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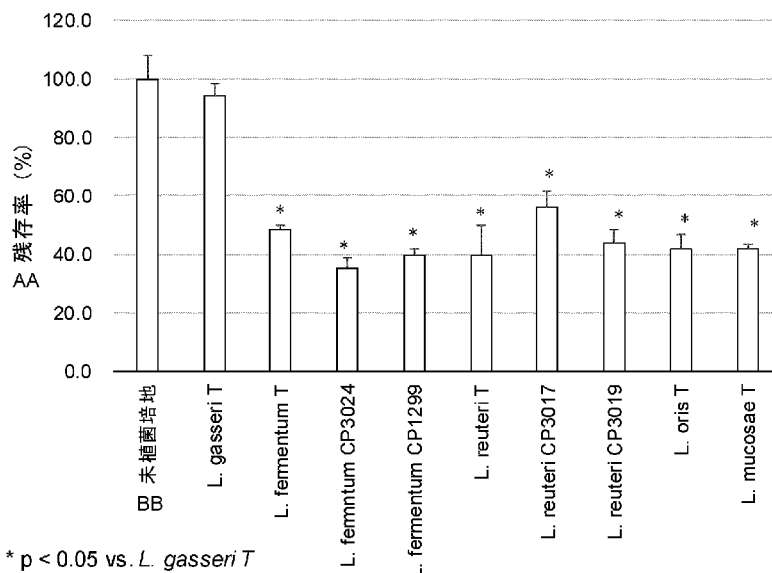
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(54) Title: METHOD FOR PRODUCING PLANT MILK-FERMENTED LIQUID

(54) 発明の名称: 植物性ミルク発酵液の製造方法



AA Survival rate (%)
BB Uninoculated medium

(57) Abstract: Provided is a method for producing a plant milk-fermented liquid having reduced plant smell. This method for producing a plant milk-fermented liquid comprises bringing, into contact with plant milk, a Lactobacillus strain including at least one selected from the group consisting of a Lactobacillus fermentum CP1299 strain, a Lactobacillus fermentum CP3024 strain, a Lactobacillus reuteri CP3017 strain, and a Lactobacillus reuteri CP3019 strain.



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- 国際調査報告 (条約第21条(3))
- 規則13の2に基づいて明細書とは別に提出された、寄託された生物材料に関する表示 (規則13の2.4(d)(i)及び48.2(a)(viii))

(57) 要約 : 植物臭が低減された植物性ミルク発酵液の製造方法が提供される。ラクトバチルス・ファーマンタムC P 1 2 9 9株、ラクトバチルス・ファーマンタムC P 3 0 2 4株、ラクトバチルス・ロイテリC P 3 0 1 7株及びラクトバチルス・ロイテリC P 3 0 1 9株からなる群から選択される少なくとも1種を含む乳酸菌株と、植物性ミルクとを接触させることを含む植物性ミルク発酵液の製造方法である。

DESCRIPTION

METHOD FOR PRODUCING PLANT-BASED MILK-FERMENTED LIQUID

TECHNICAL FIELD

5 [0001] The present invention relates to a method for producing a plant-based milk-fermented liquid.

BACKGROUND ART

[0002] Plant-based milk is a product prepared by size
10 reduction and liquefaction of a plant material such as rice, wheat, or soybeans (see, for example, J. Food Sci. Technol (September 2016) 53(9): 3408-3423). Examples of plant-based milk include rice milk, which is a food prepared by saccharifying rice with enzyme, malted
15 rice(koji), and/or the like, seasoning the saccharified rice by addition of a vegetable oil and/or the like, and then making the resulting product into the form of a milk.

[0003] In general, plant-based milk contains aldehydes as flavor components. Aldehydes may be favorable flavor
20 components in beverages having plant flavors. However, in beverages having fermented-milk-like flavors, aldehydes are known to be a cause of unpleasant odors due to unfavorable plant odors. Examples of methods of reducing the amount of aldehydes contained in plant-based milk include the
25 following methods.

[0004] For example, JP H9-205999 A describes a method in which the lactic acid bacterium *Leuconostoc mesenteroides* or the lactic acid bacterium *Lactobacillus brevis* is brought into contact with a food containing medium-chain aldehydes while preventing the growth of the lactic acid bacterium, to cause reduction of the medium-chain aldehydes to alcohols. Further, JP 2020-17 A describes a lactic acid fermented beverage obtained by subjecting a plant-based milk obtained using a rice material to lactic acid fermentation using the lactic acid bacterium *Lactobacillus sakei* strain No. 7 or the lactic acid bacterium *Lactobacillus sakei* strain No. 16, and describes that the beverage contained reduced amounts of diacetyl and hexanal.

15 SUMMARY OF THE INVENTION

PROBLEMS TO BE SOLVED BY THE INVENTION

[0005] An object of the present invention is to provide a method for producing a plant-based milk-fermented liquid with reduced plant odor.

20

MEANS FOR SOLVING PROBLEMS

[0006] The present invention provides a method for producing a plant-based milk-fermented liquid, the method including bringing a plant-based milk into contact with a lactic acid bacterial strain(s) including at least one

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selected from the group consisting of the *Lactobacillus fermentum* CP1299 strain, the *Lactobacillus fermentum* CP3024 strain, the *Lactobacillus reuteri* CP3017 strain, and the *Lactobacillus reuteri* CP3019 strain.

5 [0007] In an embodiment, the plant-based milk in the production method may be derived from a cereal. In an embodiment, the plant-based milk in the production method may be derived from rice. In an embodiment, the plant-based milk in the production method may be a saccharified
10 plant-based milk. In an embodiment, the contact between the lactic acid bacterial strain(s) and the plant-based milk in the production method may involve fermentation of the plant-based milk by the lactic acid bacterial strain(s).

15 [0008] The present invention provides a method for reducing the amount of aldehyde in a plant-based milk, the method including bringing a plant-based milk containing aldehyde into contact with a lactic acid bacterial strain(s) including at least one selected from the group
20 consisting of the *Lactobacillus fermentum* CP1299 strain, the *Lactobacillus fermentum* CP3024 strain, the *Lactobacillus reuteri* CP3017 strain, and the *Lactobacillus reuteri* CP3019 strain.

[0009] In an embodiment, the plant-based milk in the
25 reducing method may be derived from a cereal. In an

embodiment, the plant-based milk in the reducing method may be derived from rice. In an embodiment, the plant-based milk in the reducing method may be a saccharified plant-based milk. In an embodiment, the contact between the lactic acid bacterial strain(s) and the plant-based milk in the reducing method may involve fermentation of the plant-based milk by the lactic acid bacterial strain(s).

[0010] The present invention also provides the *Lactobacillus fermentum* CP3024 strain of the accession number: NITE BP-3428, the *Lactobacillus reuteri* CP3017 strain of the accession number: NITE BP-3426, and the *Lactobacillus reuteri* CP3019 strain of the accession number: NITE BP-3427.

15 ADVANTAGEOUS EFFECTS OF THE INVENTION

[0011] According to the present invention, a method for producing a plant-based milk-fermented liquid with reduced plant odor may be provided.

20 BRIEF DESCRIPTION OF THE DRAWINGS

[0012] Fig. 1 is a graph showing the abilities of lactic acid bacteria to reduce the amount of acetaldehyde at low concentration.

Fig. 2 is a graph showing the abilities of lactic acid bacteria to reduce the amount of acetaldehyde at high

concentration.

Fig. 3 is a graph showing the abilities of lactic acid bacteria to reduce the amount of hexanal at low concentration.

5 Fig. 4 is a graph showing the abilities of lactic acid bacteria to reduce the amount of hexanal at high concentration.

10 Fig. 5 is a graph showing the abilities of lactic acid bacteria to reduce the amount of benzaldehyde at low concentration.

Fig. 6 is a graph showing the abilities of lactic acid bacteria to reduce the amount of benzaldehyde at high concentration.

15 Fig. 7 is a graph showing the abilities of lactic acid bacteria to reduce the amount of benzaldehyde contained in a plant-based milk derived from brown rice.

Fig. 8 is a graph showing the abilities of lactic acid bacteria to reduce the amount of 2,4-decadienal contained in a plant-based milk derived from brown rice.

20 Fig. 9 is a graph showing the abilities of lactic acid bacteria to reduce the amount of 2-methylbutanal contained in a plant-based milk derived from brown rice.

25 Fig. 10 is a graph showing the abilities of lactic acid bacteria to reduce the amount of 2-methylpropanal contained in a plant-based milk derived from brown rice.

Fig. 11 is a graph showing the abilities of lactic acid bacteria to reduce the amount of 3-methylbutanal contained in a plant-based milk derived from brown rice.

Fig. 12 is a graph showing the abilities of lactic acid bacteria to reduce the amount of hexanal contained in a plant-based milk derived from brown rice.

Fig. 13 is a graph showing the abilities of lactic acid bacteria to reduce the amount of benzaldehyde contained in a plant-based milk derived from white rice.

Fig. 14 is a graph showing the abilities of lactic acid bacteria to reduce the amount of hexanal contained in a plant-based milk derived from white rice.

Fig. 15 is a graph showing the abilities of lactic acid bacteria to reduce the amount of benzaldehyde contained in a plant-based milk derived from barley.

Fig. 16 is a graph showing the abilities of lactic acid bacteria to reduce the amount of 2,4-decadienal contained in a plant-based milk derived from barley.

MODE FOR CARRYING OUT THE INVENTION

[0013] In the present specification, the term "step" includes not only an independent step, but also a step that is not clearly distinguishable from another step, as long as the desired purpose of the step can be achieved. Unless otherwise specified, when a plurality of substances

corresponding to a certain component is present in a composition, the content of the component in the composition means the total amount of the plurality of substances present in the composition. The upper limit and lower limit of a numerical range described in the present description may be an arbitrary combination of values selected from those exemplified for the numerical range. Embodiments of the present invention are described below in detail. The embodiments described below, however, are merely examples of methods of producing a plant-based milk-fermented liquid for realization of the technological thought of the present invention. Therefore, the present invention is not limited to the methods of producing a plant-based milk-fermented liquid described below.

[0014] Method for Producing Plant-based milk-fermented liquid

The method for producing a plant-based milk-fermented liquid includes bringing a plant-based milk into contact with a lactic acid bacterial strain(s) including at least one selected from the group consisting of the *Lactobacillus fermentum* CP1299 strain, the *Lactobacillus fermentum* CP3024 strain, the *Lactobacillus reuteri* CP3017 strain, and the *Lactobacillus reuteri* CP3019 strain.

[0015] By bringing a particular lactic acid bacterial strain(s) into contact with a plant-based milk, the content

of aldehyde in the plant-based milk may be effectively reduced to reduce the plant odor derived from the aldehyde, to enable production of a plant-based milk-fermented liquid having an excellent yogurt-like flavor. The plant-based milk-fermented liquid produced has an excellent balance between sweetness and sourness, and an improved mellow taste as a fermented milk. This may be thought to be due to, for example, the excellent activity of the particular lactic acid bacterial strain(s) to reduce the amount of aldehyde contained in the plant-based milk.

[0016] In the method for producing a plant-based milk-fermented liquid, the content of aldehyde in a plant-based milk may be reduced by bringing a particular lactic acid bacterium/bacteria into contact with the plant-based milk. In other words, the method for producing a plant-based milk-fermented liquid may be a method for reducing the amount of aldehyde in a plant-based milk.

[0017] Lactic Acid Bacterial Strains

In the method for producing a plant-based milk-fermented liquid, a particular lactic acid bacterial strain(s) is/are used. The particular lactic acid bacterial strain(s) include(s) at least one selected from the group consisting of the *Lactobacillus fermentum* CP1299 strain, the *Lactobacillus fermentum* CP3024 strain, the *Lactobacillus reuteri* CP3017 strain, and the *Lactobacillus*

reuteri CP3019 strain. The *Lactobacillus fermentum* CP1299 strain herein is a lactic acid bacterium belonging to *Lactobacillus fermentum*, and has been internationally deposited with Patent Microorganisms Depositary, NITE
5 National Institute of Technology and Evaluation (Room 122, 2-5-8 Kazusakamatari, Kisarazu-shi, Chiba, 292-0818 Japan), which is an international depository institution in accordance with the Budapest Treaty, under the accession number: NITE BP-1512 as of January 18, 2013. The
10 *Lactobacillus fermentum* CP3024 strain is a lactic acid bacterium belonging to *Lactobacillus fermentum*, and has been internationally deposited under the accession number: NITE BP-03428 as of March 1, 2021. The *Lactobacillus reuteri* CP3017 strain is a lactic acid bacterium belonging
15 to *Lactobacillus reuteri*, and has been internationally deposited under the accession number: NITE BP-03426 as of March 1, 2021. The *Lactobacillus reuteri* CP3019 strain is a lactic acid bacterium belonging to *Lactobacillus reuteri*, and has been internationally deposited under the accession
20 number: NITE BP-03427 as of March 1, 2021.

[0018] In the method for producing a plant-based milk-fermented liquid, the particular lactic acid bacterium/bacteria may be used in combination with other lactic acid bacteria, bifidobacteria, and the like.

25 Examples of the lactic acid bacteria other than the

particular lactic acid bacterium/bacteria include the genus *Lactobacillus*, such as *Lactobacillus fermentum* (for example, the JCM1173 strain) other than the CP1299 strain and the CP3024 strain, *Lactobacillus reuteri* (for example, 5 the JCM1112 strain) other than the CP3017 strain and the CP3019 strain, *Lactobacillus gasseri*, *Lactobacillus oris*, *Lactobacillus mucosae*, *Lactobacillus fructivorans*, *Lactobacillus casei*, *Lactobacillus delbrueckii*, *Lactobacillus bulgaricus*, *Lactobacillus helveticus*, 10 *Lactobacillus brevis*, *Lactobacillus plantarum*, *Lactobacillus lactis*, *Lactobacillus acidophilus*, *Lactobacillus amylovorus*, *Lactobacillus crispatus*, *Lactobacillus gallinarum*, *Lactobacillus jensenii*, *Lactobacillus johnsonii*, *Lactobacillus kefirgranum*, 15 *Lactobacillus mali*, *Lactobacillus paracasei*, *Lactobacillus rhamnosus*, *Lactobacillus salivarius*, and *Lactobacillus tolerans*; the genus *Brucella*, such as *Brucella adolescentis*, *Brucella angulatum*, *Brucella animalis*, *Brucella bifidum*, *Brucella breve*, *Brucella catenulatum*, 20 *Brucella dentium*, *Brucella gallicum*, *Brucella infantis*, *Brucella longum*, *Brucella pseudocatenulatum*, *Brucella pseudolongum*, *Brucella suis*, and *Brucella thermophilum*; the genus *Streptococcus*, such as *Streptococcus thermophilus*, *Streptococcus lactis*, *Streptococcus cremoris*, *Streptococcus faecalis*, and *Streptococcus faecium*; the genus *Leuconostoc*, 25

such as *Leuconostoc mesenteroides*, *Leuconostoc dextranicum*,
Leuconostoc cremoris, and *Leuconostoc oenos*; and the genus
Pediococcus, such as *Pediococcus cerevisiae*, *Pediococcus*
acidilactici, and *Pediococcus halophilus*. Examples of the
5 bifidobacteria include *Bifidobacterium longum*,
Bifidobacterium bifidum, *Bifidobacterium breve*,
Bifidobacterium adolescentis, *Bifidobacterium infantis*, and
Bifidobacterium animalis.

[0019] In cases where the lactic acid bacterium/bacteria
10 used in the method for producing a plant-based milk-
fermented liquid include(s) at least one lactic acid
bacterium other than the particular lactic acid
bacterium/bacteria, and at least one bifidobacterium,
its/their content ratio may be not more than 50%,
15 preferably not more than 10%, or less than 5% with respect
to the total of the lactic acid bacterium and/or
bifidobacterium. The content ratio of the lactic acid
bacterium and/or bifidobacterium may be determined from the
number of bacterial cells of the at least one lactic acid
20 bacterium and/or bifidobacterium used. Alternatively, in
cases where a preculture is used as the at least one lactic
acid bacterium and/or bifidobacterium, the content ratio
may be determined based on the volume of the preculture.

[0020] Plant-based milk

25 The plant-based milk may be a product prepared by size

reduction and liquefaction of a plant material by physical crushing, and/or saccharification using an enzyme, malted rice, and/or the like. The plant-based milk may be a product prepared by liquefaction of a plant material by a known method, or may be a product selected from commercially available products as appropriate. Examples of the plant material include cereals such as rice, barley, wheat, oats, rye, soybeans, and peanuts; nuts and seeds such as almonds, coconuts, cashew nuts, macadamia nuts, and walnuts; and vegetables such as carrot, potato, sweet potato, cassava, and tomato. In particular, the plant material preferably contains a cereal since a cereal contains a large amount of carbohydrates. The plant material more preferably contains at least one selected from the group consisting of at least rice and barley, still more preferably contains at least rice or barley.

[0021] In cases where rice is used as the plant material, the rice may be at least one selected from the group consisting of unhulled rice, brown rice (including germinated brown rice), polished rice, red bran (akanuka), middle bran (chunuka), white bran (shironuka), and higher white bran (joshironuka). Examples of the polished rice include those with different degrees of polishing, such as 30%-polished rice, 50%-polished rice, 70%-polished rice, and white rice, and the polished rice may contain residual

germs. The rice as the plant material is preferably at least one selected from the group consisting of brown rice, polished rice, middle bran, and higher white bran, more preferably at least one selected from the group consisting of brown rice, polished rice, and higher white bran. The rice is still more preferably brown rice or white rice. The rice as the plant material may be pulverized to a predetermined grain size by a known method, or the pulverization may be omitted to leave the rice as it is.

The rice may also be in a state where it has been saccharified by a microorganism, an enzyme treatment, and/or the like. Depending on the starch contained, rice may be classified into non-glutinous rice, glutinous rice, and the like. Any of these may be used. In cases where the plant material contains rice, plant materials other than rice may also be contained. The plant material may contain, in addition to rice, for example, other cereals such as barley, wheat, oats, rye, soybeans, and peanuts; and nuts and seeds such as almonds, coconuts, cashew nuts, macadamia nuts, and walnuts. The content of the other cereals and the like in the plant material containing rice may be, for example, not more than 30% by mass, not more than 20% by mass, or not more than 10% by mass. The lower limit of the content of the other cereals and the like in the plant material containing rice may be, for example, not

less than 0.1% by mass or not less than 1% by mass.

[0022] In cases where barley is used as the plant material, it may be at least one selected from the group consisting of two-rowed barley, four-rowed barley, six-
5 rowed barley, naked barley, and wild barley, or may be at least one selected from the group consisting of two-rowed barley and six-rowed barley. The barley as the plant material may be pulverized to a predetermined grain size by a known method, or the pulverization may be omitted to
10 leave the barley as it is. The barley may be in a state where it has been saccharified by a microorganism, an enzyme treatment, and/or the like. In cases where the plant material contains barley, the plant material may also contain plant materials other than barley. The plant
15 material may contain, in addition to barley, for example, other cereals such as rice, wheat, oats, rye, soybeans, and peanuts; and nuts and seeds such as almonds, coconuts, cashew nuts, macadamia nuts, and walnuts. The content of the other cereals and the like in the plant material
20 containing barley may be, for example, not more than 30% by mass, not more than 20% by mass, or not more than 10 % by mass. The lower limit of the content of the other cereals in the plant material containing barley may be, for
25 example, not less than 0.1% by mass or not less than 1% by mass.

[0023] The plant-based milk may be, for example, a product prepared by performing saccharification treatment in which, for example, starch contained in the plant material (preferably a cereal) is hydrolyzed by the use of an appropriate enzyme. In other words, the plant-based milk may be a saccharified plant-based milk, and is preferably a plant-based milk prepared, from the viewpoint of increasing the monosaccharide content, by subjecting a plant-based milk that uses rice as a plant material to saccharification treatment. The plant-based milk is more preferably a plant-based milk prepared by subjecting a plant-based milk that uses brown rice or white rice as a plant material to saccharification treatment. As the enzyme in the saccharification treatment, for example, α -amylase (EC 3.2.1.1) may be used. The amount of the enzyme added may be, for example, 0.01% to 1% with respect to the mass of the plant material. The conditions of the hydrolysis reaction may be appropriately selected in accordance with, for example, the enzyme and the plant material used. The reaction temperature may be, for example, 40°C to 100°C, preferably 50°C to 70°C. The reaction time may be, for example, 1 hour to 24 hours, preferably 2 hours to 12 hours.

[0024] In the preparation of the plant-based milk by saccharification treatment, hydrolysis with α -amylase may

be further followed by saccharification with glucoamylase (EC 3.2.1.3) or the like. The plant-based milk may be a product obtained by saccharification treatment in which malted rice or the like is allowed to act on the plant material.

[0025] The plant-based milk prepared by the saccharification treatment contains various saccharides including monosaccharides such as glucose, disaccharides such as maltose, trisaccharides such as maltotriose, and tetra- and higher saccharides. The saccharide composition of the plant-based milk is preferably a glucose-rich composition since, while glucose is degraded by a lactic acid bacterium in the later-described fermentation step, glucose eventually becomes the main source of sweetness. The content of monosaccharides such as glucose may be, for example, not less than 30% by mass, preferably not less than 50% by mass, more preferably not less than 70% with respect to the total saccharides contained in the plant-based milk.

[0026] The plant-based milk may be in the form of, for example, a milk-like emulsion. After the size reduction and liquefaction of the plant material, the plant-based milk may be seasoned with a vegetable oil, an animal oil, salt, sugar, amino acid, a flavor, a thickener, a thickening agent, an emulsifier, a pH-adjusting agent,

and/or the like.

[0027] Aldehydes

Plant-based milk contains aldehydes derived from a plant material. The aldehydes may have, for example, a plant-odor-like flavor. Examples of the aldehydes include aliphatic aldehydes such as short-chain aliphatic aldehydes, medium-chain aliphatic aldehydes, and long-chain aliphatic aldehydes; and aromatic aldehydes. Short-chain aliphatic aldehydes have not more than 5 carbon atoms; medium-chain aliphatic aldehydes have 6 to 12 carbon atoms; and long-chain aliphatic aldehydes have not less than 13 carbon atoms. Specific examples of the aliphatic aldehydes include acetaldehyde, malondialdehyde, butanal, 2-methylpropanal, pentanal, 2-methylbutanal, 3-methylbutanal, hexanal, crotonaldehyde, 4-hydroxy-2-hexenal, 4-hydroxy-2-nonenal, 2,4-nonadienal, heptanal, octanal, nonanal, decanal, 2,4-decadienal, and tetradecanal. Specific examples of the aromatic aldehydes include benzaldehyde, anisaldehyde, and cuminaldehyde.

[0028] The aldehyde whose content is to be reduced by the method for producing a plant-based milk-fermented liquid may be aliphatic aldehydes and aromatic aldehydes, or may be at least one selected from a group consisting of short-chain aliphatic aldehydes, medium-chain aliphatic aldehydes, and aromatic aldehydes, or may be at least one

selected from a group consisting of acetaldehyde,
malondialdehyde, butanal, 2-methylpropanal, pentanal, 2-
methylbutanal, 3-methylbutanal, hexanal, crotonaldehyde, 4-
hydroxy-2-hexenal, 4-hydroxy-2-nonenal, 2,4-nonadienal,
5 heptanal, octanal, nonanal, decanal, 2,4-decadienal, and
benzaldehyde.

[0029] The method for producing a plant-based milk-
fermented liquid includes a step of bringing a particular
lactic acid bacterium/bacteria into contact with a plant-
10 based milk. The step of bringing a particular lactic acid
bacterium/bacteria into contact with a plant-based milk may
be a fermentation step of adding the particular lactic acid
bacterium/bacteria to the plant-based milk and carrying out
lactic acid fermentation. More specifically, by
15 maintaining a plant-based milk at a predetermined
fermentation temperature, adding a predetermined amount of
a culture liquid of a lactic acid bacterium/bacteria
thereto, and then carrying out lactic acid fermentation for
a predetermined fermentation time, a desired plant-based
20 milk-fermented liquid may be obtained.

[0030] The lactic acid bacterium/bacteria used for the
contact with the plant-based milk may be a preculture
preliminarily obtained by culturing in an appropriate
medium, or may be a frozen product obtained by mixing the
25 preculture with a cryoprotective agent and freezing the

resulting mixture, or may be a powder obtained by freeze-drying the preculture. The medium used for the preparation of the preculture may be selected as appropriate from media normally used for culture of lactic acid bacteria.

5 Examples of the medium include MRS Medium (manufactured by BD Japan). Regarding the culture conditions, the culture may be carried out, for example, under static or anaerobic conditions at 30°C to 40°C for 12 hours to 32 hours.

[0031] The amount of the preculture of the lactic acid
10 bacterium/bacteria added to the plant-based milk may be, for example, 0.1% by volume to 30% by volume, preferably 0.5% by volume to 20% by volume, more preferably 1% by volume to 10% by volume, with respect to the liquid volume of the plant-based milk.

15 [0032] The fermentation temperature and the fermentation time may be appropriately selected in accordance with the culture conditions, the lactic acid bacterial strain(s) added, and the like. The fermentation temperature may be, for example, 10°C to 45°C, preferably 20°C to 40°C, more
20 preferably 30°C to 40°C. The fermentation time may be, for example, 1 hour to 72 hours, preferably 6 hours to 48 hours, more preferably 12 hours to 32 hours.

[0033] In the method for producing a plant-based milk-fermented liquid, the aldehyde content in the plant-based
25 milk is reduced by the fermentation step. The residual

ratio (%), obtained by dividing the aldehyde content in the resulting plant-based milk-fermented liquid by the aldehyde content in a plant-based milk obtained by the same process except that no lactic acid bacterium is brought into
5 contact, in the case of aromatic aldehydes is, for example, not more than 85%, preferably not more than 70%, more preferably not more than 60%, still more preferably not more than 50%. The lower limit of the residual ratio (%) is not limited, and may be, for example, not less than 1%,
10 or may be preferably not less than 5%. The residual ratio (%) of aldehydes in the case of aliphatic aldehydes is, for example, not more than 55%, preferably not more than 45%, more preferably not more than 40%, still more preferably not more than 35%, still more preferably not more than 30%,
15 or not more than 20%. The lower limit of the residual ratio (%) is not limited, and may be, for example, not less than 0%, preferably not less than 1%, or may be not less than 2%.

[0034] The lactic acid bacterium/bacteria contained in
20 the plant-based milk-fermented liquid obtained in the fermentation step may be in the state of viable cells as they are, or may be killed through high-temperature treatment or the like. The high-temperature treatment may be carried out, for example, by treating the resulting
25 plant-based milk-fermented liquid for a predetermined time

at a temperature of as high as not less than 63°C.

[0035] The plant-based milk-fermented liquid obtained by the production method may be provided as a lactic acid fermented beverage as it is, or the plant-based milk-fermented liquid may be subjected to a predetermined post-treatment to provide a lactic acid fermented beverage or lactic acid fermented food in a desired form. Examples of the post-treatment include a concentration step for making the plant-based milk-fermented liquid into the form of a syrup; a step of adding a desired additive(s) (such as a flavor); a step of mixing with a carbonated beverage, a fruit juice beverage, an alcoholic beverage, or the like; and a food processing step.

[0036] Examples of the additives in the post-treatment step include sugar alcohols such as sorbitol, erythritol, maltitol, and xylitol; high-intensity sweeteners such as aspartame, stevioside, sucralose, and acesulfame K; organic acids such as citric acid, tartaric acid, malic acid, succinic acid, and lactic acid; vitamins such as L-ascorbic acid, dl- α -tocopherol, vitamin Bs, nicotinamide, and calcium pantothenate; surfactants such as glycerin fatty acid esters, polyglycerin fatty acid esters, sucrose fatty acid esters, sorbitan fatty acid esters, and propylene glycol fatty acid esters; thickening agents such as gum arabic, carrageenan, pectin, and agar; stabilizers such as

casein and gelatin; amino acids; minerals such as calcium salts; additives such as sodium ascorbate, sodium erythorbate, glycerin, and propylene glycol; dyes; flavors; and preservatives.

5 [0037] Examples of the lactic acid fermented beverage obtained by the post-treatment step include carbonated beverages, fruit juice beverages, alcoholic beverages, syrups, and fruit-flavored beverages. To the lactic acid fermented beverage, other beverages such as soy milk may be
10 added as long as the effect of the present invention is not impaired. Examples of the lactic acid fermented food obtained by the post-treatment step include cold confectionery such as jelly, yogurt, pudding, and ice cream; candy; soft candy; gum; and jam.

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EXAMPLES

[0038] The present invention is described below more concretely by way of Examples. However, the present invention is not limited to these Examples.

20 [0039] Reference Example 1

As lactic acid bacteria, the bacterial strains shown in Table 1 were provided. All bacterial strains provided were type strains. Using a medium prepared by adding 0.05% by mass cysteine hydrochloride to MRS Medium (manufactured
25 by BD Japan), each lactic acid bacterium was cultured in an

environment at 37°C under anaerobic conditions for 16 hours to prepare a preculture of the lactic acid bacterium.

Subsequently, 0.05% by mass cysteine hydrochloride and 0.05% by volume acetaldehyde were added to MRS Medium

5 (manufactured by BD Japan), and the preculture of the

lactic acid bacterium was added thereto at 5% by volume,

followed by performing culture in an environment at 37°C

under anaerobic conditions for 16 hours. Subsequently, the

bacterial cells were removed by centrifuging the culture,

10 to obtain a culture supernatant. The concentration of

acetaldehyde contained in the culture supernatant was

quantified using F-kit Acetaldehyde (manufactured by JK

International). The residual ratio (%) of acetaldehyde was

calculated by dividing the acetaldehyde concentration in

15 each culture supernatant by the average acetaldehyde

concentration in the uninoculated medium, and multiplying

the resulting value by 100. The sample data number (n

number) was set to 3. The results are shown in Table 1 and

Fig. 1.

20 [0040] [Table 1]

	Strain name	Residual ratio (%)			Strain name	Residual ratio (%)	
		Average	Standard deviation			Average	Standard deviation
uninoculated medium	-	100.0	6.2	<i>L. paracasei T</i>	JCM8130	46.8	1.9
<i>L. acidophilus T</i>	JCM1132	40.5	4.8	<i>L. plantarum T</i>	JCM1149	55.6	0.8
<i>L. amylovorus T</i>	JCM1126	39.3	1.5	<i>L. reuteri T</i>	JCM1112	0.0	0.0
<i>L. brevis T</i>	JCM1059	0.1	0.1	<i>L. rhamnosus T</i>	JCM1136	21.0	2.2
<i>L. bulgaricus T</i>	JCM1002	73.6	2.2	<i>L. salivarius T</i>	JCM1231	27.0	3.7
<i>L. casei T</i>	JCM1134	14.1	4.2	<i>L. tolerans T</i>	JCM1171	70.9	9.2
<i>L. crispatus T</i>	JCM1185	15.3	3.0	<i>B. adolescentis T</i>	JCM1275	35.7	1.5
<i>L. delbrueckii T</i>	JCM1012	61.3	4.9	<i>B. angulatum T</i>	JCM7096	98.2	1.5
<i>L. fermentum T</i>	JCM1173	0.4	0.6	<i>B. animalis T</i>	JCM1190	40.8	9.0
<i>L. fructivorans T</i>	JCM1117	0.0	0.0	<i>B. bifidum T</i>	JCM1255	78.7	9.6
<i>L. gallinarum T</i>	JCM2011	40.2	3.0	<i>B. breve T</i>	JCM1192	26.4	0.7
<i>L. gasseri T</i>	JCM1131	42.6	2.7	<i>B. catenulatum T</i>	JCM1194	62.5	1.5
<i>L. helveticus T</i>	JCM1120	46.5	10.0	<i>B. dentium T</i>	JCM1195	31.8	0.7
<i>L. jensenii T</i>	JCM15953	42.0	2.2	<i>B. gallicum T</i>	JCM8224	82.9	4.1
<i>L. johnsonii T</i>	JCM2012	55.9	1.9	<i>B. infantis T</i>	JCM1222	65.2	0.7
<i>L. kefirgranum T</i>	JCM8572	49.5	2.1	<i>B. longum T</i>	JCM1217	50.2	2.1
<i>L. lactis T</i>	JCM1248	55.6	5.5	<i>B. pseudocatenulatum T</i>	JCM1200	10.2	1.5
<i>L. mali T</i>	JCM1116	0.4	0.6	<i>B. pseudolongum T</i>	JCM1205	73.3	3.9
<i>L. mucosae T</i>	JCM12515	0.0	0.0	<i>B. suis T</i>	JCM1269	61.3	2.4
<i>L. oris T</i>	JCM11028	0.0	0.0	<i>B. thermophilum T</i>	JCM1207	48.0	2.7

[0041] It can be seen that the ability to reduce the amount of acetaldehyde varies among the lactic acid bacterial species.

[0042] Reference Example 2

The ability of each lactic acid bacterium to reduce the amount of acetaldehyde was evaluated in the same manner as in Reference Example 1 except that the bacterial strains shown in Table 2 were used, and that the concentration of the acetaldehyde added was 0.2% by volume. The results are shown in Table 2 and Fig. 2. For statistical processing, the Student t-test was used. The test was carried out based on comparison with the residual ratio in the case of *Lactobacillus brevis T*. A significant difference was

assumed at a level of $P < 0.05$. The sample data number (n number) was set to 3.

[0043] [Table 2]

	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. brevis</i> T)
uninoculated medium	-	100.0	3.4	0.025
<i>L. gasseri</i> T	JCM1131	116.7	3.6	0.065
<i>L. fructivorans</i> T	JCM1117	107.0	1.9	0.233
<i>L. brevis</i> T	JCM1059	109.6	1.9	
<i>L. fermentum</i> T	JCM1173	1.4	0.4	0.000
<i>L. fermentum</i> CP1299	CP1299	0.0	0.0	0.000
<i>L. fermentum</i> CP3024	CP3024	0.0	0.0	0.000
<i>L. reuteri</i> T	JCM1112	0.0	0.0	0.000
<i>L. reuteri</i> CP3017	CP3017	0.0	0.0	0.000
<i>L. reuteri</i> CP3019	CP3019	0.0	0.0	0.000
<i>L. mucosae</i> T	JCM12515	0.0	0.0	0.000
<i>L. oris</i> T	JCM11028	0.0	0.0	0.000

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[0044] Under the high-concentration conditions, particular bacterial species were found to show higher abilities to reduce the amount of acetaldehyde compared to *Lactobacillus brevis* T. Under the low-concentration

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conditions, *Lactobacillus fructivorans* T was found to show a high ability to reduce the amount of acetaldehyde similarly to particular bacterial species. However, under the high-concentration conditions, the ability of this species was equivalent to that of *Lactobacillus brevis* T.

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It can thus be seen that the ability to reduce the amount

of acetaldehyde varies among the bacterial species and bacterial strains.

[0045] Reference Example 3

As lactic acid bacteria, the bacterial strains shown
5 in Table 3 were provided. All strains provided were type strains. Using a medium prepared by adding 0.05% by mass cysteine hydrochloride to MRS Medium (manufactured by BD Japan), each lactic acid bacterium was cultured in an environment at 37°C under anaerobic conditions for 16 hours
10 to prepare a preculture of the lactic acid bacterium. Subsequently, 0.05% by mass cysteine hydrochloride and 0.05% by volume hexanal were added to MRS Medium (manufactured by BD Japan), and the preculture of the lactic acid bacterium was added thereto at 5% by volume,
15 followed by performing culture in an environment at 37°C under anaerobic conditions for 16 hours. Subsequently, the bacterial cells were removed by centrifuging the culture, to obtain a culture supernatant. The concentration of hexanal contained in the culture supernatant was quantified
20 by a modification of the LC-MS/MS method described in Drug Test. Analysis, 8, 458-464 (2016). Details of the quantification method are described later. The residual ratio (%) of hexanal was calculated by dividing the hexanal concentration in each culture supernatant by the average
25 hexanal concentration in the uninoculated medium, and

multiplying the resulting value by 100. The sample data number (n number) was set to 3. The results are shown in Table 3 and Fig. 3.

[0046] [Table 3]

	Strain name	Residual ratio (%)			Strain name	Residual ratio (%)		
		Average	Standard deviation			Average	Standard deviation	
uninoculated medium	-	100.0	12.1		<i>L. jensenii T</i>	JCM15953	45.2	2.9
<i>L. acidophilus T</i>	JCM1132	81.6	3.5		<i>L. johnsonii T</i>	JCM2012	41.0	0.7
<i>L. amylovorus T</i>	JCM1126	66.1	5.9		<i>L. kefirgranum T</i>	JCM8572	79.8	3.1
<i>L. brevis T</i>	JCM1059	7.7	0.4		<i>L. lactis T</i>	JCM1248	29.5	1.4
<i>L. bulgaricus T</i>	JCM1002	91.4	0.9		<i>L. mali T</i>	JCM1116	8.7	5.1
<i>L. casei T</i>	JCM1134	30.3	2.3		<i>L. mucosae T</i>	JCM12515	13.3	1.8
<i>L. crispatus T</i>	JCM1185	37.4	0.6		<i>L. oris T</i>	JCM11028	15.5	3.4
<i>L. delbrueckii T</i>	JCM1012	46.9	1.8		<i>L. paracasei T</i>	JCM8130	77.2	3.5
<i>L. fermentum T</i>	JCM1173	2.2	0.6		<i>L. plantarum T</i>	JCM1149	44.0	5.3
<i>L. fructivorans T</i>	JCM1117	5.8	2.1		<i>L. reuteri T</i>	JCM1112	14.5	0.3
<i>L. gallinarum T</i>	JCM2011	100.2	13.2		<i>L. rhamnosus T</i>	JCM1136	30.0	5.4
<i>L. gasseri T</i>	JCM1131	69.0	3.5		<i>L. salivarius T</i>	JCM1231	62.7	1.2
<i>L. helveticus T</i>	JCM1120	77.9	1.2		<i>L. tolerans T</i>	JCM1171	86.8	7.1

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[0047] It could be confirmed that the ability to reduce the amount of hexanal varies among the lactic acid bacterial species.

10 [0048] Reference Example 4

The ability of each lactic acid bacterium to reduce the amount of hexanal was evaluated in the same manner as in Reference Example 3 except that the bacterial strains shown in Table 4 were used, and that the concentration of the hexanal added was 0.2% by volume. The results are shown in Table 4 and Fig. 4. For statistical processing, the Student t-test was used. The test was carried out based on comparison with the residual ratio in the case of

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Lactobacillus brevis T. A significant difference was assumed at a level of $P < 0.05$. The sample data number (n number) was set to 3.

[0049] [Table 4]

	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. brevis</i> T)
uninoculated medium	-	100.0	4.8	0.000
<i>L. gasseri</i> T	JCM1131	78.3	5.7	0.001
<i>L. fructivorans</i> T	JCM1117	43.8	1.1	0.420
<i>L. brevis</i> T	JCM1059	42.9	1.0	
<i>L. fermentum</i> T	JCM1173	0.8	0.5	0.000
<i>L. fermentum</i> CP1299	CP1299	1.0	1.0	0.000
<i>L. fermentum</i> CP3024	CP3024	1.5	2.1	0.000
<i>L. reuteri</i> T	JCM1112	2.6	0.2	0.000
<i>L. reuteri</i> CP3017	CP3017	5.2	0.8	0.000
<i>L. reuteri</i> CP3019	CP3019	0.6	0.5	0.000
<i>L. mucosae</i> T	JCM12515	6.5	1.1	0.000
<i>L. oris</i> T	JCM11028	5.9	0.9	0.000

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[0050] Under the high-concentration conditions, particular bacterial species were found to show higher abilities to reduce the amount of hexanal compared to *Lactobacillus brevis* T. Under the low-concentration conditions, *Lactobacillus fructivorans* T was found to show a high ability to reduce the amount of hexanal similarly to particular bacterial species. However, under the high-concentration conditions, the ability of this species to reduce the amount of hexanal was equivalent to that of

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Lactobacillus brevis T. It can thus be seen that the ability to reduce the amount of hexanal varies among the bacterial species and bacterial strains.

[0051] Reference Example 5

5 As lactic acid bacteria, the bacterial strains shown in Table 5 were provided. All strains provided were type strains. Using a medium prepared by adding 0.05% by mass cysteine hydrochloride to MRS Medium (manufactured by BD Japan), each lactic acid bacterium was cultured in an
10 environment at 37°C under anaerobic conditions for 16 hours to prepare a preculture of the lactic acid bacterium. Subsequently, 0.05% by mass cysteine hydrochloride and 0.05% by volume benzaldehyde were added to MRS Medium (manufactured by BD Japan), and the preculture of the
15 lactic acid bacterium was added thereto at 5% by volume, followed by performing culture in an environment at 37°C under anaerobic conditions for 16 hours. Subsequently, the bacterial cells were removed by centrifuging the culture, to obtain a culture supernatant. The concentration of
20 benzaldehyde contained in the culture supernatant was quantified by a modification of the LC-MS/MS method described in Drug Test. Analysis, 8, 458-464 (2016). Details of the quantification method are described later. The residual ratio (%) of benzaldehyde was calculated by
25 dividing the benzaldehyde concentration in each culture

supernatant by the average benzaldehyde concentration in the uninoculated medium, and multiplying the resulting value by 100. The sample data number (n number) was set to 3. The results are shown in Table 5 and Fig. 5.

5 [0052] [Table 5]

	Strain name	Residual ratio (%)			Strain name	Residual ratio (%)	
		Average	Standard deviation			Average	Standard deviation
uninoculated medium	-	100.0	7.8	<i>L. jensenii T</i>	JCM15953	74.0	1.0
<i>L. acidophilus T</i>	JCM1132	92.3	1.6	<i>L. johnsonii T</i>	JCM2012	74.1	0.6
<i>L. amylovorus T</i>	JCM1126	50.8	9.5	<i>L. kefirgranum T</i>	JCM8572	69.5	3.7
<i>L. brevis T</i>	JCM1059	3.3	1.3	<i>L. lactis T</i>	JCM1248	41.1	0.5
<i>L. bulgaricus T</i>	JCM1002	113.5	5.3	<i>L. mali T</i>	JCM1116	39.6	3.6
<i>L. casei T</i>	JCM1134	43.5	5.7	<i>L. mucosae T</i>	JCM12515	2.6	0.3
<i>L. crispatus T</i>	JCM1185	56.0	1.3	<i>L. oris T</i>	JCM11028	2.5	0.2
<i>L. delbrueckii T</i>	JCM1012	66.0	4.8	<i>L. paracasei T</i>	JCM8130	92.0	7.3
<i>L. fermentum T</i>	JCM1173	1.5	0.6	<i>L. plantarum T</i>	JCM1149	42.9	1.6
<i>L. fructivorans T</i>	JCM1117	4.9	0.5	<i>L. reuteri T</i>	JCM1112	2.1	0.3
<i>L. gallinarum T</i>	JCM2011	84.7	4.4	<i>L. rhamnosus T</i>	JCM1136	52.0	1.2
<i>L. gasseri T</i>	JCM1131	90.8	6.1	<i>L. salivarius T</i>	JCM1231	100.0	3.1
<i>L. helveticus T</i>	JCM1120	107.4	5.7	<i>L. tolerans T</i>	JCM1171	103.4	7.2

[0053] It could be confirmed that the ability to reduce the amount of benzaldehyde varies among the lactic acid bacterial species.

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[0054] Reference Example 6

The ability of each lactic acid bacterium to reduce the amount of benzaldehyde was evaluated in the same manner as in Reference Example 5 except that the bacterial strains shown in Table 6 were used, and that the concentration of the benzaldehyde added was 0.2% by volume. The results are shown in Table 6 and Fig. 6. For statistical processing, the Student t-test was used. The test was carried out

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based on comparison with the residual ratio in the case of *Lactobacillus brevis* T. A significant difference was assumed at a level of $P < 0.05$. The sample data number (n number) was set to 3.

5 [0055] [Table 6]

	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. brevis</i> T)
uninoculated medium	-	100.0	6.5	0.002
<i>L. gasseri</i> T	JCM1131	79.4	2.6	0.004
<i>L. fructivorans</i> T	JCM1117	80.8	18.1	0.249
<i>L. brevis</i> T	JCM1059	63.4	2.9	
<i>L. fermentum</i> T	JCM1173	7.8	1.3	0.000
<i>L. fermentum</i> CP1299	CP1299	5.5	0.6	0.000
<i>L. fermentum</i> CP3024	CP3024	5.6	0.2	0.000
<i>L. reuteri</i> T	JCM1112	4.8	0.8	0.000
<i>L. reuteri</i> CP3017	CP3017	4.7	0.1	0.000
<i>L. reuteri</i> CP3019	CP3019	5.2	0.6	0.000
<i>L. mucosae</i> T	JCM12515	1.1	0.1	0.000
<i>L. oris</i> T	JCM11028	4.7	0.6	0.000

[0056] Under the high-concentration conditions, particular bacterial species were found to show higher benzaldehyde-reducing abilities compared to *Lactobacillus brevis* T. Under the low-concentration conditions, *Lactobacillus fructivorans* T was found to show a high ability to reduce the amount of benzaldehyde similarly to particular bacterial species. However, under the high-concentration conditions, the ability of this species to

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reduce the amount of benzaldehyde was equivalent to that of *Lactobacillus brevis* T. It can thus be seen that the ability to reduce the amount of benzaldehyde varies among the bacterial species and bacterial strains.

5 [0057] Example 1

As lactic acid bacteria, the bacterial strains shown in Table 7 to Table 12 were provided. Using MRS Medium (manufactured by BD Japan) supplemented with 0.05% by mass cysteine hydrochloride, each lactic acid bacterium was
10 cultured in an environment at 37°C under anaerobic conditions for 16 hours, to prepare a preculture of each lactic acid bacterial strain. Concentrated rice milk (manufactured by Kikkoman Corporation; raw material: processed brown rice) was diluted 1.67 times with pure
15 water, and sterilized at 95°C for 15 minutes, to obtain a brown rice milk medium. To the brown rice milk medium, the preculture of the lactic acid bacterium was added at 5% by volume, and static culture was performed in an environment at 37°C for 16 hours, to obtain a culture as a plant-based
20 milk-fermented liquid. The concentration of each of benzaldehyde, 2,4-decadienal, 2-methylbutanal, 2-methylpropanal, 3-methylbutanal, and hexanal in the obtained culture was quantified by a modification of the LC-MS/MS method described in Drug Test. Analysis, 8, 458-
25 464 (2016). Details of the quantification method are

described later. The residual ratio (%) of each aldehyde was calculated by dividing the concentration of the aldehyde in each culture by the average of the concentration of the aldehyde in the uninoculated medium, and multiplying the resulting value by 100. The results are shown in Table 7 to Table 12, and Fig. 7 to Fig. 12. For statistical processing, the Student t-test was used. The test was carried out based on comparison with the residual ratio in the case of *Lactobacillus gasseri* T. The sample data number (n number) was set to 3. A significant difference was assumed at a level of $P < 0.05$.

[0058] [Table 7]

Benzaldehyde	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. gasseri</i> T)
uninoculated medium	-	100.0	8.1	0.413
<i>L. gasseri</i> T	JCM1131	94.1	4.2	
<i>L. fermentum</i> T	JCM1173	48.4	1.7	0.000
<i>L. fermentum</i> CP3024	CP3024	35.2	3.4	0.000
<i>L. fermentum</i> CP1299	CP1299	40.0	2.0	0.000
<i>L. reuteri</i> T	JCM1112	39.6	10.2	0.002
<i>L. reuteri</i> CP3017	CP3017	56.0	5.7	0.002
<i>L. reuteri</i> CP3019	CP3019	43.6	4.9	0.000
<i>L. oris</i> T	JCM11028	42.0	4.7	0.000
<i>L. mucosae</i> T	JCM12515	42.1	1.3	0.000

[0059] [Table 8]

2,4-Decadienal	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. gasseri T</i>)
uninoculated medium	-	100.0	0.8	0.438
<i>L. gasseri T</i>	JCM1131	111.5	18.8	
<i>L. fermentum T</i>	JCM1173	1.3	0.7	0.001
<i>L. fermentum CP3024</i>	CP3024	1.2	0.0	0.001
<i>L. fermentum CP1299</i>	CP1299	1.0	0.3	0.001
<i>L. reuteri T</i>	JCM1112	1.3	0.5	0.001
<i>L. reuteri CP3017</i>	CP3017	1.9	0.1	0.001
<i>L. reuteri CP3019</i>	CP3019	1.3	0.3	0.001
<i>L. oris T</i>	JCM11028	1.7	0.6	0.001
<i>L. mucosae T</i>	JCM12515	1.0	0.4	0.001

[0060] [Table 9]

2-Methylbutanal	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. gasseri T</i>)
uninoculated medium	-	100.0	3.5	0.340
<i>L. gasseri T</i>	JCM1131	92.9	8.5	
<i>L. fermentum T</i>	JCM1173	21.1	8.2	0.001
<i>L. fermentum CP3024</i>	CP3024	27.6	9.6	0.002
<i>L. fermentum CP1299</i>	CP1299	44.4	2.4	0.002
<i>L. reuteri T</i>	JCM1112	34.4	11.0	0.004
<i>L. reuteri CP3017</i>	CP3017	31.7	9.1	0.002
<i>L. reuteri CP3019</i>	CP3019	51.3	1.3	0.002
<i>L. oris T</i>	JCM11028	28.4	13.4	0.005
<i>L. mucosae T</i>	JCM12515	38.0	11.8	0.006

[0061] [Table 10]

2-Methylpropanal	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. gasseri T</i>)
uninoculated medium	-	100.0	5.7	0.276
<i>L. gasseri T</i>	JCM1131	94.7	1.6	
<i>L. fermentum T</i>	JCM1173	23.9	4.8	0.000
<i>L. fermentum CP3024</i>	CP3024	30.4	7.4	0.000
<i>L. fermentum CP1299</i>	CP1299	41.8	1.4	0.000
<i>L. reuteri T</i>	JCM1112	32.5	5.5	0.000
<i>L. reuteri CP3017</i>	CP3017	37.1	7.3	0.000
<i>L. reuteri CP3019</i>	CP3019	48.6	0.9	0.000
<i>L. oris T</i>	JCM11028	32.5	12.6	0.002
<i>L. mucosae T</i>	JCM12515	37.5	8.5	0.001

[0062] [Table 11]

3-Methylbutanal	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. gasseri T</i>)
uninoculated medium	-	100.0	3.5	0.015
<i>L. gasseri T</i>	JCM1131	112.4	2.4	
<i>L. fermentum T</i>	JCM1173	14.9	3.6	0.000
<i>L. fermentum CP3024</i>	CP3024	9.3	1.4	0.000
<i>L. fermentum CP1299</i>	CP1299	12.1	1.5	0.000
<i>L. reuteri T</i>	JCM1112	33.9	2.5	0.000
<i>L. reuteri CP3017</i>	CP3017	34.9	9.7	0.000
<i>L. reuteri CP3019</i>	CP3019	38.6	2.2	0.000
<i>L. oris T</i>	JCM11028	9.4	4.5	0.000
<i>L. mucosae T</i>	JCM12515	30.9	4.3	0.000

[0063] [Table 12]

Hexanal	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. gasseri</i> T)
uninoculated medium	-	100.0	2.4	0.000
<i>L. gasseri</i> T	JCM1131	50.0	2.1	
<i>L. fermentum</i> T	JCM1173	36.0	3.0	0.006
<i>L. fermentum</i> CP3024	CP3024	28.3	2.3	0.001
<i>L. fermentum</i> CP1299	CP1299	28.4	1.8	0.000
<i>L. reuteri</i> T	JCM1112	31.5	5.3	0.010
<i>L. reuteri</i> CP3017	CP3017	39.6	2.4	0.010
<i>L. reuteri</i> CP3019	CP3019	28.9	2.1	0.001
<i>L. oris</i> T	JCM11028	27.1	7.5	0.015
<i>L. mucosae</i> T	JCM12515	31.2	1.3	0.000

[0064] It could be confirmed that the amounts of aldehydes were significantly reduced by particular bacterial strains compared to the case of the control, *Lactobacillus gasseri* T.

[0065] Example 2

As lactic acid bacteria, the bacterial strains shown in Table 13 and Table 14 were provided. Using MRS Medium (manufactured by BD Japan) supplemented with 0.05% by mass cysteine hydrochloride, each lactic acid bacterium was cultured in an environment at 37°C under anaerobic conditions for 16 hours, to prepare a preculture of each lactic acid bacterial strain. A commercially available confectionery rice flour (Tomizawa Shoten; material: non-glutinous rice) and pure water were mixed together at a weight ratio of 3:7, and amylase (manufactured by Novozymes, BAN480L) was added thereto at 0.03% by volume.

Thereafter, the mixture was allowed to react at 60°C for 6 hours, and then stirred, followed by sterilization at 95°C for 15 minutes to provide a white-rice milk medium. To the white-rice milk medium, the preculture of the lactic acid bacterium was added at 5% by volume, and static culture was performed in an environment at 37°C for 16 hours, to obtain a culture as a plant-based milk-fermented liquid. The concentration of each of benzaldehyde and hexanal contained in the resulting culture was quantified by a modification of the LC-MS/MS method described in Drug Test. Analysis, 8, 458-464 (2016). Details of the quantification method are described later. The residual ratio (%) of each aldehyde was calculated by dividing the concentration of the aldehyde in each culture by the average of the concentration of the aldehyde in the uninoculated medium, and multiplying the resulting value by 100. The results are shown in Table 13 and Table 14, and Fig. 13 and Fig. 14. For statistical processing, the Student t-test was used. The test was carried out based on comparison with the residual ratio in the case of *Lactobacillus gasseri* T. The sample data number (n number) was set to 3. A significant difference was assumed at a level of $P < 0.05$.

[0066] [Table 13]

Bezaldehyde	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. gasseri T</i>)
uninoculated medium	-	100.0	8.1	0.002
<i>L. gasseri T</i>	JCM1131	143.1	2.3	
<i>L. fermentum T</i>	JCM1173	42.7	4.8	0.000
<i>L. fermentum CP3024</i>	CP3024	35.8	5.4	0.000
<i>L. fermentum CP1299</i>	CP1299	57.5	8.1	0.000
<i>L. reuteri T</i>	JCM1112	82.9	2.0	0.000
<i>L. reuteri CP3017</i>	CP3017	65.1	1.9	0.000
<i>L. reuteri CP3019</i>	CP3019	49.6	2.2	0.000
<i>L. oris T</i>	JCM11028	42.7	8.3	0.000
<i>L. mucosae T</i>	JCM12515	44.0	4.9	0.000

[0067] [Table 14]

Hexanal	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. gasseri T</i>)
uninoculated medium	-	100.0	12.1	0.025
<i>L. gasseri T</i>	JCM1131	69.8	2.3	
<i>L. fermentum T</i>	JCM1173	14.9	2.1	0.000
<i>L. fermentum CP3024</i>	CP3024	18.1	2.4	0.000
<i>L. fermentum CP1299</i>	CP1299	22.6	7.5	0.001
<i>L. reuteri T</i>	JCM1112	31.0	4.4	0.000
<i>L. reuteri CP3017</i>	CP3017	21.1	3.9	0.000
<i>L. reuteri CP3019</i>	CP3019	16.3	0.1	0.000
<i>L. oris T</i>	JCM11028	20.7	1.6	0.000
<i>L. mucosae T</i>	JCM12515	21.8	3.4	0.000

[0068] It could be confirmed that the amounts of
5 aldehydes were significantly reduced by particular
bacterial strains compared to the case of the control,
Lactobacillus gasseri T.

[0069] Example 3

As lactic acid bacteria, the bacterial strains shown in Table 15 and Table 16 were provided. Using MRS Medium (manufactured by BD Japan) supplemented with 0.05% by mass cysteine hydrochloride, each lactic acid bacterium was

5 cultured in an environment at 37°C under anaerobic conditions for 16 hours, to prepare a preculture of each lactic acid bacterial strain. A commercially available barley flour (Oomugi-club; material: whole grain flour) and pure water were mixed together at a weight ratio of 2:8,

10 and amylase (manufactured by Novozymes, BAN480L) was added thereto at 0.02% by volume. Thereafter, the mixture was allowed to react with stirring at 60°C for 6 hours, and then sterilized at 95°C 15 minutes to provide a barley milk medium. To the barley milk medium, the preculture of the

15 lactic acid bacterium was added at 5% by volume, and static culture was performed in an environment at 37°C for 16 hours, to obtain a culture as a plant-based milk-fermented liquid. The concentration of each of benzaldehyde and 2,4-decadienal contained in the resulting culture was

20 quantified by a modification of the LC-MS/MS method described in Drug Test. Analysis, 8, 458-464 (2016). Details of the quantification method are described later. The residual ratio (%) of each aldehyde was calculated by dividing the concentration of the aldehyde in each culture

25 by the average of the concentration of the aldehyde in the

uninoculated medium, and multiplying the resulting value by 100. The results are shown in Table 15 and Table 16, and Fig. 15 and Fig. 16. For statistical processing, the Student t-test was used. The test was carried out based on comparison with the residual ratio in the case of *Lactobacillus gasseri* T. The sample data number (n number) was set to 3. A significant difference was assumed at a level of $P < 0.05$.

[0070] [Table 15]

Bezaldehyde	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. gasseri</i> T)
uninoculated medium	—	100.0	1.0	0.798
<i>L. gasseri</i> T	JCM1131	98.0	10.6	
<i>L. fermentum</i> T	JCM1173	36.5	12.3	0.006
<i>L. fermentum</i> CP3024	CP3024	32.8	9.4	0.003
<i>L. fermentum</i> CP1299	CP1299	40.3	6.8	0.003
<i>L. reuteri</i> T	JCM1112	63.3	8.3	0.022
<i>L. reuteri</i> CP3017	CP3017	65.2	6.5	0.020
<i>L. reuteri</i> CP3019	CP3019	56.3	8.3	0.012

[0071] [Table 16]

2,4-Decadienal	Strain name	Residual ratio (%)		
		Average	Standard deviation	p-value (vs. <i>L. gasseri</i> T)
uninoculated medium	—	100.0	32.8	0.844
<i>L. gasseri</i> T	JCM1131	108.1	43.7	
<i>L. fermentum</i> T	JCM1173	0.0	0.0	0.025
<i>L. fermentum</i> CP3024	CP3024	9.7	4.1	0.034
<i>L. fermentum</i> CP1299	CP1299	5.4	7.6	0.031
<i>L. reuteri</i> T	JCM1112	0.0	0.0	0.025
<i>L. reuteri</i> CP3017	CP3017	7.4	5.4	0.032
<i>L. reuteri</i> CP3019	CP3019	1.7	2.5	0.026

[0072] It could be confirmed that the amounts of aldehydes were significantly reduced by particular bacterial strains compared to the case of the control, *Lactobacillus gasseri* T.

[0073] Method of Quantifying Aldehydes

Hexanal d-12 (manufactured by Sigma-Aldrich) was used alone as an internal standard. An internal standard solution was prepared by dissolving hexanal d-12 at a concentration of 25 ng/ μ l in 50 % by volume aqueous ethanol solution. After mixing 40 μ l of the internal standard solution, 80 μ l of a solution prepared by dissolving dibutylhydroxytoluene (manufactured by Tokyo Chemical Industry Co., Ltd.) at a concentration of 1 mg/ml in 100 % by volume ethanol, 440 μ l of 100 % by volume ethanol, and 440 μ l of a culture supernatant or culture for which aldehydes were to be quantified, the resulting mixture was centrifuged at an acceleration of 3700 g for 10 minutes,

followed by collecting 750 µl of the supernatant. After mixing 325 µl of this supernatant with 325 µl of a solution prepared by dissolving 2,4-dinitrophenylhydrazine hydrochloride (manufactured by Tokyo Chemical Industry Co., Ltd.) at a concentration of 0.05 mol/l in the mixed solvent "acetonitrile : acetic acid : pure water = 80 : 10 : 10 (volume ratio)", the resulting mixture was allowed to react in an environment at 60°C for 2 hours. Thereafter, the resulting reaction product was mixed with 1 ml of pure water, and then liquid-liquid extraction was carried out a total of four times using 500 µl of hexane. The hexane solution obtained by the liquid-liquid extraction was dried and solidified using a centrifugal evaporator, and the dried and solidified product was dissolved in 1 ml of the mixed solvent "0.03 % by volume aqueous acetic acid solution : acetonitrile = 60 : 40 (volume ratio)". The resulting solution was subjected to high-performance liquid chromatography (HPLC) as a derivatized sample.

[0074] As a column and a guard column for the HPLC, an Atlantis T3 C18 column (3 µm, 2.1×150 mm), manufactured by Waters, and an Atlantis guard column T3 C18 (3 µm, 2.1×10 mm), manufactured by Waters, were used according to the method described in Drug Test. Analysis, 8, 458-464 (2016). For a modified LC-MS/MS equipment, the Nextera HPLC system (communication bus module: CBM-20A; pump: LC-30AD;

autosampler: SIL-30AC; degasser: DGU-20A5R; column oven: CTO-20AC), manufactured by Shimadzu Corporation, was used as a high-performance liquid chromatography (HPLC) system, and Sciex Triple quad 6500+, manufactured by AB SCIEX, was used as a tandem mass spectrometry (MS/MS) system.

[0075] The following modified conditions were used for the LC-MS/MS. As mobile phase A, 0.03 % by volume aqueous acetic acid solution was used. As mobile phase B, 100 % by volume acetonitrile was used. At the beginning, "mobile phase A: B = 60:40 (volume ratio)" was used as the initial concentration. Two minutes later, the concentration was set to "mobile phase A: B = 52:48 (volume ratio)", and this condition was maintained for 37 minutes. Four minutes thereafter, the concentration was set to "mobile phase A: B = 0:100 (volume ratio)", and this condition was maintained for 8 minutes. The sample injection volume for the HPLC was 1 μ l. Modified conditions commonly employed for the compounds were as shown in Table 17 below, and modified conditions specifically employed for each compound were as shown in Table 18 below.

[0076] [Table 17]

Curtain gas (psi)	Collision activated dissociation gas (psi)	Ion spray voltage floating (volts)	Temperature (°C)	Ion source gas1 (psi)	Ion source gas2 (psi)	Entrance potential (volts)
30	8	-4500	500	30	70	-10

[0077] [Table 18]

	Declustering potential (volts)	Collision energy (volts)	Collision cell exit potential (volts)	Elution time (min)	Precursor ion (m/z)	Product ion (m/z)
Benzaldehyde	-35	-20	-13	31.3	284.8	162.8
Hexanal d-12	-5	-20	-1	45.6	291.0	163.1
Hexanal	-35	-32	-11	45.7	278.9	152.0
2-Methylpropanal	-40	-16	-11	25.5	251.0	162.8
3-Methylbutanal	-5	-26	-11	37.2	264.9	152.1
2,4-Decadienal	-40	-26	-19	47.7	331.0	180.9
2-Methylbutanal	-25	-18	-13	38.6	265.0	100.9
2,4-Nonadienal	-5	-24	-9	47.2	317.0	181.1

[0078] Based on the above conditions, a quantitative method was established in the multiple reaction monitoring mode using hexanal d-12 as an internal standard.

Benzaldehyde (manufactured by Tokyo Chemical Industry Co., Ltd.), hexanal (manufactured by Tokyo Chemical Industry Co., Ltd.), 2-methylpropanal (manufactured by Tokyo Chemical Industry Co., Ltd.), 3-methylbutanal (manufactured by Tokyo Chemical Industry Co., Ltd.), 2,4-decadienal (manufactured by Tokyo Chemical Industry Co., Ltd.), 2-methylbutanal (manufactured by Tokyo Chemical Industry Co., Ltd.), and 2,4-nonadienal (manufactured by Tokyo Chemical Industry Co., Ltd.) were derivatized under the conditions described above, to prepare calibration curves for the aldehydes. Thereafter, the aldehydes contained in the culture or culture supernatant were derivatized under the above conditions, and the concentrations of the aldehydes

contained in the culture supernatant (Reference Examples 3 to 6) or in the culture (Examples 1 to 3) were quantified.

[0079] The disclosure of Japanese patent application No. 2021-052277 (filing date: March 25, 2021) is incorporated
5 herein by reference in its entirety. All documents, patent applications, and technical standards cited in the present description are incorporated herein by reference to the same extent as in cases where the individual documents, patent applications, and technical standards are
10 specifically and individually described to be incorporated by reference.

Accession Numbers

[0080] NITE BP-1512
15 NITE BP-03426
NITE BP-03427
NITE BP-03428

CLAIMS

[Claim 1] A method for producing a plant-based milk-fermented liquid, the method comprising bringing a plant-based milk into contact with a lactic acid bacterial strain(s) including at least one selected from the group consisting of the *Lactobacillus fermentum* CP1299 strain, the *Lactobacillus fermentum* CP3024 strain, the *Lactobacillus reuteri* CP3017 strain, and the *Lactobacillus reuteri* CP3019 strain.

10 [Claim 2] The method for producing a plant-based milk-fermented liquid according to claim 1, wherein the plant-based milk is derived from a cereal.

[Claim 3] The method for producing a plant-based milk-fermented liquid according to claim 1 or 2, wherein the plant-based milk is derived from rice.

[Claim 4] The method for producing a plant-based milk-fermented liquid according to any one of claims 1 to 3, wherein the plant-based milk is a saccharified plant-based milk.

20 [Claim 5] The method for producing a plant-based milk-fermented liquid according to any one of claims 1 to 4, wherein the contact between the lactic acid bacterial strain(s) and the plant-based milk comprises fermentation of the plant-based milk by the lactic acid bacterial strain(s).

25

[Claim 6] A method for reducing the amount of aldehyde in a plant-based milk, the method comprising bringing a plant-based milk containing aldehyde into contact with a lactic acid bacterial strain(s) including at least one
5 selected from the group consisting of the *Lactobacillus fermentum* CP1299 strain, the *Lactobacillus fermentum* CP3024 strain, the *Lactobacillus reuteri* CP3017 strain, and the *Lactobacillus reuteri* CP3019 strain.

[Claim 7] The method for reducing the amount of aldehyde
10 in a plant-based milk according to claim 6, wherein the plant-based milk is derived from a cereal.

[Claim 8] The method for reducing the amount of aldehyde in a plant-based milk according to claim 6 or 7, wherein the plant-based milk is derived from rice.

15 [Claim 9] The method for reducing the amount of aldehyde in a plant-based milk according to any one of claims 6 to 8, wherein the plant-based milk is a saccharified plant-based milk.

[Claim 10] The method for reducing the amount of
20 aldehyde in a plant-based milk according to any one of claims 6 to 9, wherein the contact between the lactic acid bacterial strain(s) and the plant-based milk involves fermentation of the plant-based milk by the lactic acid bacterial strain(s).

25 [Claim 11] A *Lactobacillus fermentum* CP3024 strain of

the accession number: NITE BP-3428.

[Claim 12] A *Lactobacillus reuteri* CP3017 strain of the
accession number: NITE BP-3426.

[Claim 13] A *Lactobacillus reuteri* CP3019 strain of the
5 accession number: NITE BP-3427.

DRAWINGS

Fig. 1

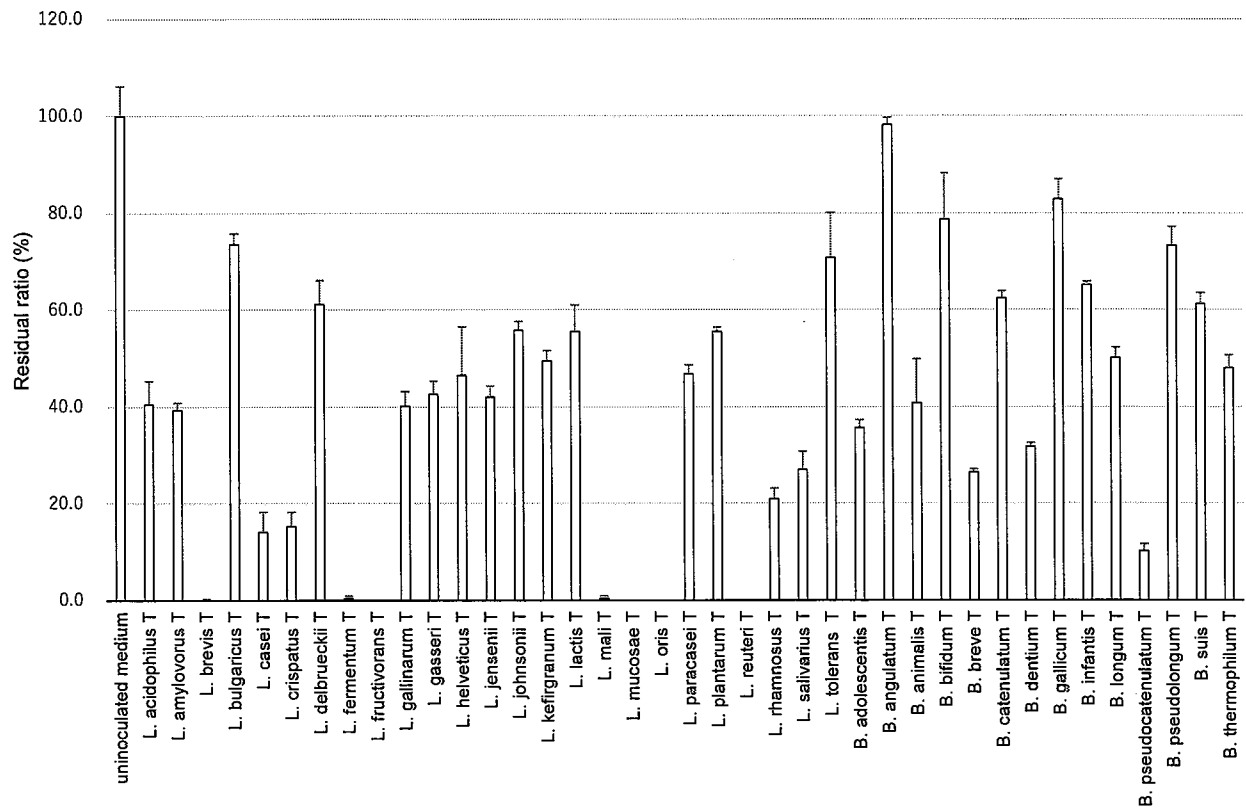


Fig. 2

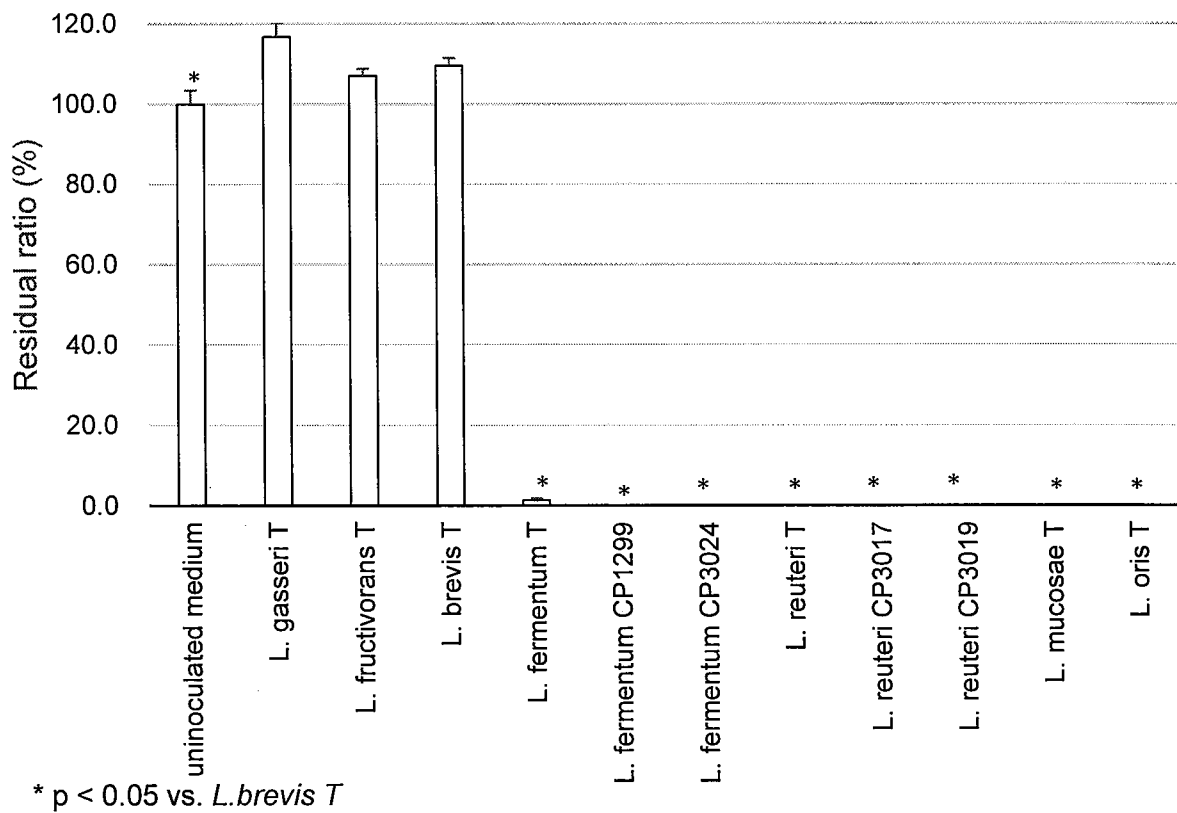
* p < 0.05 vs. *L. brevis* T

Fig. 3

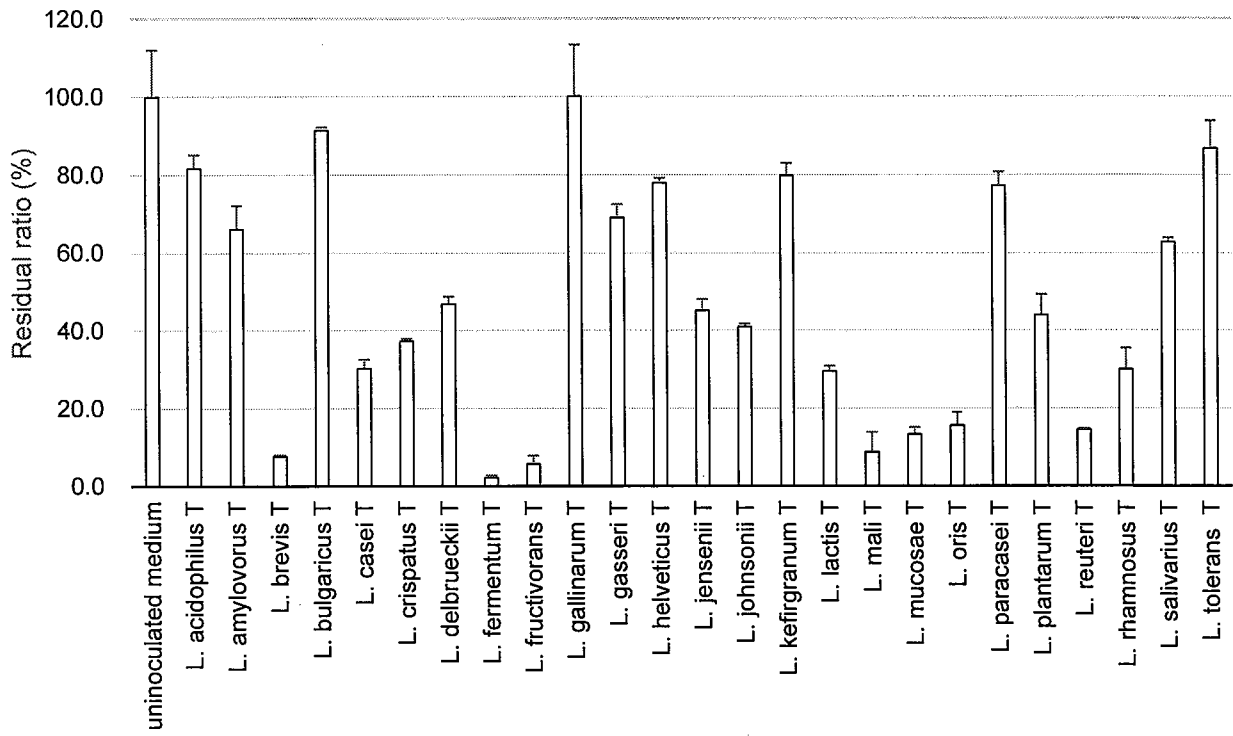
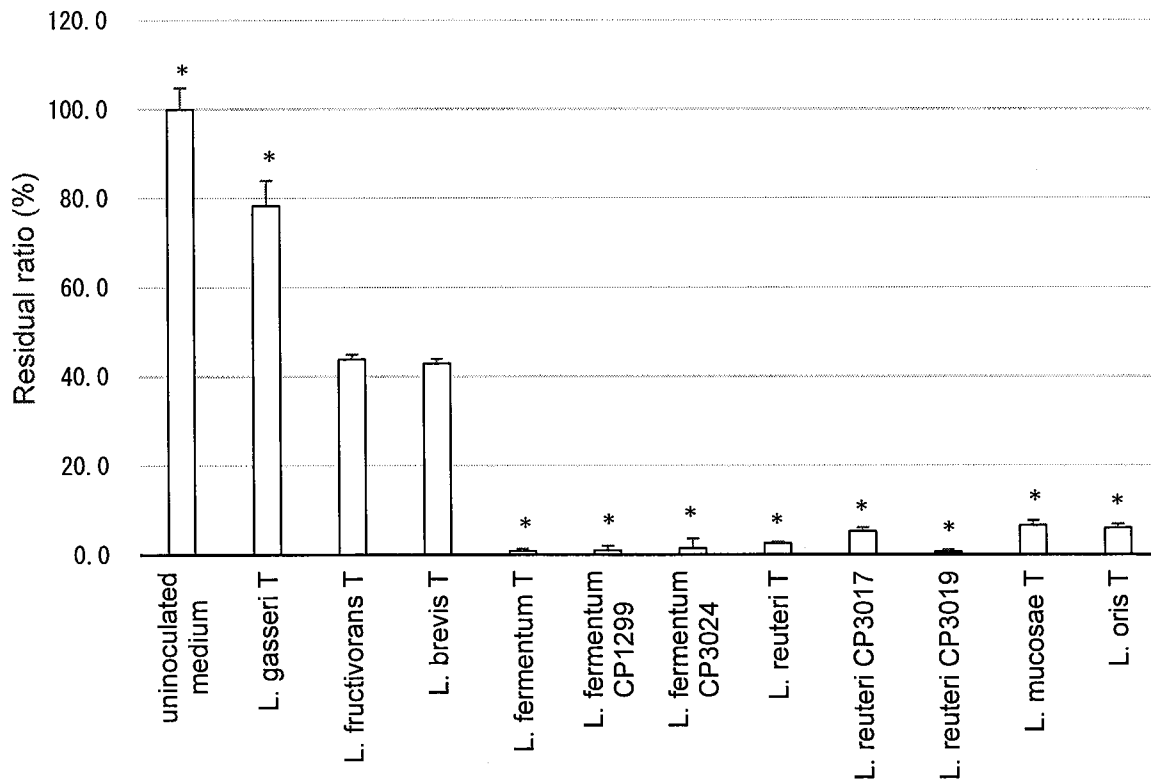


Fig. 4



* $p < 0.05$ vs. *L. brevis* T

Fig. 5

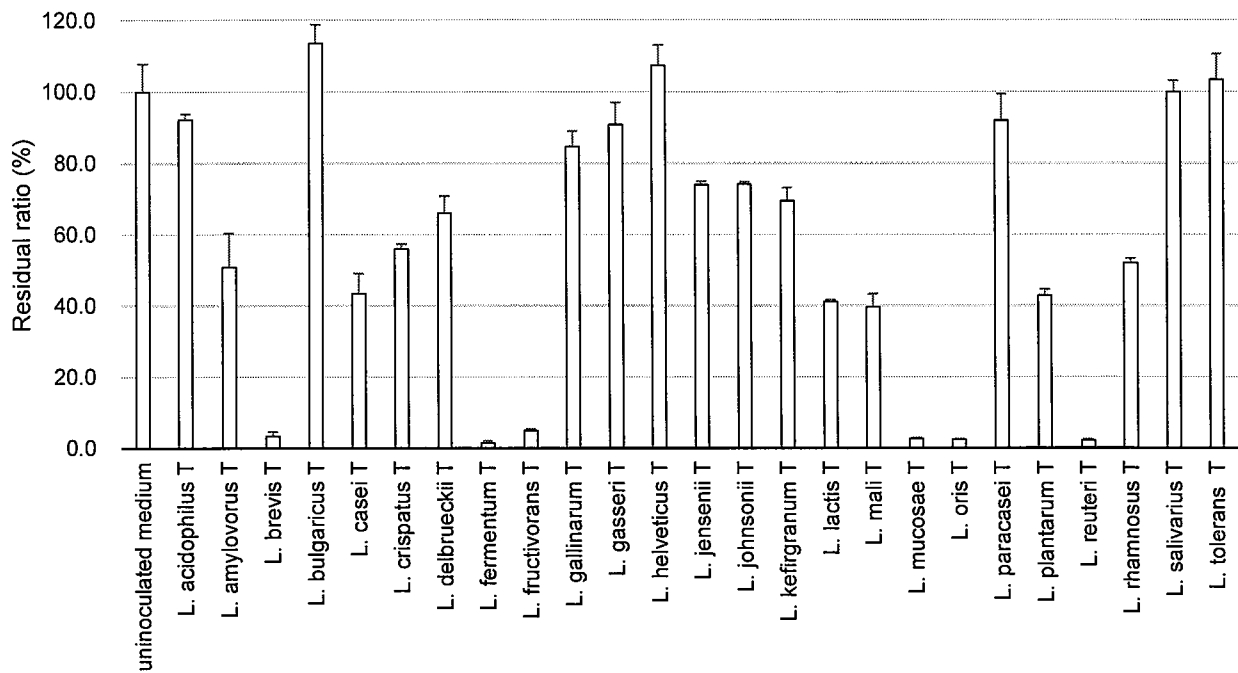
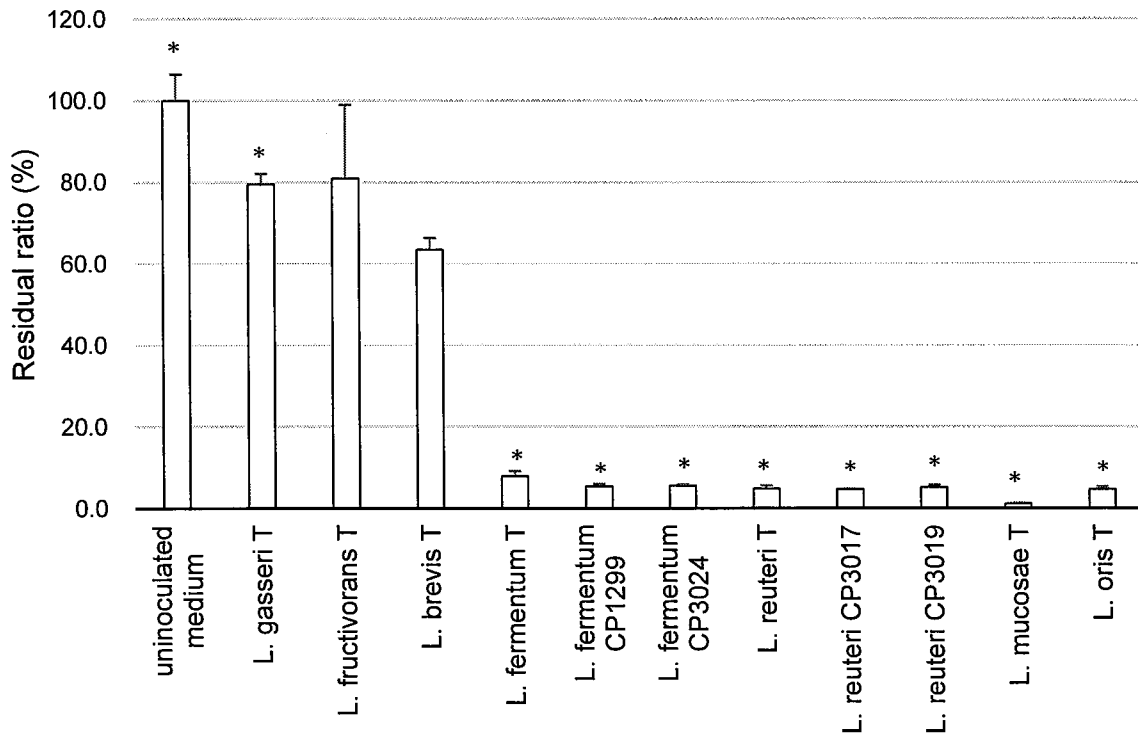


Fig. 6



* $p < 0.05$ vs. *L. brevis* T

Fig. 7

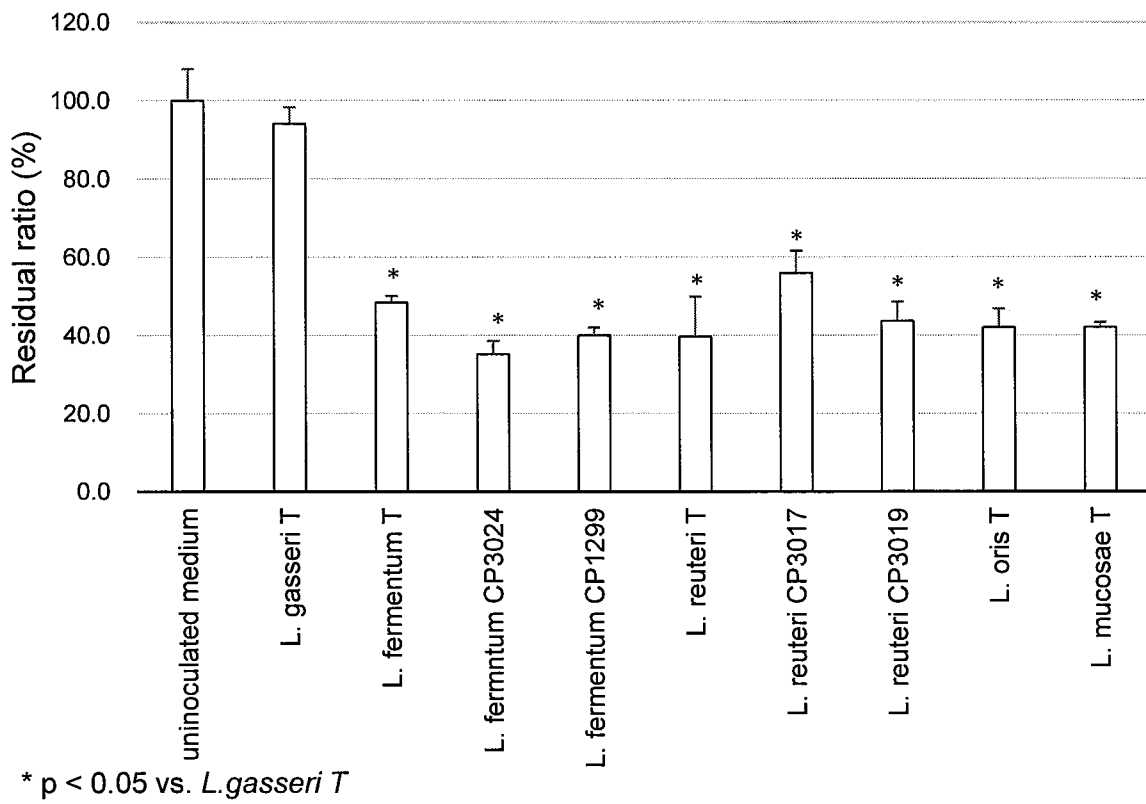


Fig. 8

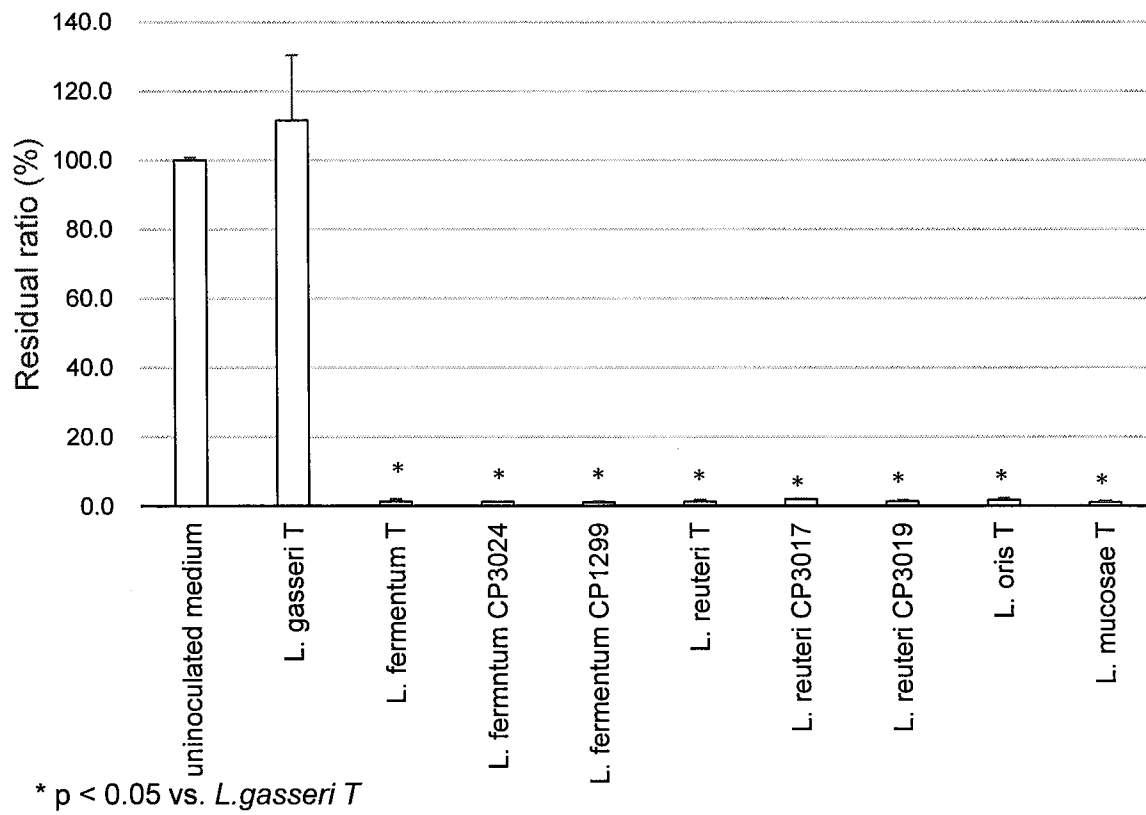
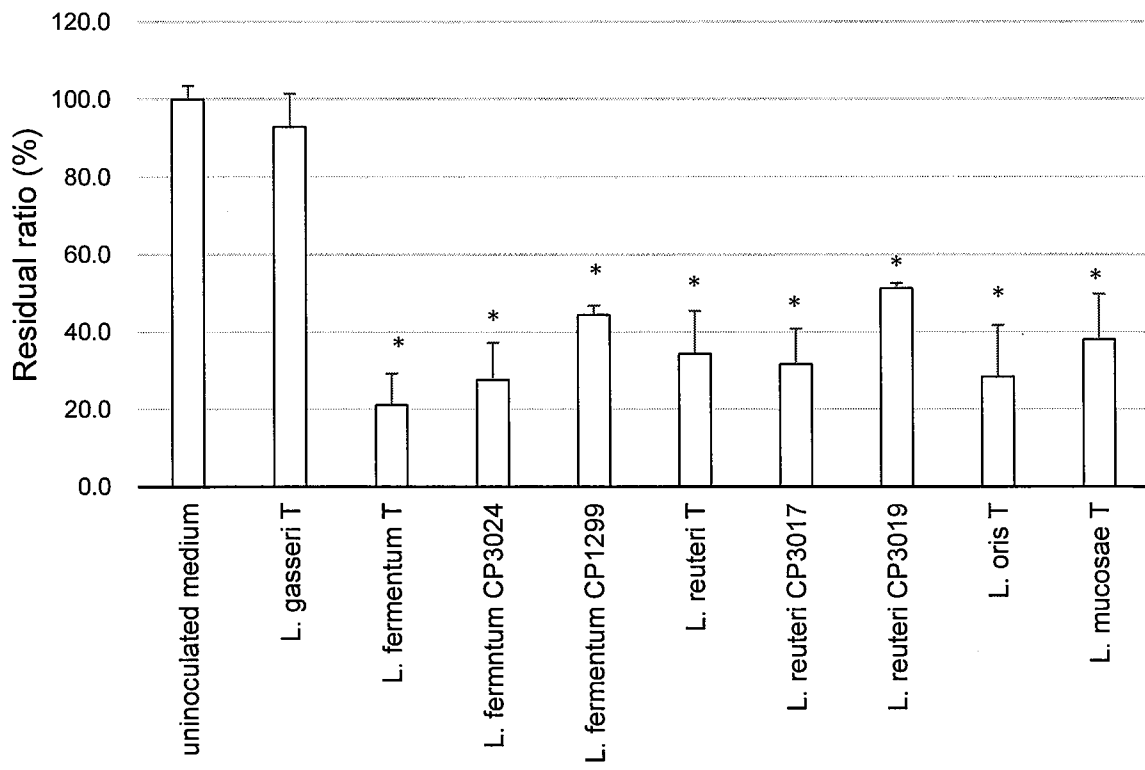
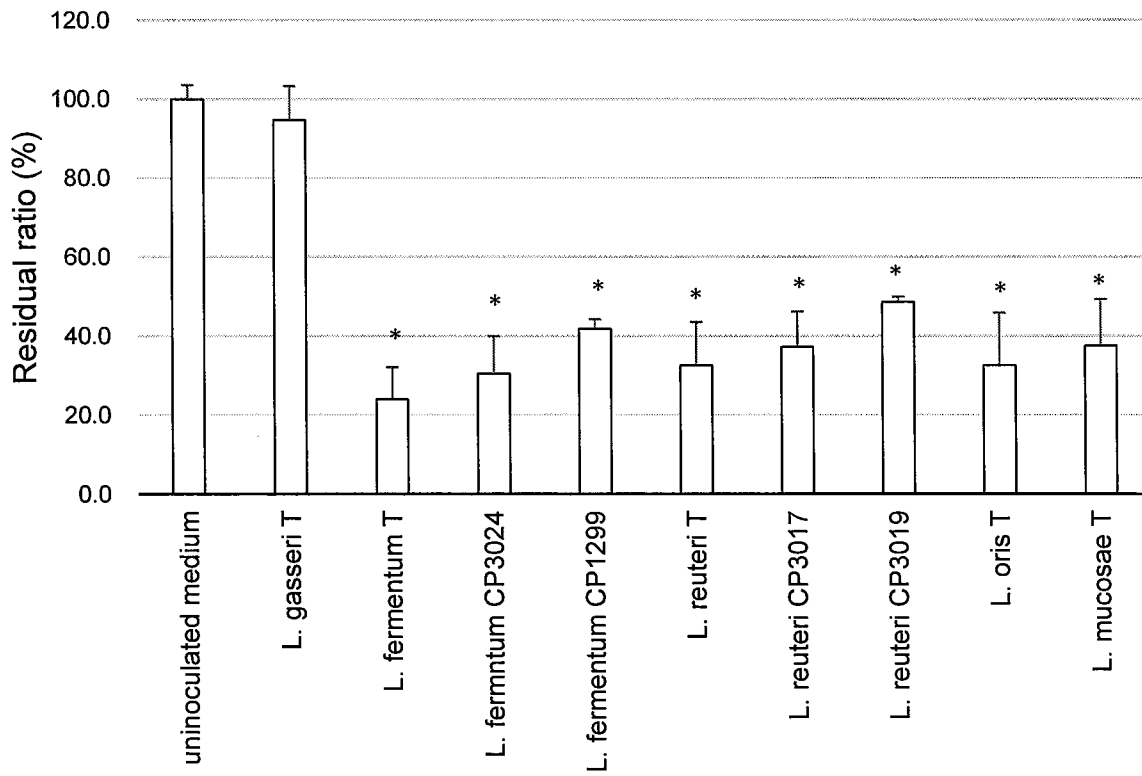


Fig. 9



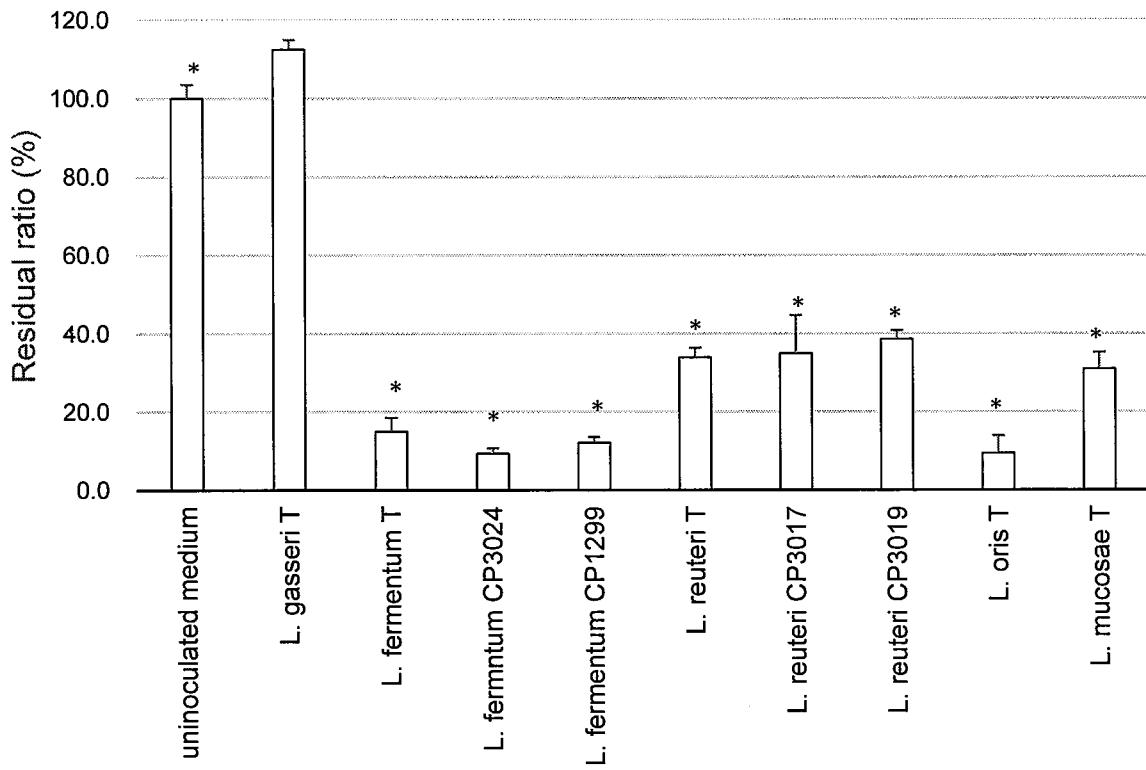
* $p < 0.05$ vs. *L. gasseri* T

Fig. 10



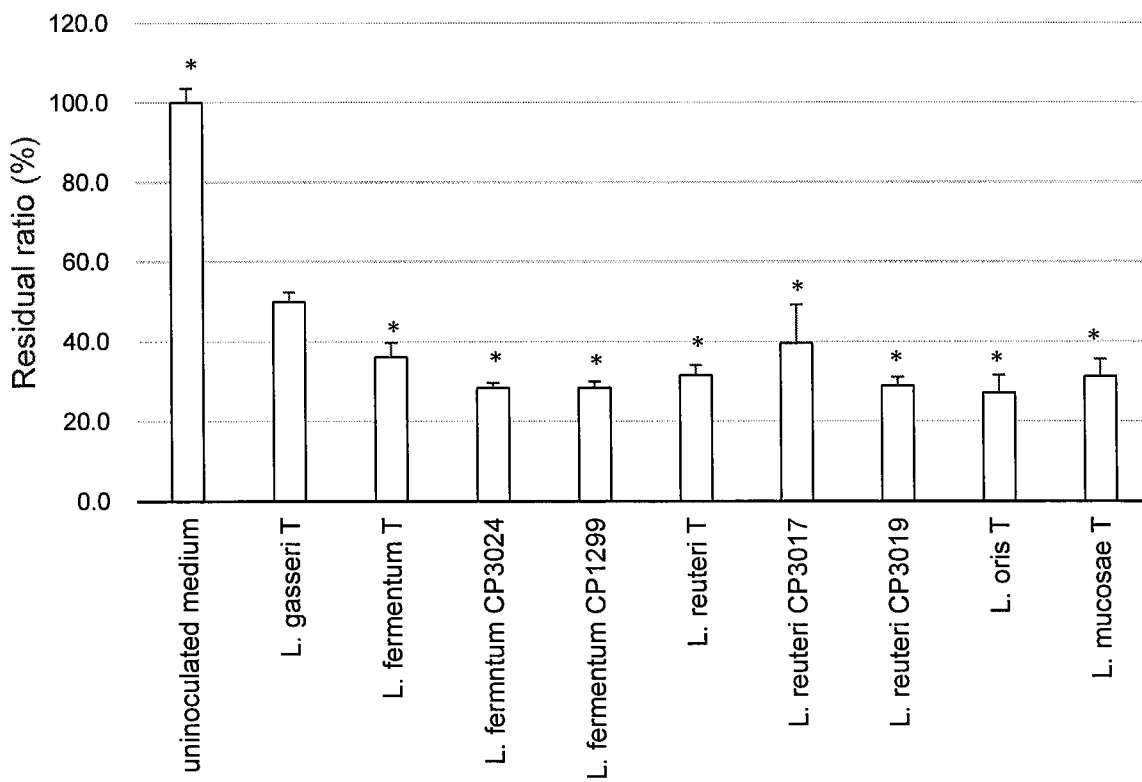
* $p < 0.05$ vs. *L. gasseri* T

Fig. 11



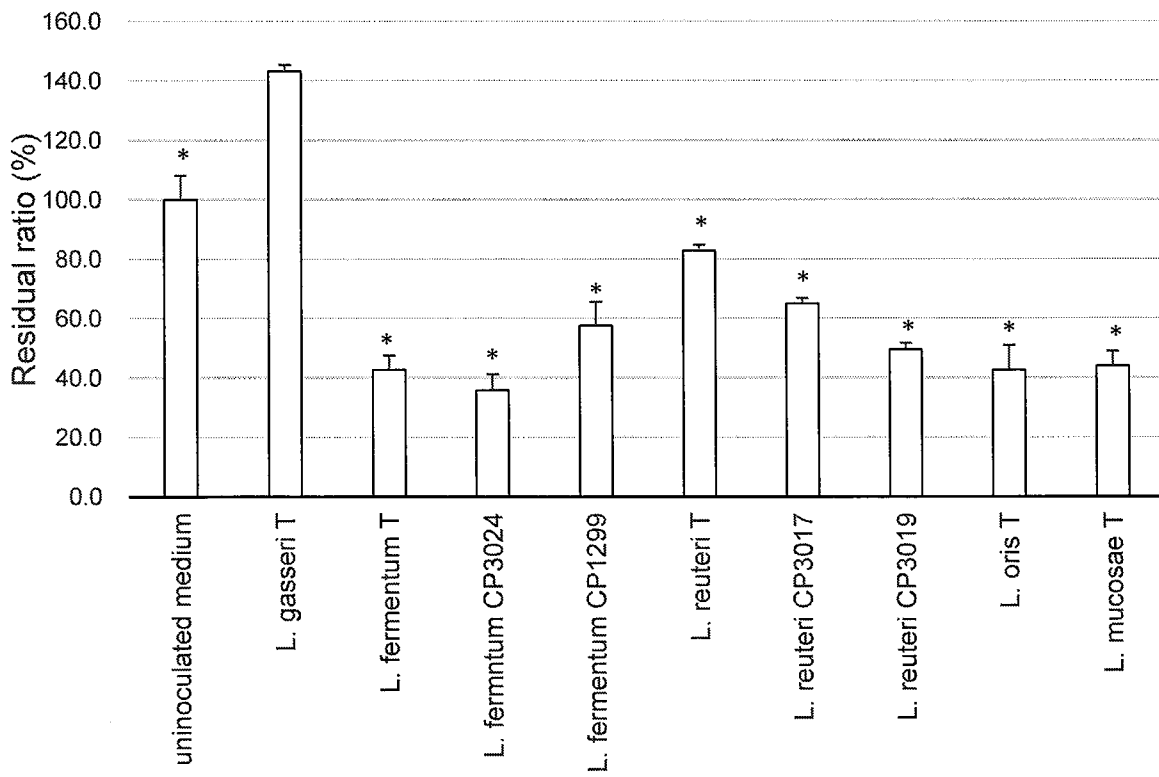
* $p < 0.05$ vs. *L. gasseri* T

Fig. 12



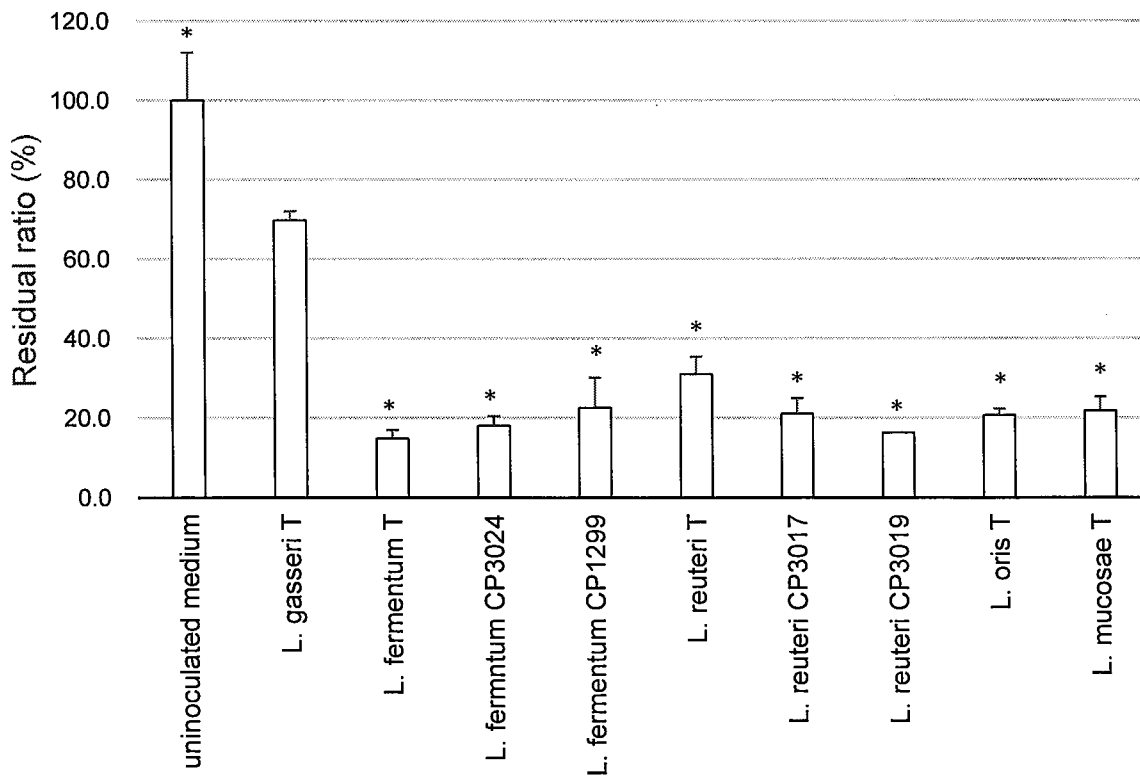
* $p < 0.05$ vs. *L. gasseri* T

Fig. 13



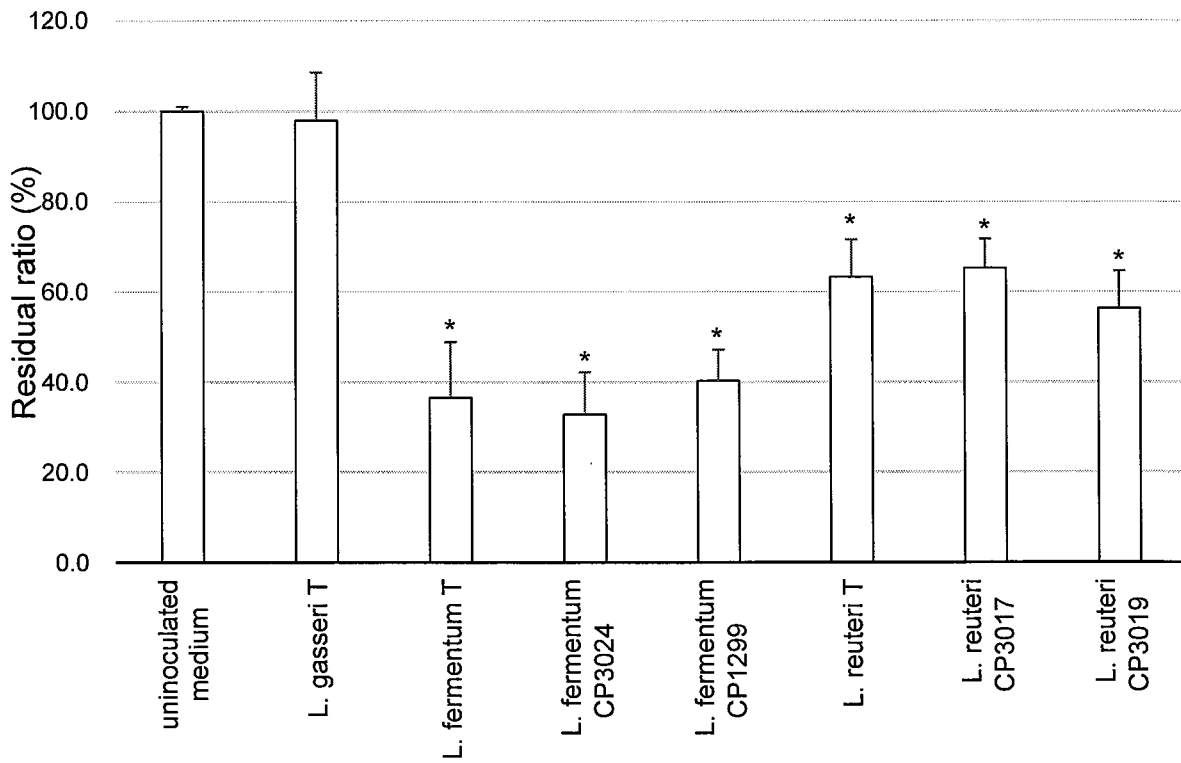
* $p < 0.05$ vs. *L. gasseri* T

Fig. 14



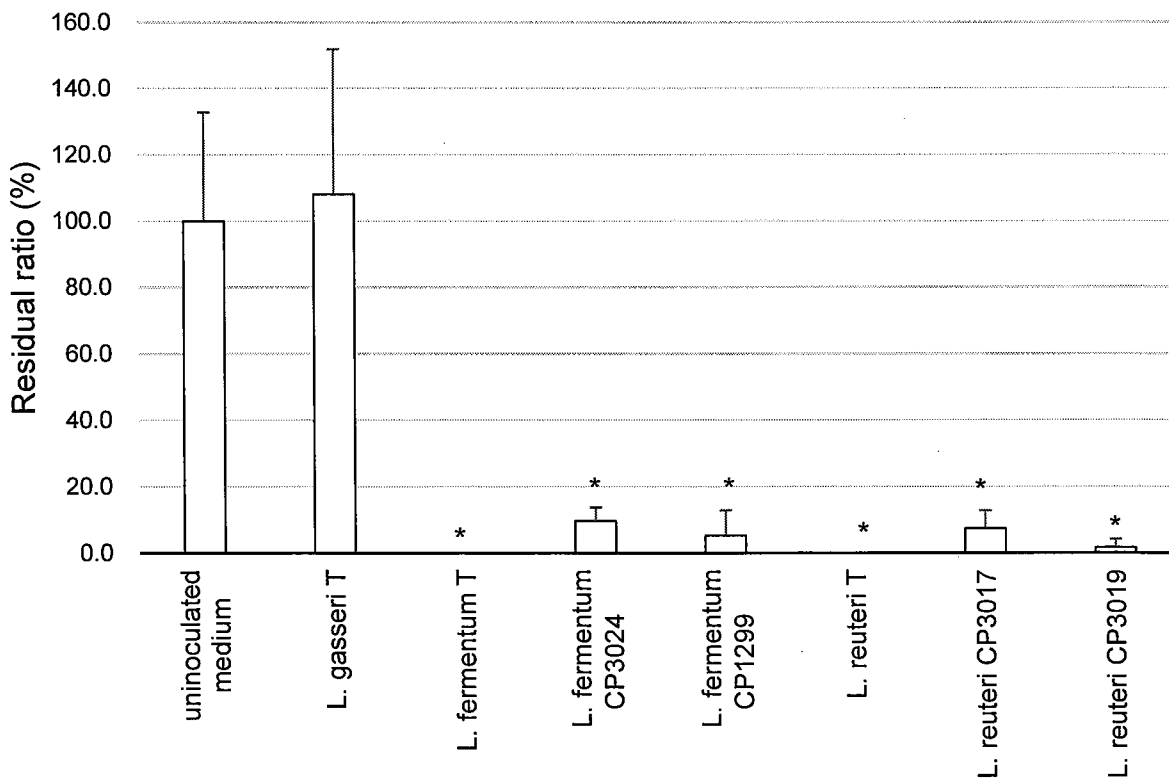
* $p < 0.05$ vs. *L. gasseri* T

Fig. 15



* $p < 0.05$ vs. *L. gasseri* T

Fig. 16



* $p < 0.05$ vs. *L. gasseri* T