

WO 2017/220525 A1

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization

International Bureau



(10) International Publication Number

WO 2017/220525 A1

(43) International Publication Date
28 December 2017 (28.12.2017)

(51) International Patent Classification:

<i>A61Q 19/00</i> (2006.01)	<i>C12R 1/225</i> (2006.01)
<i>C12R 1/25</i> (2006.01)	<i>A61P 17/00</i> (2006.01)
<i>A61K 35/747</i> (2015.01)	
<i>A61K 8/99</i> (2017.01)	

KR, KW, KZ, LA, LC, LK, LR, LS, LU, LY, MA, MD, ME, MG, MK, MN, MW, MX, MY, MZ, NA, NG, NI, NO, NZ, OM, PA, PE, PG, PH, PL, PT, QA, RO, RS, RU, RW, SA, SC, SD, SE, SG, SK, SL, SM, ST, SV, SY, TH, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, ZA, ZM, ZW.

(21) International Application Number:

PCT/EP2017/065006

(84) **Designated States** (unless otherwise indicated, for every kind of regional protection available): ARIPO (BW, GH, GM, KE, LR, LS, MW, MZ, NA, RW, SD, SL, ST, SZ, TZ, UG, ZM, ZW), Eurasian (AM, AZ, BY, KG, KZ, RU, TJ, TM), European (AL, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HR, HU, IE, IS, IT, LT, LU, LV, MC, MK, MT, NL, NO, PL, PT, RO, RS, SE, SI, SK, SM, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, KM, ML, MR, NE, SN, TD, TG).

(22) International Filing Date:

20 June 2017 (20.06.2017)

(25) Filing Language:

English

(26) Publication Language:

English

(30) Priority Data:

2016/5454 21 June 2016 (21.06.2016) BE

(71) **Applicants:** YUN NV [BE/BE]; Berkenlaan 4, 2630 Aartselaar (BE). UNIVERSITEIT ANTWERPEN [BE/BE]; Prinsstraat 13, 2000 Antwerpen (BE).

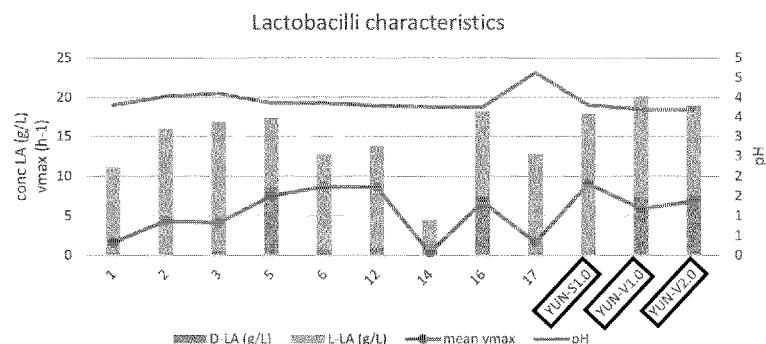
(72) **Inventors:** LEBEER, Sarah; Groenenborgerlaan 171, 2020 Antwerpen (BE). CLAES, Ingmar; Groenenborgerlaan 171, 2020 Antwerpen (BE). OERLEMANS, Eline; Groenenborgerlaan 171, 2020 Antwerpen (BE). VAN DEN BROEK, Marianne; Groenenborgerlaan 171, 2020 Antwerpen (BE).

(74) **Agent:** LC PATENTS; Kempische Steenweg 542A, 3500 Hasselt (BE).

(81) **Designated States** (unless otherwise indicated, for every kind of national protection available): AE, AG, AL, AM, AO, AT, AU, AZ, BA, BB, BG, BH, BN, BR, BW, BY, BZ, CA, CH, CL, CN, CO, CR, CU, CZ, DE, DJ, DK, DM, DO, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, GT, HN, HR, HU, ID, IL, IN, IR, IS, JO, JP, KE, KG, KH, KN, KP,

(54) **Title:** DERMATOLOGICAL PREPARATIONS FOR MAINTAINING AND/OR RESTORING HEALTHY SKIN MICROBIO-
TA

Fig. 1



(57) **Abstract:** The present invention is directed to the direct application of beneficial or probiotic bacteria to the skin for maintenance of a healthy skin microbiota and to help restore an unbalanced skin microbiota. The application is based on the use of selected Lactobacillus strains as anti-5 pathogenic agents, in particular *L. plantarum*, *L. pentosus* and/or *L. rhamnosus*, against common skin pathogens, whereby produced acids such as lactic acid are important antimicrobial factors.

**DERMATOLOGICAL PREPARATIONS FOR MAINTAINING AND/OR RESTORING
HEALTHY SKIN MICROBIOTA**

FIELD OF THE INVENTION

5 The present invention is directed to the direct topical application of beneficial or probiotic bacteria to the skin for maintenance of a healthy skin microbiota and to help restore an unbalanced skin microbiota. This restoration of a healthy microbiota falls under the term probiotherapy, defined as the use of beneficial micro-organisms or probiotics to restore a healthy microbiota at a site where microbial dysbiosis occurs. The application is based on the
10 use of selected *Lactobacillus* strains as anti-pathogenic agents, in particular *L. plantarum*, *L. pentosus* and/or *L. rhamnosus*, against common skin pathogens, whereby produced acids such as lactic acid are important antimicrobial factors.

BACKGROUND TO THE INVENTION

15 Hence, it was an object of the present invention to provide a solution for subjects suffering from skin conditions due to an aberrant microbial balance on the skin. Thereto, it was found that the topical use of *L. plantarum*, *L. pentosus* and/or *L. rhamnosus* species on the skin is very effective in restoring and/or maintaining a healthy skin microbiota, and is thus very suitable in
20 relieving skin conditions in subjects in need thereof.

Oral formulations comprising *Lactobacillus* strains have been used before in the treatment of skin conditions like atopic dermatitis. However, oral administration versus direct topical administration are different administration routes and each have a completely different
25 underlying mechanism. In oral administration, in particular a beneficial effect on the general health via immuno-stimulation is intended, whereas by direct dermatological (skin) administration, competition with 'unwanted' microorganisms occurs.

Like the gastrointestinal tract, our skin harbours a unique microbial ecosystem. The type of
30 micro-organisms found on the skin depends on a combination of host factors, environmental factors but also topographical location. The role of this microbiota in skin disorders is still not completely unravelled. However, it seems that, at least, some skin disorders are linked to a disturbed microbiota as antimicrobial treatments can improve clinical symptoms (Grice & Segre 2011). For example, in acne vulgaris a correlation has been found with the presence of
35 *Propionibacterium acnes* (Beylot et al. 2014). Although acne vulgaris is a multifactorial condition and is, among other factors, influenced by hormonal factors, these *P. acnes* bacteria seem to induce inflammation resulting in inflamed pimples also called papules or pustules. As *P. acnes* is also found on a healthy skin not causing acne, this suggests that other factors are involved, tipping the balance of the composition of the skin microbiota towards an overgrowth of this
40 bacteria.

- 2 -

Another example of a skin disorder where the microbiota seems to be important is dandruff (Wang et al. 2015; Sugita et al. 2015; Grice & Segre 2011). In people with dandruff, the fungus *Malassezia* is often overrepresented. Indications that it is this fungus that is a possible cause of the condition, come from the fact that antimycotic treatment improves the symptoms. In 5 contrast, antibacterial therapies do not improve dandruff. Again, other factors are expected to be involved in this skin disorder but the correlation with *Malassezia* is intriguing.

Similarly as for dandruff, fungal skin infections with *Candida albicans* or dermatophytes, like *Trichophyton* spp., seem to be skin disorders linked to a dysbiosis in the skin microbiota as 10 these species are also present on healthy subjects. In the case of *Tinea pedis* or 'athlete's foot' overgrowth of *Trichophyton rubrum* or *T. mentagrophytes* is often observed.

The production of lactic acid in combination with possibly other antimicrobial compounds like 15 bacteriocins seems to give protection against aforementioned infections and dysbiotic conditions and lactic acid seems to be active against bacterial, fungal and even viral pathogens. It is for this reason that lactobacilli are considered to be important in the homeostasis of the dynamical dermatological ecosystem. Potential health promoting mechanisms of lactobacilli are 20 i) to preserve a healthy skin pH (+/- 5.5), mainly by production of lactic acid; ii) production of antimicrobial compounds and competitive exclusion of pathogens; iii) modulation of immune response and iv) strengthening of the epithelial barrier.

Hence, it was an object of the present invention to provide a solution for subjects suffering from 25 dermatological conditions due to an aberrant microbial balance of the skin. Thereto, it was found that the topical dermatological use of *L. plantarum*, *L. pentosus* and/or *L. rhamnosus* species is very effective in restoring and/or maintaining a healthy microbiota on the skin, and is thus very suitable in relieving dermatological conditions in subjects in need thereof.

Oral formulations comprising *Lactobacillus* strains have been used before in the treatment of 30 dermatological disorders. However, oral administration versus direct topical administration are different administration routes and each have a completely different underlying mechanism. In oral administration, in particular a beneficial effect on the general health via immuno-stimulation is intended, whereas by direct administration on the skin, competition with 'unwanted' microorganisms occur.

35

40

SUMMARY OF THE INVENTION

In a first aspect, the present invention provides a topical skin composition comprising one or more live *Lactobacillus* species; wherein at least one of said *Lactobacillus* species is *L. plantarum*; more in particular a *L. plantarum* strain having at least 97% sequence similarity with 5 SEQ ID N° 4 in its 16S rRNA gene.

In a further aspect, the present invention provides a live *Lactobacillus* species for use in restoring and/or maintaining a healthy skin microbiota, by topical route, said *Lactobacillus* species being *L. plantarum*; more in particular a *L. plantarum* strain having at least 97% 10 sequence similarity with SEQ ID N° 4 in its 16S rRNA gene.

In yet a further aspect, the present invention provides the use of one or more live *Lactobacillus* species, in the preparation of a topical skin composition for restoring and/or maintaining a healthy skin microbiota; wherein at least one of said *Lactobacillus* species is *L. plantarum*; more 15 in particular a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene.

The present invention also provides a method for restoring and/or maintaining a healthy skin microbiota; comprising at least one step of administering by topical route, to an individual, an 20 effective amount of one or more live *Lactobacillus* species; wherein at least one of said *Lactobacillus* species is *L. plantarum*; more in particular a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene.

In yet another aspect, the present invention provides a composition comprising one or more live 25 *Lactobacillus* species for use in restoring and/or maintaining a healthy skin microbiota, by topical route, said *Lactobacillus* species being selected from the list comprising *L. plantarum*, *L. pentosus* and *L. rhamnosus*; more in particular a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene, a *L. pentosus* strain having at least 97% sequence similarity with SEQ ID N° 1 in its 16S rRNA gene and a *L. rhamnosus* strain 30 having at least 97% sequence similarity with SEQ ID N° 5 in its 16S rRNA gene.

The present invention further provides a *Lactobacillus* strain being *L. rhamnosus* YUN-S1.0 deposited under accession number LMG P-29611 (deposited at BCCM on May, 12 2016).

35 In a particular aspect, the present invention provides a composition comprising one or more *Lactobacillus* strains as defined herein above.

In a particular embodiment, the composition of the present invention is a topical skin composition, more in particular in the form of a gel, cream, foam, lotion or ointment.

- 4 -

In another particular embodiment, the present invention provides the *Lactobacillus* strain as defined herein above or the compositions as defined herein above; for use in restoring and/or maintaining a healthy skin microbiota, by topical route.

5 In a particular aspect, the present invention provides a topical use of one or more live *Lactobacillus* species in probiotherapy of the skin; wherein said *Lactobacillus* species are selected from the list comprising *L. plantarum*, *L. pentosus* and *L. rhamnosus*; more in particular, said probiotherapy consists of restoring and/or maintaining a healthy skin microbiota in a subject in need thereof.

10 In another particular embodiment, said *Lactobacillus* species in the topical uses, methods and compositions as disclosed herein, is a *Lactobacillus* strain selected from the list comprising *L. plantarum* YUN-V2.0 deposited under accession number LMG P-29456 (deposited at BCCM on Mar, 09 2016), *L. pentosus* YUN-V1.0 deposited under accession number LMG P-29455 15 (deposited at BCCM on Mar, 09 2016); and *L. rhamnosus* YUN-S1.0 deposited under accession number LMG P-29611 (deposited at BCCM on May, 12 2016).

BRIEF DESCRIPTION OF THE DRAWINGS

With specific reference now to the figures, it is stressed that the particulars shown are by way of example and for purposes of illustrative discussion of the different embodiments of the present invention only. They are presented in the cause of providing what is believed to be the most useful and readily description of the principles and conceptual aspects of the invention. In this regard no attempt is made to show structural details of the invention in more detail than is necessary for a fundamental understanding of the invention. The description taken with the drawings making apparent to those skilled in the art how the several forms of the invention may be embodied in practice.

Fig. 1: Characteristics of lactobacilli in reference to growth, production of D- and L-lactic acid (LA) and lowering of the pH of the medium.

30 **Fig. 2:** Time course experiment for the analysis of the antipathogenic effect of spent culture supernatant of lactobacilli against *Propionibacterium acnes*. Growth of the bacteria (optical density at 600nm; Y-axis) is measured in time (X-axis). Each graph shows replicates of growth of *P. acnes*. It can be clearly noted that without any addition of antibiotic or SCS, *P. acnes* 35 quickly starts to grow (NC1). Similar as when erythromycin at 50 μ g/ml is added, SCS of all lactobacilli prevents growth of *P. acnes* while SCS of streptococci or staphylococci does not inhibit growth. *Erythromycine (50 μ g/ml); #Erythromycine (5 μ g/ml); $^{\$}$ Minocycline (20 μ g/ml) NC1=medium control; NC2=MRS at pH4.3; Numbers 1 to 22 = lactobacilli strains (for details see table 1); St=*Streptococcus thermophilus*; Ss=*Streptococcus salivarius*; Se=*Staphylococcus epidermidis*; T0.5=0.5% Tween 80; T1=1% Tween 80.

DETAILED DESCRIPTION OF THE INVENTION

The present invention is based on the discovery of specific *Lactobacillus* strains that can compete with growth of *Propionibacterium acnes*, *Candida albicans*, *Malassezia spp.*, 5 *Trichophyton spp.* and bacteria or fungi that are linked with skin conditions like acne vulgaris, dandruff, tinea pedis or other fungal skin infections. These selected strains are herein generally termed "YUN" strains and are capable of competing with skin pathogens and thereby restore a healthy skin microbiota. This restoration of a healthy microbiota falls under the term probiotherapy, defined as the use of beneficial micro-organisms or probiotics to restore a 10 healthy microbiota at a site where microbial dysbiosis occurs.

Hence, in a first aspect, the present invention provides a topical skin composition comprising one or more live *Lactobacillus* species; wherein at least one of said *Lactobacillus* species is *L. plantarum*; more in particular a *L. plantarum* strain having at least 97% sequence similarity with 15 SEQ ID N° 4 in its 16S rRNA gene.

Said composition according to the present invention may comprise further *Lactobacillus* species such as for example selected from the non-limiting list comprising *L. pentosus*, *L. gasseri*, *L. crispatus*, *L. acidophilus*, *L. jensenii*, *L. fermentum*, *L. rhamnosus*. 20

In the context of the present invention, the term "topical" is meant to be the local delivery at a specified location of the body, in particular the application to a particular place on the body. In particular, it includes the application via non-solid formulations such as creams, foams, gels, lotions or ointments. The term "topical" is not meant to include the delivery in the form of solid 25 preparations such as capsules, tablets, ...

Hence, the term "topical skin" is meant to include the local delivery using non-solid formulations directly onto the skin of the body. Preferably, the compositions according to the present invention are applied over a large area of the skin in order to be most effective. 30

In the context of the present invention the term "live *Lactobacillus* species" is meant to be viable *Lactobacillus* species, and is not meant to be fragments, culture supernatants, or killed forms thereof.

35 In a further aspect, the present invention provides a live *Lactobacillus* species for use in probiotherapy of the skin, by topical route, said *Lactobacillus* species being *L. plantarum*; more in particular a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene. As already defined herein above, said probiotherapy is meant to be the restoration and/or maintainance of a healthy skin microbiota in a subject in need thereof.

Subjects that may benefit from such probiotherapy are for example people/persons with a skin conditions linked to a disturbed skin microbiota possibly due to bacterial or yeast infections and/or any dysbiosis caused by overgrowth of specific pathogenic micro-organisms, like *acne vulgaris*, *tinea pedis*, *dandruff*, *rosaceae*, *impetigo*,...

5

Hence, in a further aspect, the present invention provides the use of one or more live *Lactobacillus* species, in the preparation of a topical skin composition for restoring and/or maintaining a healthy skin microbiota; wherein at least one of said *Lactobacillus* species is *L. plantarum*; more in particular a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene.

10 The present invention also provides a method for restoring and/or maintaining a healthy skin microbiota; comprising at least one step of administering by topical route, to an individual, an effective amount of one or more live *Lactobacillus* species; wherein at least one of said 15 *Lactobacillus* species is *L. plantarum*; more in particular a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene.

20 In yet another aspect, the present invention provides a composition comprising one or more live *Lactobacillus* species for use in restoring and/or maintaining a healthy skin microbiota, by topical route, said *Lactobacillus* species being selected from the list comprising *L. plantarum*, *L. pentosus* and *L. rhamnosus*; more in particular a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene; a *L. pentosus* strain having at least 97% sequence similarity with SEQ ID N° 1 in its 16S rRNA gene and a *L. rhamnosus* strain having at least 97% sequence similarity with SEQ ID N° 5 in its 16S rRNA gene.

25

30 The present invention further provides a *Lactobacillus* strain selected from the list comprising *L. pentosus* YUN-V1.0 deposited under accession number LMG P-29455 (deposited at BCCM on Mar, 09 2016); *L. plantarum* YUN-V2.0 deposited under accession number LMG P-29456 (deposited at BCCM on Mar, 09 2016); and *L. rhamnosus* YUN-S1.0 deposited under accession number LMG P-29611 (deposited at BCCM on May, 12 2016)

35 The microbiological deposits mentioned herein, have been made with the BCCM/LMG Bacteria collection ("Belgian co-ordinated collections of micro-organism") with correspondence address: Laboratorium voor Microbiologie, Universiteit Gent, K.L. Ledeganckstraat 35 – 9000 Gent, Belgium

40 *Lactobacillus pentosus* YUN-V1.0 is a single colony isolate obtained in our lab after subculturing of a strain, that was originally a vaginal isolate of healthy woman. The 16S rRNA gene sequence (SEQ ID N° 1) for strain *L. pentosus* YUN-V1.0 was determined by PCR using primers 8F (5'-AGAGTTGATCCTGGCTCAG-3' – SEQ ID N° 2) and 1525R (5'-

- 7 -

AAGGAGGTGATCCAGCCGCA-3' – SEQ ID N° 3).

YUN-V2.0 and YUN-V3.0 are single colony isolates obtained in our lab after subculturing of *Lactobacillus plantarum* strains that were originally isolated from human saliva and a maize silage respectively. The 16S rRNA gene sequence (SEQ ID N° 4) for strain *L. plantarum* YUN-V2.0 was determined by PCR using primers 8F (5'-AGAGTTGATCCTGGCTCAG-3' – SEQ ID N° 2) and 1525R (5'-AAGGAGGTGATCCAGCCGCA-3' – SEQ ID N° 3).

YUN-S1.0 is a single colony isolate obtained in our lab after subculturing of a *Lactobacillus rhamnosus* strain that was originally isolated from a healthy person. The 16S rRNA gene sequence (SEQ ID N° 5) for strain *L. rhamnosus* YUN-S1.0 was determined by PCR using primers 8F (5'-AGAGTTGATCCTGGCTCAG-3' – SEQ ID N° 2) and 1525R (5'-AAGGAGGTGATCCAGCCGCA-3' – SEQ ID N° 3).

15 These particular "YUN" strains can either be used as such, or are preferably formulated in a composition comprising such strains. Said compositions are topical skin compositions more in particular in the form of non-solid formulations such as creams, foams, gels, lotions or ointments.

20 In particular, the present invention provides the above defined "YUN" strains for use in probiotherapy of the skin, i.e. for restoring and/or maintaining a healthy skin microbiota.

In yet a further aspect, the present invention provides a topical use of one or more live *Lactobacillus* species in probiotherapy of the skin; wherein said *Lactobacillus* species are 25 selected from the list comprising *L. plantarum*, *L. pentosus* and *L. rhamnosus*; more in particular, said probiotherapy consists of restoring and/or maintaining a healthy skin microbiota in a subject in need thereof.

30 In a specific embodiment, the *Lactobacillus* species in the topical uses, methods and compositions as disclosed herein, is a *Lactobacillus* strain selected from the list comprising *L. plantarum* YUN-V2.0 deposited under accession number LMG P-29456 (deposited at BCCM on Mar, 09 2016); *L. pentosus* YUN-V1.0 deposited under accession number LMG P-29455 (deposited at BCCM on Mar, 09 2016); and *L. rhamnosus* YUN-S1.0 deposited under accession number LMG P-29611 (deposited at BCCM on May, 12 2016).

EXAMPLES**MATERIALS AND METHODS**

5

Bacterial strains and growth conditions

Lactobacillus strains (Table 1) were grown at 37°C in de Man, Rogosa and Sharpe (MRS) medium (Carl Roth). All bacteria were grown in non-shaking conditions and inoculated from glycerol stocks (-80°C). Solid media contained 1.5% (w/v) agar.

10

Table 1: Bacterial strains used in this research

Species	#	Strain	Relevant genotype or description	Reference and/or Source
LACTOBACILLI				
<i>Lactobacillus casei</i>	1	ATCC334	Single colony isolate obtained in our lab from a stock culture of ATCC334	ATCC
<i>Lactobacillus casei</i>	2	DN-114001	Single colony isolate obtained in our lab from a commercially available fermented drink (Actimel®) containing <i>L. casei</i> DN-114001, confirmed by sequencing	Commercial probiotic product
<i>Lactobacillus casei</i>	3	Shirota	Single colony isolate obtained in our lab from a commercially available fermented drink containing <i>L. casei</i> Shirota (Yakult®), confirmed by sequencing	Commercial probiotic product
<i>Lactobacillus pentosus</i>	4	YUN-V1.0	Single colony isolate	
<i>Lactobacillus plantarum</i>	5	LMG1284	Single colony isolate from <i>L. plantarum</i> ATCC8014 or LMG1284	ATCC
<i>Lactobacillus reuteri</i>	6	RC-14	Single colony isolate obtained in our lab from a commercially available probiotic supplement containing <i>L. reuteri</i> RC-14, confirmed by sequencing	Commercial probiotic product
<i>Lactobacillus rhamnosus</i>	7	YUN-S1.0	Clinical isolate	
<i>Lactobacillus rhamnosus</i>	12	GR-1	Single colony isolate obtained in our lab from a commercially available probiotic supplement containing <i>L. rhamnosus</i> GR-1	(Chan et al. 1984; 1985; Reid 1999; Reid & Bruce 2001), ATCC
<i>Lactobacillus helveticus</i>	14	AMB-2	single colony isolate	Commercial probiotic product
<i>Lactobacillus plantarum</i>	15	YUN-V2.0	Single colony isolate	
<i>Lactobacillus plantarum</i>	16	5057	Single colony isolate	
<i>Lactobacillus paracasei</i>	17	LMG12586	Single colony isolate obtained in our lab from a stock culture of LMG12586	BCCM/LMG
<i>Lactobacillus plantarum</i>	22	/	Single colony isolate	
<i>Lactobacillus pentosus</i>	25	LMG8041	Single colony isolate	BCCM/LMG

PATHOGENS				
<i>Trichophyton rubrum</i>	2	/	Clinical isolate	BCCM/LMG
<i>Malassezia furfur</i>		/	Clinical isolate	BCCM/LMG
<i>Candida albicans</i>	/	/	Clinical isolate	

Preparation of spent culture supernatant (SCS) of selected strains

To obtain spent culture supernatant (SCS) containing the secreted active antimicrobial products, growth medium specific for each species was inoculated from a preculture and 5 incubated for 24h. SCS was obtained by centrifugation for 30 min. at 6797 g (8000 rpm) at 4°C. Afterwards, the SCS was filter sterilized (0.20 µm cellulose acetate, VWR).

Antimicrobial activity assays for co-cultures of live lactobacilli against Malassezia furfur, Trichophyton rubrum, Propionibacterium acnes and Candida albican.

10 The antimicrobial activity of the selected bacteria was explored by standard antimicrobial tests with some minor modifications. The antimicrobial activity of the selected bacteria was explored by spot assay (Schillinger and Lücke 1989). Briefly, 1-3 µL of each culture was spotted on an agar plate. These plates were incubated for 24h up to 72h depending on the strain. Next, an overnight culture of the pathogen was diluted into 7 mL of soft agar of the medium of the 15 pathogen and poured over the plates with the spots of the selected strains. The plates were incubated overnight at 30-37°C, after which the inhibition zones were measured. A spot of miconazole (for fungi) and/or 0.1% hexetidine and/or tetracycline (for Propionibacterium acnes) was added to the spot plate as positive control before the soft agar was poured.

20 ***Radial diffusion test of SCS of lactobacilli***

In addition, the antimicrobial activity of spent culture supernatant (SCS) was investigated with a 25 protocol as previously described for the competition assays between lactobacilli and gastrointestinal pathogens (Coconnier et al. 1997). Miconazol (for fungi) and tetracycline (for Propionibacterium acnes) was used as a positive control. Sterile growth medium was used as a negative control.

Time course analysis of the antimicrobial activity of SCS of the selected strains against Candida, Propionibacterium acnes, Malassezia furfur and Trichophyton spp (further referred to as 'pathogens').

30 The time course analysis was performed similarly as described previously (De Keersmaecker et al. 2006) with minor modifications. Briefly, an overnight culture of the pathogen was added to the wells of a microplate filled with 50-80% the proper medium supplemented with 50-5% SCS of lactobacilli. MRS at pH 4,3 and antibiotics or antimycotics at the proper concentration were used as a negative and a positive control, respectively. Bacteria or fungi were grown and the 35 optical density (OD) was measured at 590 nm each 30 min during 3 days using a Synergy HTX

- 10 -

multi-mode reader (Biotek). Each test was measured at least in triplicate and the average OD was calculated. The antimicrobial activity was expressed as the relative optical density reached after 24h (stationary phase) compared to the negative controls.

5 ***Antibiotic susceptibility***

Antibiotic sensitivity was evaluated using the Kirby-Bauer disc diffusion test. In short, antibiotics were spotted on paper discs and the bacterial inhibition zone was measured on agar plates. The antibiotics tested were erythromycin, normocin, tetracyclin, ampicillin and clindamycin at relevant concentrations.

10

Proof-of-concept human clinical trial in patients with acne vulgaris

15

A proof-of-concept clinical trial was performed on 20 patients with acne vulgaris. Patients were men between 12-25 years with mild inflammatory acne. The aim of this proof-of-concept trial was to assess the impact of a topical probiotic cream (containing +10-8 colony forming units (CFU) of L. pentosus YUN-V1.0, +10-8 CFU of L. plantarum YUN-V2.0 and +10-8 CFU L. rhamnosus YUN-S1.0 per application of 1g of the topical cream ACN) on the skin microbiota and on the acne severity. Patients were asked to apply the cream twice daily for 56 days (8 weeks). The patients were seen by a dermatologist at start (before the therapy), week 4, week 8 and week 10. A skin swab was taken at each visit. Bacterial DNA was isolated from these samples by the commercial MoBio Powersoil kit (cfr. Human Microbiome Project). Isolated DNA was analysed via 16S rRNA amplicon sequencing with MiSeq Illumina and a bio-informatical analysis was performed. Moreover, a clinical scoring was performed and a photograph taken at each visit.

20

Proof-of-concept human clinical trial in patients with tinea pedis (Athlete's foot)

25

A proof-of-concept clinical trial was performed on 20 patients with tinea pedis. Patients were between 18-65 years having tinea pedis. The aim of this proof-of-concept trial was to assess the impact of a topical probiotic cream (containing +10-8 colony forming units (CFU) of L. pentosus YUN-V1.0, +10-8 CFU of L. plantarum YUN-V2.0 and +10-8 CFU L. rhamnosus YUN-S1.0 per application of 1g of the topical cream FNG) on the skin microbiota and on the Trichophyton infection. Patients were asked to apply the cream twice daily for 56 days (8 weeks). The patients were seen by a dermatologist at start (before the therapy), week 4, week 8 and week 10. A skin swab was taken at each visit. Bacterial DNA was isolated from these samples by the commercial MoBio Powersoil kit (cfr. Human Microbiome Project). Isolated DNA was analysed via 16S rRNA amplicon sequencing with MiSeq Illumina and a bio-informatical analysis was performed. For analysis of the presence of the fungi, swabs were also plated out on Trichophyton specific medium (medium suggested by BCCM). Colony PCR using universal ITS ('internal transcribed region') primers ITS1 (SEQ ID N°6) (5'-TCCGTAGGTGAAACCTGC-3') and ITS4 (SEQ ID N° 7) (5'-TCCTCCGCTTATTGATATGC-3') followed by sequencing was performed to identify the fungi. Moreover, a clinical scoring was performed and a photograph taken at each visit.

30

RESULTS***Growth characteristics and lactate production***

5 Possible beneficial or probiotic strains were characterized in terms of growth characteristics, lactate production and ability of lowering of the pH of the medium. These characteristics are expected to be important for the antipathogenic activity. These data show that *Lactobacillus pentosus* YUN-V1.0 and *L. plantarum* YUN-V2.0 and *L. rhamnosus* YUN-S1.0 produce the highest amount of lactic acid (Fig 1).

10 ***Antipathogenic activity against *Propionibacterium acnes****

Time course experiments were performed analyzing the antimicrobial activity of spent culture supernatant (SCS) of the selected strains against *Propionibacterium acnes*. SCS of all tested strains inhibited the growth of *Propionibacterium acnes* while SCS of other bacterial species like *Streptococcus thermophilus* and *S. salivarius*, both also lactic acid bacteria, and *Staphylococcus epidermidis* did not inhibit growth of *P. acnes*. This suggests species and perhaps strain specific properties of the selected lactobacilli to be important for the antipathogenic activity against *P. acnes* (Fig 2).

Antipathogenic activity against *Malassezia*, *Trichophyton* and *Candida*

20 In a next phase, the beneficial or probiotic bacteria were screened for their antipathogenic effect against specific skin pathogens. The results of a spot assays against *Malassezia furfur*, *Trichophyton rubrum* and *Candida albicans* are shown in table 1, 2 and 3 respectively.

Table 1: Spot assay of selected lactobacilli against *Malassezia furfur*.

Strain	<i>Malassezia furfur</i>		
	Exp 1	Exp 2	Exp 3
1	++	-	+
2	++	-	+
3	+	+	-
4	++	+++	++
5	+++	++	++
6	+	++	++
7	++	-	-
12	+	-	-
13	+	-	+
14	-	+	-
15	+++	+++	+++
16	++	++	++
17	-	+	-
22	+++	++	++
25	++	++	+

25 *three independent repeats are shown

Table 2: Spot assay of selected lactobacilli against *Trichophyton rubrum*.

Strain	<i>Trichophyton rubrum</i>		
	Exp 1	Exp 2	Exp 3
1	+	++	+++
2	+	++	++
3	+	++	++
4	++	++	+++
5	++	++	+++
6	-	-	+++
7	++	+++	+
12	++	+++	+++
13	++	-	-
14	+	++	++
15	+++	+++	+++
16	++	+++	+++
17	+	+++	++
22	++	+++	+++
25	++	+++	+++

*three independent repeats are shown

5 **Table 3:** Radial diffusion assay of selected lactobacilli against *Candida albicans*.

Strain	<i>Candida albicans</i>		
	Exp 1	Exp 2	Exp 3
1	-	-	-
2	+	+	+
3	+	+	+
4	++	++	++
5	+	+	+
6	-	-	-
7	+	+	++
12	+	+	+
13	/	/	/
14	+	-	-
15	+	+	++
16	+	+	+
17	-	-	-
22	/	/	/
25	/	/	/

*three independent repeats are shown

Spent culture supernatant from *L. pentosus* YUN-V1.0 and *L. plantarum* YUN-V2.0 was also tested in radial diffusion assays and demonstrated to be efficient in inhibiting *Malassezia*, 10 *Trichophyton* and *Candida* growth. *L. rhamnosus* YUN-S1.0 was not as efficient in inhibiting growth of *Malassizia* but was able to inhibit growth of *Trichophyton* and *Candida*.

Antibiotic susceptibility

The selected bacteria were also tested for their antibiotic susceptibility as to prevent spreading of antibiotic resistance genes. All lactobacilli were susceptible to erythromycin, normocin, 5 tetracyclin, ampicillin and clindamycin, except for *L. plantarum* 5057, which was susceptible to tetracyclin. For this reason, strain *L. plantarum* 5057 was found not to be suitable to use as a strain for probiotherapy.

REFERENCES

5 Beylot, C. et al., 2014. *Propionibacterium acnes*: an update on its role in the pathogenesis of acne. *Journal of the European Academy of Dermatology and Venereology : JEADV*, 28(3), pp.271–8.

10 Chan, R.C. et al., 1985. Competitive exclusion of uropathogens from human uroepithelial cells by *Lactobacillus* whole cells and cell wall fragments. *Infection and immunity*, 47(1), pp.84–9.

15 Chan, R.C., Bruce, A.W. & Reid, G., 1984. Adherence of cervical, vaginal and distal urethral normal microbial flora to human uroepithelial cells and the inhibition of adherence of gram-negative uropathogens by competitive exclusion. *The Journal of urology*, 131(3), pp.596–601.

20 Grice, E.A. & Segre, J.A., 2011. The skin microbiome. *Nature reviews. Microbiology*, 9(4), pp.244–53.

25 Reid, G., 1999. The Scientific Basis for Probiotic Strains of *Lactobacillus*. *Appl. Environ. Microbiol.*, 65(9), pp.3763–3766.

Reid, G. & Bruce, A.W., 2001. Selection of *lactobacillus* strains for urogenital probiotic applications. *The Journal of infectious diseases*, 183 Suppl , pp.S77–80.

Sugita, T. et al., 2015. Temporal changes in the skin *Malassezia* microbiota of members of the Japanese Antarctic Research Expedition (JARE): A case study in Antarctica as a pseudo-space environment. *Medical mycology*, 53(7), pp.717–24.

25 Wang, L. et al., 2015. Characterization of the major bacterial-fungal populations colonizing dandruff scalps in Shanghai, China, shows microbial disequilibrium. *Experimental dermatology*, 24(5), pp.398–400.

SEQUENCE LISTING

<110> YUN NV

5 <120> DERMATOLOGICAL PREPARATIONS FOR MAINTAINING AND/OR RESTORING HEALTHY SKIN MICROBIOTA

<130> YUN-002

10 <150> BE2016/5454
<151> 2016-06-21

<170> BiSSAP 1.3.6

15 <210> 1
<211> 1406
<212> RNA
<213> Lactobacillus pentosus
<223> 16S rRNA sequence

20 <400> 1

cttaggcggc	tggttcctaa	aaggttaccc	caccgacttt	gggtgttaca	aactctcatg	60
gtgtgacggg	cggtgtgtac	aaggccgggg	aacgtattca	ccgcgcatg	ctgatccgcg	120
attactagcg	attccogactt	catgtaggcg	agttgcagcc	tacaatccga	actgagaatg	180
25 gctttaagag	attagcttac	tctcgcgagt	tcgcaactcg	ttgtaccatc	cattgttagca	240
cgtgtgttagc	ccaggtcata	aggggcatga	tgatttgacg	tcatccccac	cttcctccgg	300
tttgcaccg	gcagtctcac	cagagtgccc	aacttaatgc	tggcaactga	taataagggt	360
tgcgctcggt	gcgggactta	acccaacatc	tcacgacacg	agctgacgac	aaccatgcac	420
cacctgtatc	catgtccccg	aagggaacgt	ctaattcttt	agatttgcatt	agatgtcaa	480
30 gacctggtaa	ggttcttcgc	gtagcttcga	attaaaccac	atgctccacc	gcttgtgcgg	540
gccccccgtca	attcccttga	gtttcagcct	tgcggccgta	ctccccaggc	ggaatgctta	600
atgcgttagc	tgcagcactg	aaggcgaa	accctccaac	acttagcatt	catcgttac	660
gttatggact	accagggtat	ctaattcctgt	ttgctaccca	tacttcgag	cctcagcgtc	720
45 35 agttacagac	cagacagccg	ccttcggccac	tgggttctt	ccatatatct	acgcatttca	780
cgcgtacaca	tggagttcca	ctgtccttctt	ctgcactcaa	gtttcccagt	ttccgatgca	840
cttcttcgggt	tgagccgaag	gtttcacat	cagactaaa	aaaccgcctg	cgctcgcttt	900
acgccccata	aatccggaca	acgcttgcca	cctacgtatt	accgcggctg	ctggcacgtta	960
gttagccgtg	gctttctgggt	taaataccgt	caatacctga	acagttactc	tcagatatgt	1020
40 tcttcctttaa	caacagagtt	ttacgagccg	aaacccttct	tcactcacgc	ggcggtgctc	1080
catcagactt	tgtccatttgc	tggaaagattc	cctactgctg	cctcccgtag	gagtttgggc	1140
cgtgtctcag	tcccaatgtg	gccgattacc	ctctcaggc	ggctacgtat	cattgccatg	1200
gtgagccgtt	accccaccat	ctagctaata	cgccgcggga	ccatccagaa	gtgatagccg	1260
aagccatctt	tcaaactcggt	accatgcgggt	ccaagttgtt	atgcgttatt	agcatctgtt	1320
tccaggtgtt	atcccccgct	tctggcagg	tttcccacgt	gttactcacc	agttcgccac	1380
45 45 tcactcaaataat	gtaaatcatg	atgcaaa				1406

<210> 2
 <211> 20
 <212> DNA
5 <213> Artificial Sequence
 <223> Primer 8F

<400> 2
 agagttttagt cctggctcag 20
10

<210> 3
 <211> 20
 <212> DNA
15 <213> Artificial Sequence
 <223> Primer 1525R

<400> 3
 aaggaggtagt tccagccgca 20
20

<210> 4
 <211> 1425
 <212> RNA
25 <213> *Lactobacillus plantarum*
 <223> 16S rRNA

<400> 4
 ggttcctaaa aggttacccc accgactttg ggtgttacaa actctcatgg tgtgacgggc 60
30 ggtgtgtaca aggcccgaaa acgtattcac cgccggcatgc tgatccgcga ttactagcga 120
 ttccgacttc atgttaggcga gttgcagcct acaatccgaa ctgagaatgg cttaagaga 180
 ttagcttaact ctcgcgagtt cgcaactcgt tgtaccatcc attgttagcac gtgtgttagcc 240
 caggtcataa gggcatgat gatttgacgt catccccacc ttccctccggg ttgtcaccgg 300
 cagtctcacc agagtgcaca acttaatgtt ggcactgtt aataagggtt gcgcgcgttg 360
35 cgggacttaa cccaaacatct cacgacacga gctgacgaca accatgcacc acctgtatcc 420
 atgtccccga agggaaacgta taatcttta gatttgacata gtatgtcaag acctggtaag 480
 gttcttcgog tagcttcgaa ttaaaccaca tgctccaccc cttgtgcggg ccccccgtcaa 540
 ttccctttag gttcagcctt gcggccgtac tccccaggcg gaatgtttaa tgcgttagct 600
 gcagcactga agggcgaaaa ccctccaaca ctttagcatc atcggttacg gtatggacta 660
40 ccagggtatc taatccgtt tgctacccat actttcgagc ctcagcgtca gttacagacc 720
 agacagccgc ctccgcact ggtgttcttc catatatcta cgcatatcc acgttacat 780
 ggagttccac tgtcccttca tgcaactcaag tttcccagggtt tccgatgcac ttcttcgggt 840
 gagccgaagg ctttcacatc agacttaaaa aaccgcctgc gctcgctta cggccaaataa 900
 atccggacaa cgcttgcac ctacgttata cccggcgtgc tggcacgttag ttagccgtgg 960
45 ctttctgggtt aaataccgtc aatacgtt aacttactct cagatatgtt cttctttaac 1020
 aacagagttt tacgagccga aacccttctt cactcacgcg gcgttgctcc atcagacttt 1080

- 17 -

- 18 -

<210> 6
<211> 19
<212> DNA
<213> Artificial Sequence
5 <223> Primer Sequence

<400> 6
tccgttagtg aacctgogg 19

10 <210> 7
<211> 20
<212> DNA
<213> Artificial Sequence
15 <223> Primer Sequence

<400> 7
tcctccgctt attgatatgc 20

20

PCT

Print Out (Original in Electronic Form)

(This sheet is not part of and does not count as a sheet of the international application)

0-1	Form PCT/RO/134 Indications Relating to Deposited Microorganism(s) or Other Biological Material (PCT Rule 13bis)	
0-1-1	Prepared Using	PCT Online Filing Version 3.5.000.251e MT/FOP 20141031/0.20.5.20
0-2	International Application No.	EP2017054513
0-3	Applicant's or agent's file reference	YUN-002
1	The indications made below relate to the deposited microorganism(s) or other biological material referred to in the description on:	
1-1	page	4
1-2	line	15
1-3	Identification of deposit	
1-3-1	Name of depositary institution	BCCM Belgian Coordinated Collections of Microorganisms (BCCM)
1-3-2	Address of depositary institution	BCCM Coordination Cell, Federal Public Planning Service Science Policy, 231, avenue Louise, 1050 Brussels, Belgium
1-3-3	Date of deposit	12 May 2016 (12.05.2016)
1-3-4	Accession Number	BCCM LMG P-29611
1-4	Additional Indications	L. rhamnosus YUN-S1.0
1-5	Designated States for Which Indications are Made	All designations
2	The indications made below relate to the deposited microorganism(s) or other biological material referred to in the description on:	
2-1	page	4
2-2	line	13
2-3	Identification of deposit	
2-3-1	Name of depositary institution	BCCM Belgian Coordinated Collections of Microorganisms (BCCM)
2-3-2	Address of depositary institution	BCCM Coordination Cell, Federal Public Planning Service Science Policy, 231, avenue Louise, 1050 Brussels, Belgium
2-3-3	Date of deposit	09 March 2016 (09.03.2016)
2-3-4	Accession Number	BCCM LMG P-29456
2-4	Additional Indications	L. plantarum YUN-V2.0
2-5	Designated States for Which Indications are Made	All designations

PCT

Print Out (Original in Electronic Form)

(This sheet is not part of and does not count as a sheet of the international application)

3	The indications made below relate to the deposited microorganism(s) or other biological material referred to in the description on:	
3-1	page	4
3-2	line	14
3-3	Identification of deposit	
3-3-1	Name of depositary institution	BCCM Belgian Coordinated Collections of Microorganisms (BCCM)
3-3-2	Address of depositary institution	BCCM Coordination Cell, Federal Public Planning Service Science Policy, 231, avenue Louise, 1050 Brussels, Belgium
3-3-3	Date of deposit	09 March 2016 (09.03.2016)
3-3-4	Accession Number	BCCM LMG P-29455
3-4	Additional Indications	
3-5	Designated States for Which Indications are Made	
FOR RECEIVING OFFICE USE ONLY		

0-4	This form was received with the international application: (yes or no)	YES
0-4-1	Authorized officer	Wilson, Patrick

FOR INTERNATIONAL BUREAU USE ONLY

0-5	This form was received by the international Bureau on:	
0-5-1	Authorized officer	

CLAIMS

1. A topical dermatological composition comprising one or more live *Lactobacillus* species; wherein at least one of said *Lactobacillus* species is a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene.
5
2. A live *Lactobacillus* species for use in restoring and/or maintaining a healthy skin microbiota, by topical route, said *Lactobacillus* species being a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene.
10
3. Use of one or more live *Lactobacillus* species, in the preparation of a topical dermatological composition for restoring and/or maintaining a healthy skin microbiota; wherein at least one of said *Lactobacillus* species is a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene.
15
4. Method for restoring and/or maintaining a healthy skin microbiota; comprising at least one step of administering by topical route, to an individual, an effective amount of one or more live *Lactobacillus* species; wherein at least one of said *Lactobacillus* species is a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene.
20
5. A composition comprising one or more live *Lactobacillus* species for use in restoring and/or maintaining a healthy skin microbiota, by topical route, said *Lactobacillus* species being selected from the list comprising *L. plantarum*, *L. pentosus* and *L. rhamnosus*; wherein said *L. plantarum* is a *L. plantarum* strain having at least 97% sequence similarity with SEQ ID N° 4 in its 16S rRNA gene, said *L. pentosus* is a *L. pentosus* strain having at least 97% sequence similarity with SEQ ID N° 1 in its 16S rRNA gene, and said *L. rhamnosus* is a *L. rhamnosus* strain having at least 97% sequence similarity with SEQ ID N° 5 in its 16S rRNA gene.
25
6. A *Lactobacillus* strain being *L. rhamnosus* YUN-S1.0 deposited under accession number LMG P-29611 (deposited at BCCM on May, 12 2016).
30
7. A composition comprising a *Lactobacillus* strain as defined in claim 6.
35
8. The composition as defined in claim 7; wherein said composition is a topical skin composition.
9. The use according to claim 3, or the composition according to anyone of claims 1, 5, 7 or 8 wherein said composition is a topical skin composition in the form of a gel, cream,
40

ovule, suppository forms, foam, lotion, or ointment.

10. A *Lactobacillus* strain as defined in claim 6 or a composition as defined in any one of
5 claims 7 or 8; for use in restoring and/or maintaining a healthy skin microbiota, by
topical route.
11. Topical use of one or more live *Lactobacillus* species in probiotherapy of the skin;
wherein said *Lactobacillus* species are selected from the list comprising *L. plantarum*,
L. pentosus and *L. rhamnosus*.
- 10 12. Topical use according to claim 11; wherein said probiotherapy consists of restoring
and/or maintaining a healthy skin microbiota in a subject in need thereof.
13. Topical use as defined in anyone of claims 11 or 12, the use as defined in claim 3, the
15 method as defined in claim 4, the live *Lactobacillus* species as defined in claim 2, or
the composition as defined in claim 5; wherein said *Lactobacillus* species is a
Lactobacillus strain selected from the list comprising *L. plantarum* YUN-V2.0
deposited under accession number LMG P-29456; *L. pentosus* YUN-V1.0 deposited
under accession number LMG P-29455; and *L. rhamnosus* YUN-S1.0 deposited under
20 accession number LMG P-29611.

- 1/1 -

Fig. 1

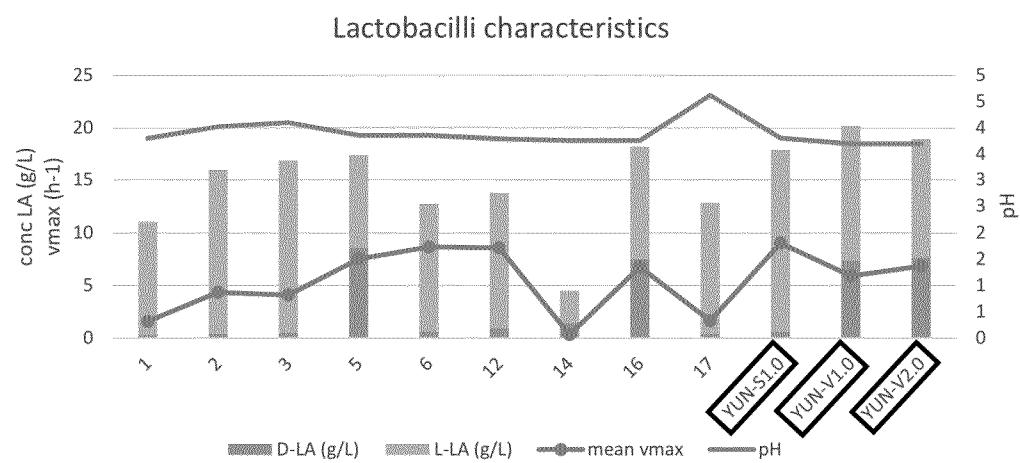
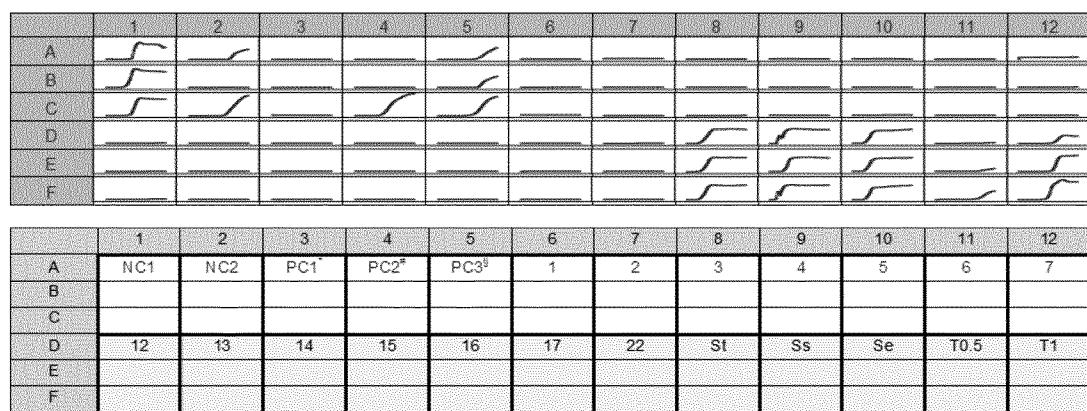


Fig. 2



INTERNATIONAL SEARCH REPORT

International application No
PCT/EP2017/065006

A. CLASSIFICATION OF SUBJECT MATTER	INV.	A61Q19/00	C12R1/25	A61K35/747	C12R1/225	A61P17/00
		A61K8/99				

ADD.

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

A61Q C12R A61K A61P

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

EPO-Internal, WPI Data, BIOSIS, EMBASE

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 01/13956 A2 (GANEDEN BIOTECH INC [US]; FARMER SEAN [US]) 1 March 2001 (2001-03-01) claims 1-3 -----	1-5,9, 11-13
X	EP 2 149 368 A1 (OREAL [FR]; NESTEC SA [CH]) 3 February 2010 (2010-02-03) paragraph [0133] - paragraph [0137] claims 1-4 -----	1-5,9, 11-13
A	EP 1 736 537 A1 (ORGANOBALANCE GMBH [DE]) 27 December 2006 (2006-12-27) paragraph [0021] claims 1-18 ----- -/-	1-5,9, 11-13



Further documents are listed in the continuation of Box C.



See patent family annex.

* Special categories of cited documents :

- "A" document defining the general state of the art which is not considered to be of particular relevance
- "E" earlier application or patent but published on or after the international filing date
- "L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- "O" document referring to an oral disclosure, use, exhibition or other means
- "P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

"&" document member of the same patent family

Date of the actual completion of the international search	Date of mailing of the international search report
7 September 2017	15/11/2017

Name and mailing address of the ISA/
European Patent Office, P.B. 5818 Patentlaan 2
NL - 2280 HV Rijswijk
Tel. (+31-70) 340-2040,
Fax: (+31-70) 340-3016

Authorized officer

Steinheimer, K

INTERNATIONAL SEARCH REPORT

International application No
PCT/EP2017/065006

C(Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 2013/188626 A2 (DOW GLOBAL TECHNOLOGIES LLC [US]) 19 December 2013 (2013-12-19) page 1 - page 5 pages 1-2; examples 1-2 -----	11,12
X	EP 2 332 521 A1 (OREAL [FR]; NESTEC SA [CH]) 15 June 2011 (2011-06-15) claims 1-13 paragraph [0074] - paragraph [0090] -----	11,12
X	KR 2012 0089530 A (KOREA RES INST OF BIOSCIENCE [KR]) 13 August 2012 (2012-08-13) paragraphs [0047], [0021]; claims 1-7; figure 5 & DATABASE Geneseq [Online] 11 April 2013 (2013-04-11), Kim MS et al.: "Lactobacillus plantarum strain MH19 16S ribosomal RNA", Database accession no. BAK32313 the whole document -----	1-5,9, 11-13
X	KR 2015 0075447 A (JUNG LAB CO LTD [KR]; KIM SUNG OK [KR]) 6 July 2015 (2015-07-06) claim 4 -----	1-5,9, 11-13
X	WO 2016/023688 A1 (NESTEC SA [CH]) 18 February 2016 (2016-02-18) claims 1-15 -----	1-5,9, 11-13
X	WO 2011/052996 A2 (CJ CHEILJEDANG CORP [KR]; KIM BONG JOON [KR]; JUNG HEON WOONG [KR]; LE) 5 May 2011 (2011-05-05) claims 6, 10 -----	1-5,9, 11-13

INTERNATIONAL SEARCH REPORT

International application No.
PCT/EP2017/065006

Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

3. Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

see additional sheet

1. As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.

2. As all searchable claims could be searched without effort justifying an additional fees, this Authority did not invite payment of additional fees.

3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:

4. No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

1-4(completely); 5, 9, 11-13(partially)

Remark on Protest

The additional search fees were accompanied by the applicant's protest and, where applicable, the payment of a protest fee.

The additional search fees were accompanied by the applicant's protest but the applicable protest fee was not paid within the time limit specified in the invitation.

No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

1. claims: 1-4(completely); 5, 9, 11-13(partially)

Claims relating to *Lactobacillus plantarum*: Topical use of *Lactobacillus plantarum* in probiotherapy of the skin (claim 11). Topical dermatological composition comprising *lactobacillus plantarum* and corresponding uses.

2. claims: 5, 9, 11-13(all partially)

Claims relating to *Lactobacillus pentosus*: Topical use of *Lactobacillus pentosus* in probiotherapy of the skin (claim 11). Topical dermatological composition comprising *lactobacillus pentosus* and corresponding uses.

3. claims: 6-8, 10(completely); 5, 9, 11-13(partially)

Claims relating to *Lactobacillus rhamnosus*: Topical use of *Lactobacillus rhamnosus* in probiotherapy of the skin (claim 11). Strain YUN-S1.0 (claim 6) and its use in restoring or maintaining healthy skin.

INTERNATIONAL SEARCH REPORT

Information on patent family members

International application No

PCT/EP2017/065006

Patent document cited in search report	Publication date	Patent family member(s)		Publication date
WO 0113956	A2	01-03-2001	AT 270558 T CA 2382670 A1 DE 60012026 D1 DE 60012026 T2 EP 1212093 A2 ES 2221624 T3 JP 2003507437 A US 2001033838 A1 US 2003170334 A1 US 2004208860 A1 US 2006147544 A1 US 2008274153 A1 WO 0113956 A2	15-07-2004 01-03-2001 12-08-2004 30-12-2004 12-06-2002 01-01-2005 25-02-2003 25-10-2001 11-09-2003 21-10-2004 06-07-2006 06-11-2008 01-03-2001
EP 2149368	A1	03-02-2010	CN 102131495 A EP 2149368 A1 US 2011014248 A1 WO 2010013182 A1	20-07-2011 03-02-2010 20-01-2011 04-02-2010
EP 1736537	A1	27-12-2006	AU 2006261100 A1 AU 2012203217 A1 CA 2607911 A1 CN 101203600 A EP 1736537 A1 EP 1893744 A2 EP 1995307 A1 JP 5495559 B2 JP 5826241 B2 JP 2008539747 A JP 2012070745 A JP 2014057600 A KR 20080031201 A US 2009317370 A1 US 2012328586 A1 US 2014072542 A1 US 2017202889 A1 WO 2006136420 A2	28-12-2006 21-06-2012 28-12-2006 18-06-2008 27-12-2006 05-03-2008 26-11-2008 21-05-2014 02-12-2015 20-11-2008 12-04-2012 03-04-2014 08-04-2008 24-12-2009 27-12-2012 13-03-2014 20-07-2017 28-12-2006
WO 2013188626	A2	19-12-2013	NONE	
EP 2332521	A1	15-06-2011	BR 112012013720 A2 CN 102762194 A EP 2332521 A1 EP 2509581 A1 FR 2953407 A1 JP 2013512947 A KR 20120116429 A SG 181510 A1 US 2012294841 A1 WO 2011070509 A1	04-04-2017 31-10-2012 15-06-2011 17-10-2012 10-06-2011 18-04-2013 22-10-2012 30-07-2012 22-11-2012 16-06-2011
KR 20120089530	A	13-08-2012	NONE	
KR 20150075447	A	06-07-2015	NONE	
WO 2016023688	A1	18-02-2016	CN 106573023 A EP 3180012 A1 US 2017224750 A1	19-04-2017 21-06-2017 10-08-2017

INTERNATIONAL SEARCH REPORT

Information on patent family members

International application No

PCT/EP2017/065006

Patent document cited in search report	Publication date	Patent family member(s)		Publication date
		WO	2016023688 A1	18-02-2016
W0 2011052996	A2	05-05-2011	AU 2010314024 A1	10-05-2012
			CA 2778372 A1	05-05-2011
			CN 102597216 A	18-07-2012
			DK 2494031 T3	01-08-2016
			EP 2494031 A2	05-09-2012
			ES 2582828 T3	15-09-2016
			HK 1173468 A1	16-10-2015
			JP 5709883 B2	30-04-2015
			JP 2013509176 A	14-03-2013
			KR 20110046020 A	04-05-2011
			MY 156849 A	15-04-2016
			PL 2494031 T3	31-10-2016
			US 2012208260 A1	16-08-2012
			WO 2011052996 A2	05-05-2011