC19. The resulting vector will be referred to as pCAT vector.

[0553] Subsequently, by using the pm Venus vector as a template and using primer set No. 17/18, a DNA fragment was amplified by PCR. Furthermore, by using the pCAT vector as a template and using primer set No. 19/20, a DNA fragment was amplified by PCR. In addition, by using primer set No. 21/22, the pURA5.3up vector was amplified by PCR. Then, by using the In-Fusion HD Cloning Kit, the amplified fragment of the pm Venus vector and the amplified fragment of the pCAT vector were cloned into the pURA5.3up vector amplified as described above. By using the resulting pURA5.3up-m Venus-CAT vector as a template and using primer set No. 23/24, a DNA fragment was amplified by PCR. The amplified DNA fragment was used as a CAT vector for transformation. The structure of the CAT vector for transformation is shown in FIG. 13 as PCR Product.

(Transformation)

[0554] The cells were transformed by a method obtained by modifying the method described in Ohnuma M et al (Plant Cell Physiol. 2008 January; 49 (1): 117-20.). The HKN1 strain (haploid) was used as a parent strain for transformation. The cells of the parent strain were diluted with 50 mL of an MA2U medium (MA2 medium containing 0.5 mg/mL uracil) so as to yield a concentration of OD750=0.3. The cells were cultured for 60 hours with aeration (600 mL ambient air/min) in a light-dark cycle (12L:12D) under light (50 μ mol/m²s) at a temperature of 42° C. Then, Tween-20 was added to the culture solution such that the final concentration became 0.002%, and the cells were collected by centrifugation (2,000 g, 5 minutes). The collected cells were suspended in 270 μ L of an MA2U medium. Transformation was performed according to the protocol using polyethylene glycol (PEG). PEG4000 (Sigma-Aldrich Co. LLC., #81240, 0.6 g) was dissolved in 450 μ L of an MA2U medium (95° C., 10 minutes), thereby preparing a 60% (w/v) PEG4000 solution. Subsequently, the PEG4000 solution was kept at 42° C. on a heat block until used.

[0555] The CAT vector (4 μ g) for transformation was dissolved in 90 μ L of water. The vector solution (90 μ L), 10 μ L of the 10xTF solution (400 mM (NH₄)₂SO₄, 40 mM MgSO₄, 0.3% H₂SO₄), and 100 μ L of the PEG4000 solution were mixed together in a 1.5 mL tube by means of pipetting, thereby preparing a TF-CAT vector-PEG 4000 mixed solution. Then, 25 μ L of the cell suspension was added to 200 μ L of the TF-CAT vector-PEG4000 mixed solution, stirred by being flipped upside down three to four times, and immediately moved to 10 mL of an MA2U medium. The cells were cultured for 2 days in a static state under continuous light (20 μ mol/m²s) at a temperature of 42° C. Thereafter, the cells were collected by centrifugation (1,500 g, 5 minutes) and suspended in 1 mL of the Tsukahara mineral spring medium or 1 mL of a modified MA medium. The cell suspension (100 μ L) was added to 1 mL of the Tsukahara mineral spring medium or the modified MA medium containing CP at 100 μ g/mL, and the cells were cultured for 7 days in a static state in a 24-well plate (TPP Techno Plastic Products AG) under the conditions of continuous light (20 μ mol/m²s), 42° C., and 3% CO₂. A green portion, which had become darker, was added to 1 mL of a new Tsukahara mineral spring medium or modified MA medium containing

CP at 100 g/mL, the cells were further cultured for 7 days in a static state, and CP-resistant transformants were selected. By using the Pasteur pipette with a sharp tip, the CP-resistant transformant cells were isolated one by one under an inverted microscope (CKX41; Olympus Corporation), and cultured in a static state in 1 mL of the Tsukahara mineral spring medium or the modified MA medium.

[0556] Table 15 shows the composition of the modified MA medium.

TABLE 15

Modified MA medium	
(NH ₄) ₂ SO ₄	2.64 g
KH ₂ PO ₄	0.54 g
MgSO ₄ •7H ₂ O	0.5 g
CaCl ₂ •2H ₂ O	0.14 g
A2 trace element	2 mL
Distilled water	998 mL
Adjusted to pH 1.2 by using sulfuric acid	
Autoclaving followed by addition of 1.39 g of	
FeSO ₄ •7H ₂ O	
A2 trace element	
H ₃ BO ₄	2.85 g
MnCl ₂ •4H ₂ O	1.8 g
ZnCl ₂	0.105 g
Na ₂ MoO ₄ •2H ₂ O	0.39 g
CoCl ₂ •6H ₂ O	40 mg
CuCl ₂ •2H ₂ O	43 mg
Distilled water	1,000 mL

(Checking Transformant)

[0557] Primers were designed to check whether the m Venus-CAT gene had been introduced into the target site in the cells of the CP-resistant transformant that had been isolated and cultured (see FIG. 13; the approximate location of the primers are indicated by the arrows in the lower figure (Genome) of FIG. 13). The sequences of the primers are as follows.

Forward primer:

CATTGCACAGCAATGAAAGCG

(SEQ ID NO: 87)

Reverse primer:

ATCGAAACTGCGTAGATAGTGTGCGG

(SEQ ID NO: 88)

[0558] The cells of the CP transformant were collected. By using the primer set described above, PCR was performed, and agarose gel electrophoresis was carried out.

[0559] The results are shown in FIG. 14. An amplified fragment was observed at about 2.5 kb in the wild-type strain (WT). In contrast, an amplified fragment was detected at about 5.5 kb in the CP-resistant transformant (TF). From this result, it has been confirmed that the m Venus-CAT gene has been inserted into the target site (see FIG. 13) in the transformant (TF). The above result shows that gene targeting can be performed on the HKN1 (haploid).

[0560] Furthermore, the CP-resistant transformant was observed using a fluorescence microscope so as to check whether green fluorescence from m Venus is observed. The results are shown in FIG. 15. In FIG. 15, the left figure (m Venus) is a fluorescence micrograph showing the detected fluorescence of m Venus, the middle figure (Chl) is a

fluorescence micrograph showing the detected autofluorescence of the chloroplast, and the right figure (merged) is an image obtained by merging the two fluorescence micrographs. As shown in FIG. 15, in the CP-resistant transformant, m Venus was expressed in the cytoplasm, and green fluorescence was detected. The above results show that the HKN1 (haploid) can express a foreign gene.

[Example 8] Cell Rupturing Treatment for YFU3 Strain and HKN1 Strain

(Drying and Swelling Treatment for Algal Cells)

[0561] The YFU3 strain (haploid), the HKN1 strain (haploid), and the HKN1 strain (diploid) were cultured in the same manner as in Example 5, and 1 mL of the culture solution was subjected to centrifugation (1,500×g, 2 minutes). The supernatant obtained after the centrifugation was discarded, the precipitate of the algal cells was suspended in an isotonic solution (10% sucrose, 20 mM HEPES, pH 7.0) and subjected to centrifugation (1,500×g, 3 minutes), and the algal cells were washed. The supernatant obtained after the centrifugation was discarded, and the precipitate of the algal cells was left in a refrigerator (4° C.) for 3 days so as to dry the algal cells.

[0562] Three days later, the algal cells were suspended in 45 μL of an isotonic solution (10% sucrose, 20 mM HEPES, pH 7.0), and subjected to centrifugation (1,500×g, 3 minutes), and centrifugal supernatant and precipitate were collected.

[0563] Furthermore, *Cyanidium caldarium* RK-1 as a control of algae having a rigid cell wall and *Cyanidioschyzon merolae* 10D as a control of algae having no rigid cell wall were cultured and subjected to the same drying and swelling treatment as that described above, and centrifugal supernatant and precipitate were obtained.

(SDS-Polyacrylamide Electrophoresis (SDS-PAGE))

[0564] In order to prepare a sample for SDS-PAGE, 15 μL of a 4×SDS-PAGE sample buffer was added to the centrifugal supernatant. Furthermore, 60 μL of 1×SDS-PAGE sample buffer was added to the centrifugal precipitate. SDS-PAGE was performed on the supernatant sample and the precipitate sample prepared as above. The gel after SDS-PAGE was stained with coomassie brilliant blue, and proteins in the centrifugal supernatant and precipitate were checked.

[0565] FIG. 16 shows the results. In the YFU3 strain (haploid), the HKN1 strain (haploid), and the *Cyanidioschyzon merolae* 10D, a plurality of kinds of proteins were detected in both the supernatant and the precipitate obtained after the drying and swelling treatment. From this result, it has been confirmed that the YFU3 strain (haploid), the HKN1 strain (haploid), and the *Cyanidioschyzon merolae* 10D undergo cell rupture by the drying and swelling treatment. On the other hand, in the HKN1 strain (diploid) and the *Cyanidium Caldarium* RK-1, no protein was detected in the supernatant obtained after the drying and swelling treatment. This result shows that the HKN1 strain (diploid) and the *Cyanidium Caldarium* RK-1 do not undergo cell rupture by the drying and swelling treatment.

[Example 9] Preparation of Diploid by Combining Haploids

(Combining HKN1 Strains (Haploids))

[0566] No. 1 and No. 5 strains of the HKN1 strain (haploid) were mixed together and cultured in an MA medium for 3 weeks under the culture conditions of 40° C., light at 50 $\mu\text{mol}/\text{m}^2/\text{s}$, a light-dark cycle (12L:12D), and 2% CO₂. After being cultured for 3 weeks, the cells were further subcultured (diluted 20x) in a new medium for 1 week. Thereafter, the algal cells were observed using a microscope. As a result, diploid-like cells having a rigid cell wall were checked (FIG. 17A).

[0567] The diploid-like cells were collected and stained with DAPI so as to quantify the fluorescence intensity. For comparison, the HKN1 strain (haploid) was also stained with DAPI. The results are shown in FIG. 17B.

[0568] As shown in FIG. 17B, the fluorescence intensity of the diploid-like cells was twice the fluorescence intensity of the haploid strain. From this result, it has been confirmed that the diploid-like cells are diploids. From the above results, it has been revealed that by mixing together and culturing haploid cells, diploid cells can be prepared.

(Combining YFU3 Strains (Haploids))

[0569] The YFU3 strain (haploid) was subjected to the same operation as that described above. As a result, the emergence of diploid cells was confirmed by an optical microscope.

[Example 10] Preparation of Haploid Cells of Algae Belonging to *Galdieria*

(Creation of Cells Having No Cell Wall: *G. sulphuraria*)

(Method (a))

[0570] The Tsukahara mineral spring medium (Hirooka et al. 2016 Front in Microbiology) or a modified MA medium (1 mL) was put into a 24-well plate, about 10 cells of *Galdieria sulphuraria* SAG108.79 were added to each of the wells, and the cells were cultured for 1 week under light at 50 $\mu\text{mol}/\text{m}^2/\text{s}$ at a temperature of 42° C. in 2% CO₂. In this medium, both the tadpole-like cells and the round cells were present. Under a microscope, the tadpole-like cells were isolated using a Pasteur pipette with a sharp tip, and these cells were further cultured in an MA medium under light at 50 $\mu\text{E}/\text{m}^2/\text{s}$ at 42° C. in 2% CO₂.

(Method (b)) (Three-Stage Culturing)

[0571] The *Galdieria sulphuraria* SAG108.79 was cultured in an MA medium until the strain reached a plateau phase, a cell population was taken out of the culture solution and diluted 100x, and the cells were further subcultured in an MA medium. After 3 days of culturing, tadpole-like cells appeared. Therefore, by using a Pasteur pipette with a sharp tip, the tadpole-like cells were isolated from the culture solution while being observed under a microscope. The isolated cells were further cultured in an MA medium at 42° C. under light at 50 $\mu\text{E}/\text{m}^2/\text{s}$ in 2% CO₂.

[0572] FIG. 18A shows the normal cells of *Galdieria sulphuraria* SAG108.79 (left figure) and the cells having no rigid cell wall checked after the culturing of the normal cells (right figure).

(Creation of Cells Having No Rigid Cell Wall: *G. partita*)

[0573] In the same manner as in (Method (a)) described above, haploids were created using *Galdieria partita* NBRC 102759 (obtained from NITE Biological Resource Center) instead of *Galdieria sulphuraria* SAG108.79. FIG. 18B shows the normal cells of the *Galdieria partita* NBRC 102759 (left figure) and the cells having no rigid cell wall checked after the culturing of the normal cells (right figure).

(Observation by Optical Microscope)

[0574] In any of the culturing processes described above, the cells having no rigid cell wall could be easily disrupted by freeze thawing or the like, and the cell contents could be extracted. The cells having no rigid cell wall that appeared during the culturing of the algae belonging to *Galdieria* included both the tadpole-like cells and round cells. The cells indicated by the arrows in the left figure of FIGS. 18A and 18B are considered to be mother cell walls thrown off after mitosis.

(Allele Analysis)

[0575] The cells having no rigid cell wall that appeared by the culturing of the *Galdieria sulphuraria* SAG 108.79 were collected, one region of the genomic DNA thereof was amplified by PCR, and the sequence thereof was analyzed by the Sanger method. For comparison, sequence analysis was also performed on the same region of the cells having a rigid cell wall. The results are shown in FIG. 19.

[0576] The cells having a rigid cell wall had two kinds of allele sequences (2N_allele1 (SEQ ID NO: 61) and 2N_allele2 (SEQ ID NO: 62)). In contrast, the cells having no rigid cell wall only had either 2N_allele1 or 2N_allele2. From this result, it has been confirmed that the cells having no rigid cell wall are haploids. Therefore, it has been revealed that the algae belonging to *Galdieria* also include diploid cells and haploid cells.

[0577] Hereinafter, the diploid and haploid of the *Galdieria sulphuraria* SAG108.79 will be described as SAG108.79 (diploid) and SAG108.79 (haploid) respectively. Furthermore, the diploid and haploid of the *Galdieria partita* NBRC 102759 will be described as NBRC 102759 strain (diploid) and NBRC 102759 (haploid) respectively.

[Example 11] Cell Rupturing Treatment for Algae Belonging to *Galdieria*

[0578] A cell rupturing treatment was performed in the same manner as in Example 8, except that instead of the YFU3 strain and the like, SAG108.79 (diploid), SAG108.79 (haploid), the NBRC 102759 strain (diploid), and NBRC #102759 (haploid) were used. Then, SDS-PAGE was performed in the same manner as in Example 8. The gel after SDS-PAGE was stained with coomassie brilliant blue, and proteins in the centrifugal supernatant and precipitate were checked.

[0579] The results are shown in FIG. 20. In SAG108.79 (haploid) and the NBRC 102759 strain (haploid), a plurality of kinds of proteins were detected in both the supernatant and the precipitate obtained after the drying and swelling treatment. From these results, it has been confirmed that SAG108.79 (haploid) and the NBRC 102759 strain (haploid) undergo cell rupture by the drying and swelling treatment. On the other hand, in SAG108.79 (diploid) and the NBRC 102759 strain (diploid), no protein was detected in

the supernatant obtained after the drying and swelling treatment. This result shows that the cells of SAG108.79 (diploid) and the NBRC 102759 strain (diploid) do not undergo cell rupture by the drying and swelling treatment.

[Example 12] Preparation of Vitamin-Producing Ability-Enhanced Strain

(Preparation of Transformant)

[0580] A pD184-O250-TC-URA_{Cm-Cm} vector was prepared in the same manner as in “<Preparation of EGFP/URA_{Cm-Cm} fragment>” in Example 2 described above, except that instead of EGFP ORF, a DNA fragment was used which was obtained by adding a 3×HA tag to the 3' terminal of *Cyanidioschyzon merolae* CML326C (Tocopherol cyclase: TC). Furthermore, the upstream sequence of CMP166C (-500 to -1), a fragment obtained by adding a 3×FLAG tag to the 3' terminal of CMN202C (homogentisate phytoltransferase: HPT), and the downstream sequence of CMK296C (+1 to +278) were amplified by PCR, and the resulting fragment was introduced into the downstream of a URA_{Cm-Cm} selection marker of pD184-O250-TC-URA_{Cm-Cm} by using the In-Fusion HD Cloning Kit, thereby preparing a pD184-O250-TC-URA_{Cm-Cm} vector. By using the pD184-O250-VE-URA_{Cm-Cm} vector as a template and using primer set No. 19/20 in Table 7, a VE/URA_{Cm-Cm} fragment was amplified by PCR. The amplified DNA fragment (VE/URA_{Cm-Cm} fragment) was used for transformation. FIG. 21 shows the constitution of the DNA fragment for transformation (VE/URA_{Cm-Cm} fragment). By using the DNA fragment (VE/URA_{Cm-Cm} fragment), *Cyanidioschyzon merolae* M4 which is a uracil auxotrophic mutant was transformed. The transformation and the culturing after the transformation were performed by the same method as that in Example 2. Then, by using an anti-HA antibody or an anti-FLAG antibody, immunoblotting was performed in the same manner as in Example 2.

[0581] The results are shown in FIG. 22. The left figure shows the results of immunoblotting using an anti-HA antibody, and the right figure shows the results of immunoblotting using an anti-FLAG antibody. In FIG. 22, “GFP” represents a cell (EGFP/URA_{Cm-Cm} fragment transformant) transformed using the EGFP/URA_{Cm-Cm} fragment (FIG. 1A). “VE” represents a cell transformed using the VE/URA_{Cm-Cm} fragment (VE/URA_{Cm-Cm} fragment transformant).

[0582] From the results in FIG. 22, it has been confirmed that the VE/URA_{Cm-Cm} fragment transformant expresses tocopherol cyclase and homogentisate phytoltransferase.

(Measurement of Intracellular Vitamin E Content in Transformant)

[0583] The EGFP/URA_{Cm-Cm} fragment transformant and the VE/URA_{Cm-Cm} fragment transformant were allowed to grow in an MA2 medium. The cells were cultured under the conditions of continuous white light and 40° C. After being cultured, the cells were collected by centrifugation, and a soluble matter was extracted from the collected cells by using 75% ethanol. Vitamin E in the extracted soluble matter was quantified. The Vitamin E quantification was depicted to Japan Food Research Laboratories and performed by high performance liquid chromatography method.

[0584] The results are shown in Table 16. Table 16 shows relative values determined under the assumption that the vitamin E concentration (per dry weight of cells) in the EGFP/URA_{Cm-Cm} fragment transformant is 1.0.

TABLE 16

Transformant	Concentration of Vitamin E (relative value)
EGFP/URA _{Cm-Cm} fragment transformant	1.0
VE/URA _{Cm-Cm} fragment transformant	1.61

[0585] As shown in Table 16, the vitamin E concentration was higher in the VE/URA_{Cm-Cm} fragment transformant than in the EGFP/URA_{Cm-Cm} fragment transformant as a negative control. From these results, it has been confirmed that the intracellular concentration of vitamin E in the *Cyanidioschyzon merolae* can be increased by transformation.

INDUSTRIAL APPLICABILITY

[0586] According to the present invention, there are provided novel microalgae which can be used industrially and use of the novel microalgae. The novel microalgae provided by the present invention can be used for a nutrient composition, a nutritional supplement, a composition for supplying nutrients, a method for producing a nutrient, and the like. Furthermore, the novel microalgae provided by the present invention can be used in various foods such as processed foods, functional foods, and dietary supplements, feeds, pet foods, cosmetics, and the like.

[0587] In addition, according to the present invention, there is provided a nutrient composition or a nutritional supplement which is rich in nutrients such as amino acids or vitamins. The nutrient composition or nutritional supplement can be used for foods such as various processed foods, functional foods, and dietary supplements, feeds, pet foods, cosmetics, and the like.

[0588] Moreover, according to the present invention, there is provided a method for producing a nutrient such as amino acids, vitamins, proteins, lipids, and dietary fiber.

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AVRAGDNFGQ RASDADQGRP VHQVGRQTEG ERFPAVRRRALR VFLKFTPRHT MLGSAVSICS 120
LSLMGSVSAQ QALGAATLPL WTQLFPVLLV GLVPALLMNII YIVGLNQLCD 1PVDRVNKPY 180
LPLASGEVSL PAAVSLVGCM LLGFSLGLFW LPOSTAALRF ALVASCLGTT LYSLPPIRLK 240
RFPLLASLVCI LVRGAVVNII GFYLHARSAV MSLRGWPWLAE LSRLGPWLAE LQFGERNIFR 300
LMDIPDAKG DNQHQLSSFT FCVTMLIFMF IAGGIFCMSS ALATVPRHRA 360
FAAGGFHFVA AAWLWRWSRA SMMEAHRSEV VYNFYMDIWK LFYLEYVVLP LLL 413

SEQ ID NO: 7      moltype = DNA length = 1701
FEATURE          Location/Qualifiers
source           1..1701
                 mol_type = other DNA
                 organism = Cyanidioschyzon merolae

SEQUENCE: 7
atgcgagaaa agctccagtgc tagagggttgc ggggtcgagg tccaaacggga gctccgaaacg 60
ggcaagggttgc cctgttaaacgc acggccagcg tgggtgtcgcc cgtcgatccc gctgtccctc 120
gttggatgttgc acgtcgatccgc gggtcgatccgc gggcactccc agagccacca 180
tggagaaaacgc ataggccacgc gggtcgatccgc ctgtcgatccgc caacgtatcc 240
ccacggaaacgc ggccgcgtgc agtttcacgc ggacgcggccgc ggacgcgttc gtcgttgcgttgc 300
gcgagccccc ttgcacagca tgcgtatccgc ggggttccgc gtcgttgcgttgc gtcgttgcgttgc 360
aagcgggttgc cagggtcgac ggcgttgcgttgc gacgttgcgttgc gtcgttgcgttgc 420
aaggagacccgc cagatccccc ggaacttccc caccgttgcgttgc ttcatgttgc gtcgttgcgttgc 480
cgtggatccgc tggggatccgc atcggttgcgttgc ccacagggatccgc gacttccgttgc 540
gcgttcatgttgc acgttgcgttgc aggttgcgttgc ttcgttgcgttgc ttcgttgcgttgc 600
gtgcaggacgc agtgcgttgc ttcgttgcgttgc atgttgcgttgc gtcgttgcgttgc gtcgttgcgttgc 660
cacctgcgttgc acgttgcgttgc ctttccatgttgc gtcgttgcgttgc gtcgttgcgttgc 720
gacgaaacgc caggacgcgc ggcgttgcgttgc gtcgttgcgttgc gtcgttgcgttgc 780
acgggatccgc acgttccgc ttcgttgcgttgc ttcgttgcgttgc ttcgttgcgttgc 840
cgagacgcgtt ccatgttgcgttgc gtcgttgcgttgc gtcgttgcgttgc gtcgttgcgttgc 900

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atcgagccgc tcttatacggt gggttcggat ggtgcgcgcg cgaactggaa ctggctcaacg 960
cgtttcaag tcttttgaaac aggctggcag atactctctg cttcgcgtcg atccacttgt 1020
atatttcggg actggcatgg tccgtttttt cgcttcgcgc gtgcacccac gtatactcgag 1080
aagaactggg gcaagtgcgtt cccatcgca tgggtctggc tgcagtgtca cgtcttgcac 1140
gtgtcccgag aggacgcatac atcgcgtggag ctgtcgcccg cagacatcga tgcaccgttg 1200
actttgcacct gtgtgggagc ggcgtcgagag ctctgcgtggc cccagcgacc gaataagggt 1260
gttagctcgaa aaaccattgg tatcatcgccg ttccactggaa agggatattt gtggggatgc 1320
gtctgcttggaa actgcgcgcg catatcctgg agtgtcccgat ggggcaactg gcagatggaa 1380
gcccacggca cgcgtatag cgtaaagggtc tccgctgaga ccgtgcgaacc agggggcgat 1440
gtactgggtc cgaccgcgtc cggcatcgag tttgtgttaa gagacgggtgc gcgtgtacg 1500
cttcgttgc acgtggaga cggacggaga aacgtgtgtg ttttggacgc actctgtcgc 1560
aacgctgtc ctgtcgagct tgggtggagc ctggcgccg accccaaatgg aacgagtgtgg 1620
acggcggagc gcgagccctt tccccccacccg gtaaaggcat tgtttttat tggcgaacgca 1680
cccgccggagc cggatcgctt c 1701

```

SEQ ID NO: 8 moltype = AA length = 567

FEATURE Location/Qualifiers
source 1..567
mol_type = protein
organism = Cyanidioschyzon merolae

SEQUENCE: 8

```

MEKQLQCRGS GFEVQRELRS GKVAVNAAAA WVPSPSSLSS VVYAYQEHA GRCRALPEPP 60
WRKHRHRARS PGTGPRAVLT GRRRSASSLK ASPVRQHVLS GGSRGSLRTL 120
KRVSGARRRR DERCHHIEAT KETADPPELP HAGVHMPSVD RGFFEGWYVR IVLPTTEATSL 180
AFMYSLEDGR RGRIQVLSSG VQDELLVSRE MMDGFFPALR HLPTSTLRIG QWGRTS DLLS 240
DERAEHARSL APSLFREHVH TGYQLSAQWH QGSIQADDVD RERSMVAHLS PDISCAFDMR 300
IEPLLSWGS GARSTGTWLT RFQVFEPGWQ ILSAFARSTG IFRDWHGRVF RFERAPTYIE 360
KNWGSAPPSR WFWLQCHVFD VLREDASSLE LSSADIDAPL TLTCVGARRE LCWPQRPNKV 420
VARETIGIIA PHWKGYLWEF AAWCRRRIW SVQWGNWQME AHGTRYSVKV SAETAEPGAY 480
VLGPTRHGMQ FVVRDGARGT LRLQLRDEQA NVVILDALCR NAAAVELGGE LASDPNGTSW 540
TAERGAFPPP VKALFLIGER PGRAVRV 567

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SEQ ID NO: 9 moltype = DNA length = 600

FEATURE Location/Qualifiers
source 1..600
mol_type = other DNA
organism = Cyanidioschyzon merolae

SEQUENCE: 9

```

cgacgagaac gtataaggag tgcgcacggc gttttgttac aataccgata gatgaggttc 60
gaacatcgca ttccacccat gaggggggc gcacgcgtcca gagagtggag atggaaaagt 120
gccagcggag ccctgaggat gcaaaaaagt acggacccgt gacggaaaggag caaatggaaa 180
ggagggcggaa acttcgaggg ctacttgcatt tagtaagtac aaccaacgtat ggttaagaaac 240
gtatggagg tgcgcacggc gacttttacg ctgcccataa tatcaggaga tttgtgggtgc 300
tggagacgg acctccagag tgaacaaggaa ggtactttt tggacaacact tctaagggtcg 360
aactatatac gataaaattt gaaagaatgt ttgtggccg gtccaaaaag acgggggggc 420
gtctgcacat cagttggaga cgttttgtcc aaggcgcgcg gategcgcact actgtacacg 480
gcccgcgaga acgtggcgag aatacgcga cggactcgccg ggctggcgtc gtcgcaccc 540
cgtcttttgc ctcacaactt cgcgatatct ttgtttctgt ttgtttttccg ttctgttgc 600

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SEQ ID NO: 10 moltype = DNA length = 599

FEATURE Location/Qualifiers
source 1..599
mol_type = other DNA
organism = Cyanidioschyzon merolae

SEQUENCE: 10

```

gcctcgaggc cctcttcgtcg tcgtcgccga agagctgaca gaaagcggtc caggtgcgcg 60
agccctcaactt gggcatatac cagtgtgtt gtcggggcac cactagtccg gtgaactcg 120
ccggctcgacc aaatcttcca cccggaaagg tccgcaccgtc actgcgcgca ggcgcgttcca 180
agtttctagg cgcgttgcggc gtgcaggatgc cgcgttccaaat tccggcaatgc ggcgtgcgcg 240
catggccgc catgtgtcg accccatccg tgcacagcgag gcacgtcgag cgatttgggg 300
cgccgtgcata gcccggcgca cgtatgtatc ttcggatcgatc tccgtacaca aacaggactt 360
ttctgaccat gataatccca ttccggatcgatc aegcatecgcc cacttcgcga acacccctgtt 420
gctggggcgca gcccggcgca gcacccgtcgcc agcacggccg cctcggcgcg ggcgcgcgc 480
tagcggatc tccgacgacg gacgtgggtc gagattcgat ttccggaggcc gcacgcgcga 540
aaaagggtcat tcgtgtttgt gtctgcggac tctgaacgtt cctcgtaaaacg cacttctgt 599

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SEQ ID NO: 11 moltype = DNA length = 500

FEATURE Location/Qualifiers
source 1..500
mol_type = other DNA
organism = Cyanidioschyzon merolae

SEQUENCE: 11

```

ccctcaactgg catggcaaaag ccgttctgtc tccgtgcgcg atctgcgttca ggggtgttaag 60
cgccgttgcg aatgtcaag gaagggttacg tgcacagtgaa atgcacgcga ataaccgtta 120
catccgaaag gaagtacaag taatggaaacc tgaaggtagg gtccagcgcg atgtggcg 180
cttccggaa tgtcaataacc gatctatcgcc ctcctatgtcc catagaacgcg 240

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cttggcata ggcgggagaa cggggcgctg ccccgcgctg cgctccatgg aacaatgctc 300
aatcacaat taaattgtac ttatataat ctgtatgtac tatgtatgtac aaaatagcat 360
tccaggatc ccgagatcac acacgcgc gagacgtgaa ctgtctcgactctcgct 420
acggcttcat ttccctcgta ttcttcgca attagatacg agtgcggtaa gattctggc 480
tgaagtttc aatattatgtg 500

SEQ ID NO: 12      moltype = DNA length = 100
FEATURE           Location/Qualifiers
source            1..100
mol_type = other DNA
organism = Cyanidium sp.

SEQUENCE: 12
attttgtcac ctgaacctat ggatggtaaa caatgggttc tggtttcacc aaatggatgg 60
atggAACCTC tcatggagcg gtgtctctcg taggttttgt 100

SEQ ID NO: 13      moltype = DNA length = 100
FEATURE           Location/Qualifiers
source            1..100
mol_type = other DNA
organism = Cyanidium sp.

SEQUENCE: 13
attttgtcac ctgaacctat ggatgataaa caatgggttc tggtttcacc aaatggatgg 60
atggTACCCCT tcagggagcg gtgtctctcg taggttttgt 100

SEQ ID NO: 14      moltype = DNA length = 100
FEATURE           Location/Qualifiers
source            1..100
mol_type = other DNA
organism = Cyanidium sp.

SEQUENCE: 14
attttgtcac ctgaacctat ggatggtaaa caatgggttc tggtttcacc aaatggatgg 60
atggTACCCCT tcagggagcg gtgtctctcg taggttttgt 100

SEQ ID NO: 15      moltype = DNA length = 21
FEATURE           Location/Qualifiers
source            1..21
mol_type = other DNA
organism = synthetic construct

SEQUENCE: 15
tagaggaagg agaagtgcgt a 21

SEQ ID NO: 16      moltype = DNA length = 24
FEATURE           Location/Qualifiers
source            1..24
mol_type = other DNA
organism = synthetic construct

SEQUENCE: 16
ttgcgttcaa agactcgatc attc 24

SEQ ID NO: 17      moltype = DNA length = 20
FEATURE           Location/Qualifiers
source            1..20
mol_type = other DNA
organism = synthetic construct

SEQUENCE: 17
aaaactttcc aaggrrccwgc 20

SEQ ID NO: 18      moltype = DNA length = 23
FEATURE           Location/Qualifiers
source            1..23
mol_type = other DNA
organism = synthetic construct

SEQUENCE: 18
gcwggttgt tytchacwaa atc 23

SEQ ID NO: 19      moltype = DNA length = 39
FEATURE           Location/Qualifiers
source            1..39
mol_type = other DNA
organism = synthetic construct

SEQUENCE: 19
tttcgttctgt tgaccatgtt cgttcagacc agtttcttt 39

SEQ ID NO: 20      moltype = DNA length = 20
FEATURE           Location/Qualifiers
source            1..20

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	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 20	
atcattcgca acgccagatg	20
SEQ ID NO: 21	moltype = DNA length = 34
FEATURE	Location/Qualifiers
source	1..34
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 21	
taaaactagct atttatctgg tacatatcat tcat	34
SEQ ID NO: 22	moltype = DNA length = 44
FEATURE	Location/Qualifiers
source	1..44
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 22	
tcgcgtgagcg gtttacact tttgcctgc acaagtttc gtac	44
SEQ ID NO: 23	moltype = DNA length = 48
FEATURE	Location/Qualifiers
source	1..48
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 23	
ggcggttgca atgatatgaa ctttaataaa attgatattt acaattgg	48
SEQ ID NO: 24	moltype = DNA length = 39
FEATURE	Location/Qualifiers
source	1..39
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 24	
taaatagcta gtttataaaa gccagtcatt aggccttac	39
SEQ ID NO: 25	moltype = DNA length = 36
FEATURE	Location/Qualifiers
source	1..36
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 25	
ggtcaacgaa cgaagaaaca cagagaacaa agatat	36
SEQ ID NO: 26	moltype = DNA length = 24
FEATURE	Location/Qualifiers
source	1..24
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 26	
gaaaccgctc agcgaccaag cgac	24
SEQ ID NO: 27	moltype = DNA length = 20
FEATURE	Location/Qualifiers
source	1..20
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 27	
acacgaatca cacggtgctg	20
SEQ ID NO: 28	moltype = DNA length = 21
FEATURE	Location/Qualifiers
source	1..21
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 28	
ttgccgataa cgcagaagag a	21
SEQ ID NO: 29	moltype = DNA length = 28
FEATURE	Location/Qualifiers
source	1..28
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 29	
cgacgagaac gtataaggag tgcgacacg	28

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SEQ ID NO: 30      moltype = DNA  length = 30
FEATURE
source
1..30
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 30
acacttttg cctgcacaag tttcgtacg                                30

SEQ ID NO: 31      moltype = DNA  length = 24
FEATURE
source
1..24
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 31
atgttcgttc agaccaggttt cttt                                24

SEQ ID NO: 32      moltype = DNA  length = 22
FEATURE
source
1..22
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 32
taaaaagccag tcattaggcc ta                                    22

SEQ ID NO: 33      moltype = DNA  length = 19
FEATURE
source
1..19
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 33
gaactgaggg gcgaacgca                                         19

SEQ ID NO: 34      moltype = DNA  length = 20
FEATURE
source
1..20
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 34
cccttagcagc tgactgtatc                                     20

SEQ ID NO: 35      moltype = DNA  length = 35
FEATURE
source
1..35
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 35
caccatcacc atcacgcgtg agtcagttca ctgac                           35

SEQ ID NO: 36      moltype = DNA  length = 34
FEATURE
source
1..34
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 36
aagctcagct aattacagct tgctgacctt accc                                34

SEQ ID NO: 37      moltype = DNA  length = 20
FEATURE
source
1..20
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 37
gtgtatggtga tgggtatggg                                     20

SEQ ID NO: 38      moltype = DNA  length = 24
FEATURE
source
1..24
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 38
taatttagctg agcttggact cctg                                         24

SEQ ID NO: 39      moltype = DNA  length = 33
FEATURE
source
1..33

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	mol_type = other DNA organism = synthetic construct	
SEQUENCE: 39	tatacggtt cgtcgctca ccctcgggac ttg	33
SEQ ID NO: 40	moltype = DNA length = 33 Location/Qualifiers	
FEATURE source	1..33 mol_type = other DNA organism = synthetic construct	
SEQUENCE: 40	gcaggcaaaa agtgtgaaac cgctcagcga cca	33
SEQ ID NO: 41	moltype = DNA length = 37 Location/Qualifiers	
FEATURE source	1..37 mol_type = other DNA organism = synthetic construct	
SEQUENCE: 41	aagctcagct aattacgacg agaacgtata aggagtg	37
SEQ ID NO: 42	moltype = DNA length = 37 Location/Qualifiers	
FEATURE source	1..37 mol_type = other DNA organism = synthetic construct	
SEQUENCE: 42	gcccttgctc accatggtca acgaacgaag aaacaca	37
SEQ ID NO: 43	moltype = DNA length = 18 Location/Qualifiers	
FEATURE source	1..18 mol_type = other DNA organism = synthetic construct	
SEQUENCE: 43	atgggtgagca agggcgag	18
SEQ ID NO: 44	moltype = DNA length = 20 Location/Qualifiers	
FEATURE source	1..20 mol_type = other DNA organism = synthetic construct	
SEQUENCE: 44	cttgtacagc tcgtccatgc	20
SEQ ID NO: 45	moltype = DNA length = 47 Location/Qualifiers	
FEATURE source	1..47 mol_type = other DNA organism = synthetic construct	
SEQUENCE: 45	gacgagctgt acaagtaaac tagtattta tctggtacat atcatc	47
SEQ ID NO: 46	moltype = DNA length = 36 Location/Qualifiers	
FEATURE source	1..36 mol_type = other DNA organism = synthetic construct	
SEQUENCE: 46	caccatcacc atcacacact ttttgctgc acaagt	36
SEQ ID NO: 47	moltype = DNA length = 22 Location/Qualifiers	
FEATURE source	1..22 mol_type = other DNA organism = synthetic construct	
SEQUENCE: 47	cgacgagaac gtataaggag tg	22
SEQ ID NO: 48	moltype = DNA length = 21 Location/Qualifiers	
FEATURE source	1..21 mol_type = other DNA organism = synthetic construct	
SEQUENCE: 48	acacttttg cctgcacaag t	21

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SEQ ID NO: 49      moltype = DNA  length = 36
FEATURE
source
1..36
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 49
ttcgccctc agttcacact tttgcctgc acaagt                         36

SEQ ID NO: 50      moltype = DNA  length = 32
FEATURE
source
1..32
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 50
gtcagctgct aggggaaacc gctcagcgcac ca                                32

SEQ ID NO: 51      moltype = DNA  length = 20
FEATURE
source
1..20
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 51
acacgaatca cacggtgctg                                         20

SEQ ID NO: 52      moltype = DNA  length = 21
FEATURE
source
1..21
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 52
ttgccgataa cgcagaagaa a                                         21

SEQ ID NO: 53      moltype = DNA  length = 17
FEATURE
source
1..17
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 53
aggtcgctga tgccggaa                                         17

SEQ ID NO: 54      moltype = DNA  length = 36
FEATURE
source
1..36
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 54
gaataacgttg aatgattcct aatgggcaga agcaag                         36

SEQ ID NO: 55      moltype = DNA  length = 32
FEATURE
source
1..32
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 55
ccgcacatcgc gaccttgcg gaacactccg cc                           32

SEQ ID NO: 56      moltype = DNA  length = 27
FEATURE
source
1..27
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 56
tcattcaacg tattttcaa gtcgttg                                     27

SEQ ID NO: 57      moltype = DNA  length = 18
FEATURE
source
1..18
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 57
tggtgccgggtt cggtttta                                         18

SEQ ID NO: 58      moltype = DNA  length = 21
FEATURE
source
1..21

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mol_type = other DNA
organism = synthetic construct
SEQUENCE: 58
ccgaatatcc tggaccagca t 21

SEQ ID NO: 59      moltype = DNA length = 20
FEATURE          Location/Qualifiers
source           1..20
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 59
cgctgatcaa cctgggactt 20

SEQ ID NO: 60      moltype = DNA length = 21
FEATURE          Location/Qualifiers
source           1..21
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 60
gtcaaaggcca agctcgatga g 21

SEQ ID NO: 61      moltype = DNA length = 100
FEATURE          Location/Qualifiers
source           1..100
mol_type = other DNA
organism = Galdieria sulphuraria
SEQUENCE: 61
gttgttaaac tcggattcgg atgcaagtac ttggaaattt accatgaaag atcctctcac 60
aaaagtggaa gttgtggtaa gaatagccag ttctccttgt 100

SEQ ID NO: 62      moltype = DNA length = 100
FEATURE          Location/Qualifiers
source           1..100
mol_type = other DNA
organism = Galdieria sulphuraria
SEQUENCE: 62
gttgttaaac tcggattcgg atgcaagtac ttggaaattt accatgaaag atcctctcac 60
aaaagtggaa gttgtggtaa gaaaagacag ttctcattgt 100

SEQ ID NO: 63      moltype = DNA length = 38
FEATURE          Location/Qualifiers
source           1..38
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 63
cggtaccctgg ggatcgtctc ctctcgaaaa atgattac 38

SEQ ID NO: 64      moltype = DNA length = 37
FEATURE          Location/Qualifiers
source           1..37
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 64
cgactctaga ggatctgtga agactatgtc ggtgttc 37

SEQ ID NO: 65      moltype = DNA length = 36
FEATURE          Location/Qualifiers
source           1..36
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 65
cggtaccctgg ggatccattc ccctatttca tgaatg 36

SEQ ID NO: 66      moltype = DNA length = 22
FEATURE          Location/Qualifiers
source           1..22
mol_type = other DNA
organism = synthetic construct
SEQUENCE: 66
gactcttcg gcgttatttg cg 22

SEQ ID NO: 67      moltype = DNA length = 40
FEATURE          Location/Qualifiers
source           1..40
mol_type = other DNA
organism = synthetic construct

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SEQUENCE: 67
aacggccaaa gagtcgatcatggtagcaagggcgaag                                40

SEQ ID NO: 68      moltype = DNA  length = 48
FEATURE          Location/Qualifiers
source           1..48
                 mol_type = other DNA
                 organism = synthetic construct

SEQUENCE: 68
ccatgaaatgcgatcgacttacatgtttcatccataacc                                48

SEQ ID NO: 69      moltype = DNA  length = 21
FEATURE          Location/Qualifiers
source           1..21
                 mol_type = other DNA
                 organism = synthetic construct

SEQUENCE: 69
acttgcatttcatggcgaggc                                              21

SEQ ID NO: 70      moltype = DNA  length = 37
FEATURE          Location/Qualifiers
source           1..37
                 mol_type = other DNA
                 organism = synthetic construct

SEQUENCE: 70
cgactctaga ggatctcaag ctcggaaagaaagtttc                                              37

SEQ ID NO: 71      moltype = DNA  length = 39
FEATURE          Location/Qualifiers
source           1..39
                 mol_type = other DNA
                 organism = synthetic construct

SEQUENCE: 71
cggttacccgg ggatctcaca gcatatgtca agagcctgc                                39

SEQ ID NO: 72      moltype = DNA  length = 27
FEATURE          Location/Qualifiers
source           1..27
                 mol_type = other DNA
                 organism = synthetic construct

SEQUENCE: 72
gttgcgttg tttccatgtca gagtttg                                27

SEQ ID NO: 73      moltype = DNA  length = 35
FEATURE          Location/Qualifiers
source           1..35
                 mol_type = other DNA
                 organism = synthetic construct

SEQUENCE: 73
ggaaaacatgc caaacatgcg gttaggcgtaaagggtc                                35

SEQ ID NO: 74      moltype = DNA  length = 31
FEATURE          Location/Qualifiers
source           1..31
                 mol_type = other DNA
                 organism = synthetic construct

SEQUENCE: 74
cttggtaaagg ttcatagccg aacggcgccc g                                              31

SEQ ID NO: 75      moltype = DNA  length = 25
FEATURE          Location/Qualifiers
source           1..25
                 mol_type = other DNA
                 organism = synthetic construct

SEQUENCE: 75
atgaacttta acaagattga cctgg                                25

SEQ ID NO: 76      moltype = DNA  length = 35
FEATURE          Location/Qualifiers
source           1..35
                 mol_type = other DNA
                 organism = synthetic construct

SEQUENCE: 76
agcgcacata caaactcaca ggagccagtc gttgg                                35

SEQ ID NO: 77      moltype = DNA  length = 24

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FEATURE	Location/Qualifiers
source	1..24
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 77	
gtttgtatgt ggcgttcgtt aagc	24
SEQ ID NO: 78	moltype = DNA length = 39
FEATURE	Location/Qualifiers
source	1..39
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 78	
cgactctaga ggatctctga cctaagagcca cacacctag	39
SEQ ID NO: 79	moltype = DNA length = 32
FEATURE	Location/Qualifiers
source	1..32
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 79	
tgattcagaa agaaacattc ccctatttca tg	32
SEQ ID NO: 80	moltype = DNA length = 37
FEATURE	Location/Qualifiers
source	1..37
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 80	
tgacatatgc tgtgatcaag ctccggaaagag aagcttc	37
SEQ ID NO: 81	moltype = DNA length = 24
FEATURE	Location/Qualifiers
source	1..24
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 81	
tcacagcata tgtcaagagc ctgc	24
SEQ ID NO: 82	moltype = DNA length = 39
FEATURE	Location/Qualifiers
source	1..39
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 82	
actgtatttc gaatgtctga cctaagagcca cacacctag	39
SEQ ID NO: 83	moltype = DNA length = 23
FEATURE	Location/Qualifiers
source	1..23
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 83	
cattcgaaat acagtgtttc gtg	23
SEQ ID NO: 84	moltype = DNA length = 24
FEATURE	Location/Qualifiers
source	1..24
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 84	
tttctttctg aatcaaacga gacg	24
SEQ ID NO: 85	moltype = DNA length = 17
FEATURE	Location/Qualifiers
source	1..17
	mol_type = other DNA
	organism = synthetic construct
SEQUENCE: 85	
gtaaaaacgac ggccagt	17
SEQ ID NO: 86	moltype = DNA length = 17
FEATURE	Location/Qualifiers
source	1..17
	mol_type = other DNA
	organism = synthetic construct

-continued

SEQUENCE: 86		
caggaaacag ctagac		17
SEQ ID NO: 87	moltype = DNA length = 22	
FEATURE	Location/Qualifiers	
source	1..22	
	mol_type = other DNA	
	organism = synthetic construct	
SEQUENCE: 87		
cattgcacag caatgaaaag cg		22
SEQ ID NO: 88	moltype = DNA length = 25	
FEATURE	Location/Qualifiers	
source	1..25	
	mol_type = other DNA	
	organism = synthetic construct	
SEQUENCE: 88		
atcgaaaactg cgtagatagt gtcgg		25

1. A method for producing a diploid alga, comprising:
 - (a) mixing two or more kinds of haploid cells of an alga belonging to Cyanidiophyceae having a diploid cell form and a haploid cell form, and culturing them; and
 - (b) isolating a diploid cell generated during the culturing.
2. The method for producing a diploid alga according to claim 1, wherein at least one of the two or more kinds of haploid cells is a transformant.
3. The method for producing a diploid alga according to claim 2, wherein the transformant is prepared by self-cloning.
4. The method for producing a diploid alga according to claim 1, wherein the culturing in the (a) is performed under a condition of temperature of 15° C. to 50° C., pH of 1.0 to 6.0, and CO₂ concentration of 1% to 3%.
5. The method for producing a diploid alga according to claim 1, wherein the alga is an alga belonging to *Galdieria* or *Cyanidium*.
6. A method for producing a nutrient, comprising: producing a diploid alga by the method or producing a diploid alga according to claim 1; and producing a nutrient composition containing the diploid alga or an extract thereof.
7. The method for producing a nutrient according to claim 6, wherein at least one of the two or more kinds of haploid cells is a transformant having an increased intracellular content of at least one kind of nutrient.
8. The method for producing a nutrient according to claim 7, wherein the transformant is prepared by self-cloning.
9. The method for producing a nutrient according to claim 6, wherein the nutrient composition comprises at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber.
10. A method for producing a food, comprising: producing a nutrient composition by the method for producing a nutrient according to claim 6; and producing a food comprising the nutrient composition.
11. The method for producing a food according to claim 10, wherein the food is a functional food or a dietary supplement.
12. A method for producing a feed or a pet food, comprising: producing a nutrient composition by the method for producing a nutrient according to claim 6; and producing a feed or a pet food comprising the nutrient composition.
13. A method for producing a cosmetic, comprising: producing a nutrient composition by the method for producing a nutrient according to claim 6; and producing a cosmetic comprising the nutrient composition.
14. A method for producing a nutrient, comprising: producing a diploid alga by the method or producing a diploid alga according to claim 1; disrupting cell of the diploid alga so as to obtain a cell disruption product; and separating at least one kind of nutrient from the cell disruption product.
15. The method for producing a nutrient according to claim 14, wherein the nutrient is at least one kind of nutrient selected from the group consisting of amino acids, vitamins, proteins, lipids, and dietary fiber.
16. The method for producing a nutrient according to claim 15, wherein the amino acids are at least one kind of amino acid selected from the group consisting of isoleucine, leucine, lysine, methionine, cystine, phenylalanine, tyrosine, threonine, tryptophan, valine, arginine, histidine, alanine, aspartic acid, glutamic acid, glycine, proline, serine, and γ-aminobutyric acid.
17. The method for producing a nutrient according to claim 15, wherein the vitamins are at least one kind of vitamin selected from the group consisting of vitamin A, β-carotene, vitamin B₁, vitamin B₂, vitamin B₆, vitamin C, vitamin E, vitamin K₁, vitamin K₂, niacin, inositol, folic acid, and biotin.
18. The method for producing a nutrient according to claim 15,

wherein at least one of the two or more kinds of haploid cells is a transformant having an increased intracellular content of at least one kind of nutrient.

- 19.** The method for producing a nutrient according to claim **18**, wherein the transformant is prepared by self-cloning.

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