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# DESCRIPTION

## Technical field of the invention

[0001] The present invention relates to a composition for use in the treatment and/or prevention of a gastrointestinal condition, comprising a combination of *Lactobacillus* strains. The gastrointestinal condition may be a bacterial infection by *Clostridium difficile*, *Salmonella* or *Escherichia coli*.

## Background

[0002] The mammalian intestine is colonized by an estimated 100 trillion bacteria, which have co-evolved with the host in a symbiotic relationship. This collection of microbial populations in the host is referred to as the microbiota. The microbiota can efficiently protect the intestine against colonization by exogenous pathogens and potentially harmful indigenous microorganisms (pathobionts). Mucosal immune responses to normal intestinal bacteria are also important for development and physiology of the host. Breakdown in immunological tolerance to microbiota lead to inappropriate local and systemic immune responses to intestinal bacterial communities that may contribute to multiple disease states such as inflammatory bowel disease (IBD) and irritable bowel syndrome (IBS). The increased prevalence of these chronic conditions has been suggested to be influenced by factors including bacterial infections, antibiotic exposure, as well as dietary factors, stress and degree of hygiene.

[0003] Alteration in the balance of the intestinal microbiota results in disrupted intestinal homeostasis, which increases the risk of pathogen infection and the overgrowth of pathobionts, particularly in immunocompromised hosts. Pathobionts are typically colitogenic in that they can trigger intestinal inflammation. Pseudomembranous colitis which results in severe diarrhea, fever and abdominal pain, is caused by overgrowth of *Clostridium difficile* (CD), a Gram-positive anaerobic bacterium, following long-term treatment with broad-spectrum antibiotics.

[0004] Long-term hospitalization, antibiotic treatment, immune deficiency, cancerous diseases, chemotherapy, and steroid treatment are the main causes of nosocomial diarrhea infections triggered usually by CD but also other enteric bacterial pathogens such as *Salmonella*, *Shigella*, *Camphylobacter*, and *Yersinia*.

[0005] CD has become one of the most serious causes of antibiotic-associated diarrhea. Conventional treatment includes vancomycin or metronidazole for ten days. However, recurrence (occurs in 10 to 25% of cases) is getting increasingly common and represent the greatest challenge associated with CD infections (CDI).

[0006] Increasing rates of CDI have been reported in Canada and the United States, with a

larger proportion of severe and recurrent cases than previously reported. In US, the rates of hospital discharges with CDI listed as any diagnosis increased from 3.82 per 1000 discharges in 2000 to 8.75 per 1000 discharges in 2008; increases were especially prominent among those  $\geq 65$  years of age. Preliminary data indicate that the number of death certificates with enterocolitis due to CDI increased from 793 in 1999 to 7483 in 2008 in US. The rate of pediatric CDI-related hospitalizations also increased between 1997 and 2006, from 0.724 to 1.28 per 1000 hospitalizations. The highest incidence was reported in children 1-4 years of age. The raised incidence and virulence of CDI have coincided with the spread of the hypervirulent CD referred to as NAP1/027, also known as CD BI/NAP1/027 (a restriction endonuclease analysis group BI, pulse-field gel electrophoresis type NAP1, and polymerase chain reaction ribotype 027). NAP 1/027 strain produces a binary toxin and up to 16-fold more toxins than most other hospital outbreak associated strains. Subsequently, epidemics of CDI caused by CD NAP1/027 have been recognized in hospitals in European countries, e.g. the United Kingdom, the Netherlands, Belgium, Austria, and Sweden. The major changes in the epidemiology of CDI during recent years, with increases in incidence and severity of disease have made it a global public health challenge.

**[0007]** The only products used to treat CDI are antibiotics, such as metronidazole, vancomycin, and the recently approved Dificid. Although their high clinical cure rates, 15-30% of patients still experience a recurrence, with each recurrence increasing the risk of further infections. Both the appropriate and the inappropriate use of antibiotics are also associated with the rise in resistance to antibiotics, the emergence of vancomycin resistant new biotypes of *Clostridium difficile*, and the increasing incidence of chronic inflammatory conditions.

**[0008]** The rates of severe cases of CDI have increased during the recent years and the hypervirulent CD NAP 1/027 has been associated with recent outbreaks throughout the world. European Centre for Disease Prevention and Control (ECDC) found (in 2011) that the prevalence of CD NAP1/027 was 5% in the 34 European countries. Large outbreaks of severe, often fatal, colitis have also been reported in North America and Europe. Such an infection may, as a last option when no other treatment has been effective, be treated by the transplantation of feces from a healthy individual. Such transplantation, in addition to the possible psychological discomfort, may also pose the risk of transplanting potentially harmful microorganisms from the donor. Thus, although fecal microbiota transplant has demonstrated some promising results for treating CDI, physicians are concerned about potential infection risks and long-term safety. Recent reports on an obesity onset followed after a fecal transplantation indicate risks and long-term safety issues. In addition, the cumbersome procedures associated with the technology limit its routine use. There are no standard protocols and procedures regarding donor screening/selection, preparation of fecal materials, sanitation issues, recipient preparation, and route of administration.

**[0009]** Thus, there is a need for more effective therapeutic treatment options as well as preventative treatments for CDI.

**[0010]** An altered composition of microbiota interferes with normal intestinal functions at

diverse levels. In addition to triggering the immune system and proinflammatory cytokines, it may also induce release of microbial metabolites, activation of hypothalamic-pituitary-adrenal (HPA) axis with increase of cortisol, leading to alterations of intestinal motility and sensation, disruption of intestinal barrier and impaired production of neurotransmitters with an increased response to stressful events. These complications may cause the IBS which is a chronic functional disorder of the gastrointestinal system and of the most common causes of illness and workplace absenteeism. No cure is available and IBS patients have to rely on treatments to relieve symptoms such as pain, diarrhea and constipation, associated with side effects. The cause of IBS is unknown but there is increasing evidence showing changes in the composition of luminal and mucosal microbiota among IBS patients compared to healthy individuals. Examples of these modifications are a decreased amount of lactobacilli and bifidobacteria along with an increased amount of *Clostridium* in IBS patients. Investigations of gut microbiota of children diagnosed with IBS also indicate a dysbiosis mainly dominated by *Clostridium*.

**[0011]** Therapeutic interventions using lactobacilli are therefore attractive in IBS. A number of studies have concluded that lactobacilli, in general, benefit patients with IBS but it has been difficult to define relative benefits of different bacterial strains. Weak effects and mixed results are primarily due to lack of proper bacterial strain selection and the often poor quality of the studies.

**[0012]** Hell et al. (Beneficial Microbes 4:39-51, 2013) describes the use of probiotics in the treatment of *Clostridium difficile* infections (CDI) and suggests that only a multistrain cocktail resembling a healthy human microbiota may be the way to address CDI. Schoster et al. (Anaerobe, 20:36-41, 2013) describes the inhibition of *Clostridium perfringens* and *C. difficile* by a number of commercial probiotic strains.

### **Summary of the invention**

**[0013]** The above and other objects of the invention are achieved, in full or at least in part, by a composition for use in the treatment and/or prevention of a gastrointestinal composition comprising a combination of *L. salivarius* CW30 (LMG P-28887), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and *L. brevis* SH111 (LMG P-28888).

**[0014]** The present document is also directed to a probiotic composition comprising *L. plantarum* Y1An734 (LMG P-28886), *L. plantarum* SH1313 (LMG P-28884), and *L. plantarum* KS11 (LMG P-28885). The present document is also directed to composition comprising *L. salivarius* CW30 (LMG P-28887), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and *L. brevis* SH111 (LMG P-28888). The bacteria in the composition may consist of *L. salivarius* CW30 (LMG P-28887), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and *L. brevis* SH111 (LMG P-28888) in combination, but the composition may also comprise additional bacterial strains.

**[0015]** The present document is also directed to a probiotic composition comprising a mixture of equal amounts of bacteria of the strains *L. plantarum* Y1An734 (LMG P-28886), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885), *L. salivarius* CW30 (LMG P-28887) and *L. brevis* SH111 (LMG P-28888). It is to be understood that in the context of the present document by the term "equal" and the like in connection with amount of bacteria it is intended that the number of bacterial cells of the different bacterial strains are present in an amount relative to each other so that no bacterial strain is present in an amount more than 5 times of the other, and preferably less than 4 times, 3 times or 2 times of the other strains. Preferably, when the bacteria are present in equal amounts, each bacterial strain is present in an amount so that the ratio to each other bacterial strain is about 1:1.

**[0016]** The present document is also directed to a composition for use in the treatment and/or prevention of a gastrointestinal condition, wherein said composition comprises at least one *Lactobacillus* strain, wherein the *Lactobacillus* strain is chosen from the group consisting of one *L. salivarius* strain, three *L. plantarum* strains and one *L. brevis* strain, wherein the *L. salivarius* strain is *L. salivarius* CW30 (LMG P-28887), wherein the *L. plantarum* strains are *L. plantarum* Y1An734 (LMG P-28886), *L. plantarum* SH1313 (LMG P-28884), and *L. plantarum* KS11 (LMG P-28885), and wherein the *L. brevis* strain is *L. brevis* SH111 (LMG P-28888).

**[0017]** The treatment may be a curative treatment, i.e. a treatment which restores the health of the subject. During a curative treatment, the composition may be administered as an enema. When administered as an enema, the composition may be administered a limited number of times, typically between one and five times, during a short period of time, e.g. during the course of one day to five weeks. When administered orally, the composition may be administered daily during e.g. 1 to 6 weeks.

**[0018]** For certain conditions or in certain patients, the treatment may have a more supportive character, where the composition is administered several times over a longer, sometimes lifelong, periods. The composition may be administered with an interval of one day to two weeks.

**[0019]** The composition may also be used as a prophylactic (preventive) treatment, for example during longer periods of intake of antibiotics in order to support and/or preserve and/or restore a healthy bacterial flora of the gastrointestinal tract.

**[0020]** The composition may comprise at least one *L. salivarius* strain, wherein the *L. salivarius* strain may be *L. salivarius* CW30 (LMG P-28887). An advantage of this is that *L. salivarius* CW30 (LMG P-28887) has a potential anti-microbial effect. The present document is therefore also directed to a composition wherein the bacterial part of the composition comprises, or consists of, *L. salivarius* CW30 (LMG P-28887).

**[0021]** Further, the composition may comprise at least *L. salivarius* CW30 (LMG P-28887) and at least one of the strains *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-

28885), *L. plantarum* Y1An734 (LMG P-28886) and *L. brevis* SH111 (LMG P-28888). The present document is therefore also directed to a composition wherein the bacterial part of the composition comprises or consists of *L. salivarius* CW30 (LMG P-28887) and at least one of the strains *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and *L. brevis* SH111 (LMG P-28888). An advantage of the *L. plantarum* strains *L. plantarum* Y1An734 (LMG P-28886), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885) is that these strains possess anti-inflammatory properties. In addition, these strains may prevent and/or treat intestinal inflammation and improve the gut barrier function. Furthermore, these *L. plantarum* strains may act in synergy with *L. salivarius* CW30 (LMG P-28887) and *L. brevis* SH111 (LMG P-28888) in inhibitory activity against a broader range of pathogenic bacteria associated with gastrointestinal complications. This results in the therapeutic correction of dysbiosis of the gut microbiota promoting to restore the homeostasis of the immune system. An advantage of *L. brevis* SH111 (LMG P-28888) is that it has a synergistic antibacterial potential. This strain has been demonstrated to have a broad antimicrobial activity against different Gram-positive pathogenic bacteria, such as *Staphylococcus aureus* and Gram-negative, such as *Escherichia coli*.

**[0022]** The composition may alternatively comprise at least *L. plantarum* Y1An734 (LMG P-28886), *L. plantarum* SH1313 (LMG P-28884), and *L. plantarum* KS11 (LMG P-28885). The present document is therefore also directed to a composition wherein the bacterial part of the composition comprises or consists of *L. plantarum* Y1An734 (LMG P-28886), *L. plantarum* SH1313 (LMG P-28884), and *L. plantarum* KS11 (LMG P-28885). This composition may further comprise *L. salivarius* CW30 (LMG P-28887) and *L. brevis* SH111 (LMG P-28888).

**[0023]** The gastrointestinal condition may be a diarrhoea, antibiotic-associated diarrhoea (AAD), gastroenteritis, acute gastroenteritis and/or infectious diarrhoea.

**[0024]** Furthermore, the gastrointestinal condition may be a bacterial infection by *Clostridium difficile*, *Salmonella* and/or *Escherichia coli*. The gastrointestinal condition may also be caused by a diarrhoeal disease due to bacteria such as *Campylobacter jejuni*, *Salmonella typhimurium*, *Yersinia enterocolitica*, enteropathogenic *E. coli* (EPEC), enterohemorrhagic *E. coli* (EHEC), *Shigella* spp., *Staphylococcus aureus*, *Listeria monocytogenes*, *Vibrio cholera*, *Vibrio parahaemolyticus* and *Bacillus cereus*. The gastrointestinal condition may also be due to a viral infection, such as Rotavirus associated diarrhoea. The gastrointestinal condition may also be post infective diarrhoea.

**[0025]** Especially, the gastrointestinal condition may be an infection by *Clostridium difficile* NAP 1/027. One advantage of the present invention is that it quickly resolves the disease by targeting the pathogenic bacteria and to reestablishing immunological tolerance and intestinal homeostasis. A second advantage is that the use of purified *Lactobacillus* strains with well defined beneficial health effects avoids the use of feces or other compositions that involve critical safety and health issues.

**[0026]** The gastrointestinal condition may be irritable bowel syndrome (IBS) or inflammatory

bowel disease (IBD). The term inflammatory bowel disease (IBD) includes, among other conditions, Crohn's disease and ulcerative colitis.

**[0027]** The gastrointestinal condition may also be pouchitis or post infection colitis.

**[0028]** The composition may be administered orally, such as in a beverage or food product. The composition may be administered as a beverage, in which case the *Lactobacillus* strain(s) are suspended in a liquid. The liquid may e.g. be water, milk, a salt-sugar solution or a gruel. Alternatively, the composition may be administered as a porridge or pudding. The composition may be included in a pill, which can be swallowed by the patient. An advantage of these routes of administration is that they do not require any advanced medical equipment. Another advantage of administering the composition as a beverage, porridge, pudding or pill is that the patient can control when the beverage, porridge, pudding or pill is swallowed.

**[0029]** The composition may be administered as an enema. An advantage of this route of administration is that it can be performed using equipment which is commonly used in hospitals. An advantage of this route of administration is that the composition may be administered at the location in the gastrointestinal tract where it is most effective. Another advantage is that it is performed by healthcare professionals; the patients can content with a few treatments and will be assessed regularly by care providers.

**[0030]** The composition may be administered as an infusion into the upper gastrointestinal tract. The composition may be administered through a medical device, such as a nasoduodenal tube. An advantage of this route of administration is that it can be performed using equipment which is commonly used in hospitals. Another advantage of this route of administration is that the composition may be administered at the location in the gastrointestinal tract where it is most effective. Another advantage is that the bacterial composition bypasses the stomach and its acidic environment and are directly infused to the intestine.

**[0031]** The composition may be administered through a medical device, such as a nasogastric tube. The composition may be administered via colonoscopy.

**[0032]** The total amount of *Lactobacillus* may be  $10^9$  to  $10^{13}$  CFU per treatment, e.g.  $10^9$ ,  $10^{10}$ ,  $10^{11}$ ,  $10^{12}$  or  $10^{13}$  CFU per treatment. In the case of nasoduodenal infusion the treatment may be repeated after one day, a few days or a week for at least two times.

**[0033]** The composition may be freeze-dried. An advantage of this is that the bacteria may be preserved for long-term storage and no advanced storage facilities are needed to handle the bacteria.

**[0034]** The freeze-dried composition may be contained in a bag, a jar, a capsule or any other kind of container such as a pill. The freeze-dried composition may be one of several components in a mixture. Further, the freeze-dried composition may also be one of several

components of a pill. Other conventional additives may naturally be added to the freeze-dried compositions or to any of the other mentioned compositions as disclosed herein in order to provide stable compositions with suitable shelf life. A skilled person would know such conventional additives and the amount to be used in the compositions.

**[0035]** The composition may further comprise a fiber component.

**[0036]** The fiber component may be chosen from the group consisting of oat fiber, wheat fiber, rye fiber, chia fiber, corn fiber, barley fiber, potato fiber, fruit fiber, vegetable fiber, cereal fiber and fiber from algae.

**[0037]** Other examples of suitable fibers include different kinds of soluble and insoluble fibers. For example, fibers originating from seeds (e.g. linseeds and psyllium seeds) or from nuts (such as walnuts, coconuts, almonds) may be used. Dietary fibers such as inulin may be used. Other conventional additives may be added to the fiber containing compositions.

**[0038]** The bacteria may be suspended in a gruel, a pap, a porridge or a pudding.

**[0039]** The gruel may be oat gruel. Other gruels which may be used include wheat gruel, barley gruel, rye gruel or corn or maize gruel.

**[0040]** Before administration of the composition according to the present invention, the fresh or freeze-dried bacterial composition may be suspended in a suitable liquid. The liquid may be a pharmaceutically acceptable liquid component or any type of suitable medium, e.g. one of the media described above. Preferably, distilled water or buffered aqueous media are used, which contain pharmaceutically acceptable salts and buffers. Suitable salt solutions are PBS (*Phosphate-buffered saline*), GBSS (*Gey's Balanced Salt Solution*), EBSS (*Earle's balanced salt solution*), HBSS (*Hank's Balanced Salt Solution*), and SBF (*Synthetic/Simulated Body Fluid*). The liquid component can also be of a more hydrophobic nature depending of the application.

**[0041]** In the case of oral administration, different flavourings may be added in order to make the mixture comprising the composition more pleasant to taste.

**[0042]** The composition may be ready to use after reactivation in the suitable medium for 24 h at room temperature or up to one week at 4 °C. However, it should be noted that the time period and temperature may vary depending on i.e. the medium used.

**[0043]** The composition may further comprise one or more therapeutic agents, such as an agent against the bacterial induced infection and/or the inflammatory condition.

**[0044]** The composition may comprise the above mentioned five strains or even more, as long as at least one of the above mentioned bacterial strains are present.

**[0045]** A composition as described above may be comprised in an enema. An advantage of such an enema is that the route of administration can be performed using equipment which is commonly used in hospitals. Another advantage of this route of administration is that the composition may be administered at the location in the gastrointestinal tract where it is most effective, i.e. in the colon.

**[0046]** The volume of composition administered will vary depending on the age and size of the person receiving the enema. However general guidelines would be: Infant: 250 ml or less, School-aged child: 200-500 ml, Adult: 200-1,000 ml.

**[0047]** The composition may be a pharmaceutical formulation.

**[0048]** The present document also provides an isolated strain chosen from the group consisting of *L. salivarius* CW30 (LMG P-28887), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and *L. brevis* (LMG P-28888). The bacterial strains are biologically pure.

**[0049]** The present document is also provides an isolated strain of *L. salivarius* for use in the treatment and/or prevention of an infection by *Clostridium difficile* NAP 1/027, wherein the strain is *L. salivarius* CW30 (LMG P-28887).

**[0050]** The present document is also directed to a method for the treatment and/or prevention of a gastrointestinal condition said method comprising the step of administering a pharmaceutically effective amount of a composition comprising one or more of a bacterial strain selected from the group consisting of *L. salivarius* CW30 (LMG P-28887), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and *L. brevis* (LMG P-28888), an enema comprising one or more of these bacteria, or one or more of an isolated bacterial strain selected from the group consisting of *L. salivarius* CW30 (LMG P-28887), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and *L. brevis* (LMG P-28888), to a subject in need thereof. Said gastrointestinal is a gastrointestinal condition as disclosed elsewhere herein.

**[0051]** Use of a composition comprising one or more of a bacterial strain selected from the group consisting of *L. salivarius* CW30 (LMG P-28887), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and *L. brevis* (LMG P-28888), an enema comprising one or more of these bacteria, or one or more of an isolated bacterial strain selected from the group consisting of *L. salivarius* CW30 (LMG P-28887), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and *L. brevis* (LMG P-28888) for the preparation of a medicament for the treatment and/or prevention of a gastrointestinal condition. Said gastrointestinal is a gastrointestinal condition as disclosed elsewhere herein.

**[0052]** Other objectives, features and advantages of the present invention will appear from the

following detailed disclosure, from the attached claims, as well as from the figures.

**[0053]** Generally, all terms used in the claims are to be interpreted according to their ordinary meaning in the technical field, unless explicitly defined otherwise herein. All references to "a/an/the [element, device, component, means, step, etc.]" are to be interpreted openly as referring to at least one instance of said element, device, component, means, step, etc., unless explicitly stated otherwise. The steps of any method disclosed herein do not have to be performed in the exact order disclosed, unless explicitly stated.

**[0054]** As used herein, the term "comprising" and variations of that term are not intended to exclude other additives, components, integers or steps.

#### **Definition of strains**

**[0055]** All strains were deposited at BCCM/LMG (Belgian Coordinated Collections of Micro-organisms/Laboratorium voor Microbiologie, Universiteit Gent (UGent)), Gent, Belgium on May 19<sup>th</sup> 2015. The depositor is Shahram Aghaibeik-Lavasani, Department of Biology, Sölvegatan 35, Building C, 223 62 Lund, Sweden.

**[0056]** The deposition numbers are as follows.

<i>L. plantarum</i> SH1313	LMG P-28884
<i>L. plantarum</i> KS11	LMG P-28885
<i>L. plantarum</i> Y1An734	LMG P-28886
<i>L. salivarius</i> CW30	LMG P-28887
<i>L. brevis</i> SH111	LMG P-28888

The strains are biologically pure.

#### **Definition of strains and terms**

CFU = colony forming units

#### **Description of the tables and figure**

**[0057]**

Figure 1

shows the percentage of CD4<sup>+</sup>Foxp3<sup>+</sup> T cells after co-incubation of lactobacilli with

freshly isolated immune cells from murine intestinal lymphoid tissues.  $1 \times 10^6$  immune cells were incubated with either single *Lactobacillus* sp. or in combination with two or several. \* represents a  $p\text{-value} \leq 0.05$ , \*\* a  $p\text{-value} \leq 0.01$  and \*\*\* a  $p\text{-value} \leq 0.001$  in comparison with the control.

Figure 2

shows the percentage of  $CD4^+Foxp3^+$  T cells after co-incubation of lactobacilli with immune cells from murine intestinal lymphoid tissues as described above.  $1 \times 10^6$  immune cells were incubated with single *Lactobacillus* species including five lactobacilli strains LMG P-28884 - P28888 and four lactobacilli strains isolated from different commercially available food products (*L. paracasei* CNCM I-1518, *L. plantarum* DSM 9843, *L. acidophilus* DSM 13241 and *L. rhamnosus* ATCC 53103). \* represents a  $p\text{-value} \leq 0.05$ , \*\* a  $p\text{-value} \leq 0.01$  and \*\*\* a  $p\text{-value} \leq 0.001$  in comparison with the *L. plantarum* SH1313 (LMG P-28884).

Figure 3

Table 1 shows the antagonistic activity of different species of lactobacilli against *Clostridium difficile* including hyper-virulent NAP1/027 strains; - indicating no inhibition, + indicating inhibition, or no growth around the lactobacilli streaks.

Figure 4

Table 2 shows a cross-streak method to investigate the inter-species inhibition of five different *Lactobacillus* strains. For each group, the strains at the top of the table were the (first-streaked) tester strain; - indicating no inhibition, + indicating inhibition, or no growth in the zone where the *Lactobacillus* streaks converged.

Figure 5

Table 3 shows the antibiotic susceptibility test of *Lactobacillus* strains.

Figure 6

shows the ability of lactobacilli strains to adhere to the human epithelial colorectal cells. *Lactobacillus* species including the five lactobacilli strains LMG P-28884 - P28888 and four lactobacilli strains isolated from different commercially available food products (*L. paracasei* CNCM I-1518, *L. plantarum* DSM 9843, *L. acidophilus* DSM 13241 and *L. rhamnosus* ATCC 53103) were used for comparison. An equal number of cells for each strain ( $5 \times 10^8$ ), was independently incubated (1 h at  $37^\circ\text{C}$ ) with a monolayer of fully differentiated Caco-2 cells. The cells were washed and stained with Giemsa stain solution. The adherent bacteria in randomly selected microscopic fields were counted and averaged. \* represents a  $p\text{-value} \leq 0.05$ , \*\* a  $p\text{-value} \leq 0.01$  and \*\*\* a  $p\text{-value} \leq 0.001$  in comparison with the *L. plantarum* SH1313 (LMG P-28884).

Figure 7

Table 4 shows the antagonistic activity of the combination of five different *Lactobacillus* strains against strains of *Salmonella* and *Escherichia coli*; +++ indicates good inhibition (zone of inhibition  $>5\text{-}10$  mm). ++++ indicates strong inhibition (zone of inhibition  $>10$  mm).

Figure 8

Carbohydrate profiling of *Lactobacillus salivarius* CW30 (LMG P-28887), *Lactobacillus brevis* SH111 (LMG P-28888), *L. plantarum* SH1313 (LMG P-28884), *L. plantarum* KS11

(LMG P-28885), and *L. plantarum* Y1An734 (LMG P-28886), respectively. Scores are related to the color intensity. A positive test is indicated by a color change from purple to yellow (1-3) due to an anaerobic production of acid (-= no change).

### **Detailed description of the invention**

**[0058]** In the present context and invention the following definition apply: The term "intestinal infection and inflammation" is intended to mean gastroenteritis, an increased inflammation of the gastrointestinal tract caused by viral, bacterial or parasitic infections. It may also mean other gastrointestinal condition e.g. irritable bowel syndrome (IBS), inflammatory bowel diseases (IBD), pouchitis or post infection colitis. The term antibiotic-associated diarrhoea (AAD) is intended to mean diarrhea that develops in a person who is taking or recently took antibiotics. One of the most serious causes of AAD is infection with a bacterium, *Clostridium difficile* (CD) a Gram-positive anaerobic bacterium. Furthermore, nosocomial diarrhea infections are mainly triggered by CD but also by other enteric bacterial pathogens such as *Salmonella*, *Shigella*, *Camphylobacter*, and *Yersinia*.

**[0059]** It is also disclosed herein a symbiotic composition, comprising an effective amount *Lactobacillus* bacteria chosen from the group comprising or consisting of one or more of *Lactobacillus salivarius* CW30 (LMG P-28887), *Lactobacillus brevis* SH111 (LMG P-28888), *Lactobacillus plantarum* SH1313 (LMG P-28884), *Lactobacillus plantarum* KS11 (LMG P-28885) and *Lactobacillus plantarum* Y1An734 (LMG P-28886). The bacterial strains are selected from novel lactic acid bacterial strains isolated from different plant sources. Samples from different vegetable plants, raw and after fermentation, were plated on Ragasa agar and incubated for 24-72h at 35-42°C in anaerobic conditions. Single clones were randomly chosen. All clones with gram positive rod shape bacteria were checked for probiotic properties such as resistance to acidic pH (pH 3) and bile salt. The isolated bacteria were identified through carbohydrate fermentation profiling (see Table 5 in Figure 8) and 16S rRNA sequencing. The lactic acid bacterial strains bacterial strains were further selected for ability to adhere to the intestine and inhibit the growth of gastrointestinal pathogens, but not the growth of each other. Typically, when a composition comprises more than one of these bacterial strains, the number of bacterial cells of the different bacterial strains are about equal (i.e. about the same number of bacterial cells, e.g. as determined by measuring the number of CFU or by microscopic calculation of the number of bacteria). A composition comprising one or more of these bacterial strains may be denoted a probiotic composition.

**[0060]** It is also disclosed herein pharmaceutical compositions comprising at least one strain as described herein, i.e. at least one strain selected from the group consisting of *Lactobacillus salivarius* CW30 (LMG P-28887), *Lactobacillus brevis* SH111 (LMG P-28888), *Lactobacillus plantarum* SH1313 (LMG P-28884), *Lactobacillus plantarum* KS11 (LMG P-28885) and *Lactobacillus plantarum* Y1An734 (LMG P-28886). It is also disclosed herein a culture, a

composition or a product comprising one or more of said bacterial strains for use in the prevention and/or treatment of gastrointestinal condition, such as a condition caused by CD.

**[0061]** In the present study more than 18 different and novel lactobacilli strains have been screened and characterized regarding antimicrobial activities against 21 pathogenic CD strains. Interestingly, it turned out that these properties are strain specific and not general for all strains of a genus. A therapeutic composition of the invention comprising bacteria from all of the *Lactobacillus salivarius* CW30 (LMG P-28887), *Lactobacillus brevis* SH111 (LMG P-28888), *Lactobacillus plantarum* SH1313 (LMG P-28884), *Lactobacillus plantarum* KS11 (LMG P-28885) and *Lactobacillus plantarum* Y1An734 (LMG P-28886) strains has shown the ability to inhibit growth of all CD strains but not the growth of each other. It demonstrates the synergy exhibited between the strains of the composition of the present invention.

**[0062]** As is demonstrated in the experimental data provided in this document *L. plantarum* SH1313 (LMG P-28884), *L. brevis* SH111 (LMG P-28888), and *L. salivarius* CW30 (LMG P-28887) have a higher colonization ability (as demonstrated by binding to Caco-2 cells) than *L. plantarum* KS11 (LMG P-28885) and *L. plantarum* Y1An734 (LMG P-28886), although the latter still show good colonization ability. *L. plantarum* SH1313, KS11, and Y1An734 on the other hand were demonstrated to have the highest anti-inflammatory effect (as measured by potential to induce Tregs). A composition comprising *L. plantarum* SH1313, KS11, and Y1An734 may thus be particularly useful in the treatment and/or prevention of a chronic inflammatory disorder such as, but not restricted to, IBD, which includes, among other conditions, Crohn's disease and Ulcerative colitis, pouchitis or post infection colitis. *L. brevis* SH111 (LMG P-28888), and *L. salivarius* CW30 (LMG P-28887) show the strongest antimicrobial effect of the five lactobacilli strains isolated. A composition comprising *L. brevis* SH111 (LMG P-28888), and *L. salivarius* CW30 (LMG P-28887) may thus be particularly useful for use in the treatment and/or prevention of a microbial gastrointestinal infection, such as an infection caused by *Clostridium difficile*, *Salmonella*, *Escherichia coli*, *Campylobacter*, *Shigella*, *Yersinia*, *Staphylococcus*, *Listeria*, *Vibrio*, and/or *Bacillus cereus*. The gastrointestinal infection may be due to a viral infection, such as Rotavirus.

**[0063]** Due to inhibitory properties of the chosen lactobacilli strains on pathogenic strains of CD and/or other gut pathogens and therapeutic effect on patients with antibiotic-associated diarrhea (AAD), the present invention relates to the medical use of the symbiotic *Lactobacillus* combinations for the preparation of pharmaceutical compositions for preventing, treating, or ameliorating symptoms of a gastrointestinal condition selected from, but not limited to, AAD, gastroenteritis, acute gastroenteritis, or infectious diarrhoea. The gastrointestinal condition may be a bacterial infection caused by, but not limited to, *Clostridium difficile*, *Salmonella*, *Escherichia coli*, *Campylobacter*, *Shigella*, *Yersinia*, *Staphylococcus*, *Listeria*, *Vibrio*, or *Bacillus cereus*. The gastrointestinal condition may be due to a viral infection, such as Rotavirus associated diarrhoea, or a post infective diarrhoea.

**[0064]** The compositions of the present document may also be used effectively in the treatment of gastroenteritis caused by *Salmonella* or *Escherichia coli*. *Salmonella* spp. cause

one of the most common forms of food poisoning worldwide. The experiments clearly demonstrated that the symbiotic composition of this invention strongly inhibited the growth of pathogenic strains of *Salmonella* (*S. enteritidis* and *S. typhimurium*) and *Escherichia coli*.

**[0065]** The present invention also relates prophylactic (preventive) use against CD infections. Patients with CD infection are usually treated with antibiotics, such as metronidazole and vancomycin. The composition of this invention has further been tested experimentally and revealed no sensitivity against metronidazole and vancomycin showing its potential to be used as a prophylactic (preventive) treatment, for example during longer periods of intake of antibiotics in order to support and/or preserve and/or restore a healthy bacterial flora of the gastrointestinal tract. It can also be used to inhibit the recurrence which is common and represent the greatest challenge associated with CD infections.

**[0066]** The present invention also relates treatment of infections caused by the hypervirulent CD referred to as NAP1/027. The raised incidence and virulence of CD infections have coincided with the spread of CD NAP1/027. Screening of the novel lactobacilli revealed that only few *Lactobacillus* strains had the potential to inhibit the growth of the 2 different hypervirulent CD NAP1/027 strains. These therapeutic strains are included in the symbiotic composition presented by this invention.

**[0067]** The present document is also directed to the use of the symbiotic composition sated in this invention for the prophylactic or therapeutic treatment of chronic inflammatory disorders such as, but not restricted to, IBD which includes, among other conditions, Crohn's disease and Ulcerative colitis, pouchitis or post infection colitis. Interaction of bacterial pathogens with the intestinal epithelial cells initiates a cascade of inflammatory processes that contribute to a gastrointestinal condition. An altered intestinal microbiota and inflammation disrupts the gut barrier function, allowing pathogens to further multiply and colonize the gut. Foxp3<sup>+</sup> Regulatory T cells (Tregs) are essential in the maintenance of immune tolerance in the gut suppressing harmful inflammatory responses. Based on our experiences, the combinations of lactobacilli have further been screened for the potential to trigger Tregs. The therapeutic composition of the invention has shown the greatest potential to trigger these anti-inflammatory cells compared to other tested combinations. It once again indicated the synergy displayed between the strains of the composition of the present invention.

**[0068]** The present document is also directed to the use of symbiotic composition sated in this invention for the prophylactic or therapeutic treatment of intestinal conditions associated with abdominal discomfort or pain such as, but not restricted to, irritable bowel syndrome (IBS). A role for gut microbiota and influence of stress has been suggested to be crucial for development IBS. The therapeutic composition presented in this invention has shown major success for treatment of IBS.

**[0069]** The composition may be administered orally. The strains of the composition of the present invention are all selected due to their unique and superior abilities to survive in the low pH of the stomach and in the high bile acid content of the small intestine, and ability to attach

to colonic mucosa and to temporary colonise the large intestine. The composition may be administered as a beverage, in which case the bacterial strain(s) are suspended in a liquid. The liquid may be water, milk, a salt-sugar solution or a gruel. Alternatively, the composition may be administered as a porridge or pudding. The composition may be included in a pill, which can be swallowed by the patient.

**[0070]** The composition may be administered as an enema. An advantage of this route of administration is that the composition may be administered at the location in the gastrointestinal tract where it is most effective. Fecal microbiota transplant administered by enema has been used for treatment of patients. Although using enema is considered to be safe and the technique demonstrated some promising results for treating CD infections, but physicians are concerned about potential infection risks and long-term safety due to the content of the therapy which is fecal bacteria from voluntary donors. However, the therapeutic composition presented in this invention consist of *Lactobacillus* strains chosen from extensive studies which have been tested on patients with great success. A further advantage of the method is that it is performed by healthcare professionals; the patients can content with a few treatments and will be assessed regularly by care providers.

**[0071]** The composition may be administered as an infusion into the upper gastrointestinal tract. The composition may be administered through a medical device, such as a nasoduodenal tube. An advantage of this route of administration is that the composition may be administered at the location in the gastrointestinal tract in gastrointestinal conditions where it is most effective.

**[0072]** The bacterial combination may be administrated together with a pharmaceutically acceptable liquid component, but also as a powder. Preferably, distilled water or buffered aqueous media are used, which contain pharmaceutically acceptable salts and buffers. The bacterial strains may be present in a so that a total concentration of bacteria of at least  $10^9$  to  $10^{13}$  CFU is administered per treatment. If more than one of the bacterial strains is to be administered, typically the same amount of all different strains is administered.

**[0073]** The composition may further comprise a fiber component. The health benefits of dietary fiber have long been appreciated. Some of them are also known to selectively stimulate the growth and/or activity of one or a limited number of commensal bacteria in the colon, thus improving host health.

**[0074]** One or more pharmaceutically acceptable liquid components may be needed. Such components are well known to those skilled in the art.

**[0075]** The composition may further comprise a fiber component. The fiber component may be chosen from the group consisting of, but not limited to, oat fiber, wheat fiber, rye fiber, chia fiber, corn fiber, barley fiber, potato fiber, fruit fiber, vegetable fiber, cereal fiber and fiber from algae. The bacterial combination may also be suspended in a gruel, a pap, a porridge or a pudding.

[0076] In order to prepare a composition according to the present document, the bacterial strains to be included in the composition may be grown separately or in combination of one or more strains. If grown separately and more than one bacterial strain is intended to be administered, typically the different bacterial strains are mixed so that the resulting composition comprises the same amount of bacteria (CFU). The lactobacilli strains disclosed herein may be grown in any media commonly used for growing lactobacilli, such as Man-Rogosa-Sharpe (MRS) broth/agar and Rogosa agar. The bacteria are typically grown at a temperature of about 25 to 42°C, such as 30 to 37°C, in particular about 37°C. The bacteria may be grown in liquid culture with or without shaking or on solid media. Typically, the bacteria are grown for a time period of 24-72 hours. The time of growth will as is known to the person skilled in the art depend upon different factors, such as the temperature, the amount of bacteria inoculated, the nutrient content of the medium etc. Typically, the bacteria are grown to the stationary phase. Before the bacteria are administered to a subject in need thereof, the bacteria are typically isolated from their used growth medium by e.g. centrifuging or filtering the bacterial culture. The bacteria may be washed one or more times in e.g. a buffer or salt solution, such as in PBS (phosphate buffered saline) buffer or 0.9% NaCl. Before use, the bacteria are then resuspended in the medium to be used for administration. The bacteria may also be freeze-dried in order to prolong their storage ability. The freeze-dried bacteria may be administered in their dry state or they may be dissolved in a liquid or liquid-containing medium before administration. It is also possible to reactivate the freeze-dried bacteria in a suitable medium for 24 h at room temperature or up to one week at 4 °C. However, it should be noted that the time period and temperature for such a reactivation may vary depending on i.e. the medium used. It is also possible to use the bacterial cultures directly without separating them from their used growth medium. It is also possible to separate the bacteria from their used growth medium and administer the used growth medium instead of the bacteria themselves.

[0077] The invention will now be described in more detail in the following sections. The examples are however only illustrative and not intended to limit the scope of the present invention.

## **Experiments**

[0078] **Assessment of interaction between lactobacilli and *Clostridium difficile*:** To assess *in vitro* interaction between lactobacilli and *Clostridium difficile*, antagonistic activity of 18 *Lactobacillus* strains against 21 pathogenic *C. difficile* including two hyper-virulent CD NAP1/027 (also known as CD BI/NAP1/027 or CD NAP1/027) strains was determined. The CD strains belonged to different genotypes and were previously isolated from feces of hospitalized patients in Sweden, mainly at Skåne University Hospital in Malmö, with random cases of antibiotic-associated diarrhea were included. Antagonistic activity of lactobacilli was detected on agar plates as inhibition of *C. difficile* growth. The results are summarized in Table 1. *Lactobacillus* strains were cultured 48 h in Man-Rogosa-Sharpe (MRS) broth at 37°C. *C. difficile* strains were cultured 24 h in brain heart infusion (BHI) broth at 37°C in an anaerobic

environment. All bacterial cultures were centrifuged at 4°C and pellet were suspended in 0.9% NaCl solution to prepare the concentration equal to 0.5 on the McFarland turbidity scale. 100 µl of *C. difficile* suspension was inoculated and spread onto Wilkins-Chalgren blood agar plates. 10 µl of each *Lactobacillus* suspension were streaked in different lines (about 2 cm). Plates were incubated in an anaerobic environment at 37°C for 48 h, and inhibition zones of *Lactobacillus* growth on *C. difficile* streak were investigated.

**[0079]** The results clearly demonstrate that *L. plantarum* strains SH1313 (LMG P-28884), KS11 (LMG P-28885) and Y1An734 (LMG P-28886), as well as *S. salivarius* CW30 (LMG P-28887) and *L. brevis* SH111 (LMG P-28888) inhibit the growth of several of the tested *C. difficile* strains. Particularly, a mixture of these five strains inhibited the growth of all tested *C. difficile* strains. In addition, *S. salivarius* CW30 (LMG P-28887) and *L. brevis* SH111 (LMG P-28888) and also *L. plantarum* KS11 (LMG P-28885) inhibited the growth of CD NAP1/027 strains. Particularly, a mixture of these five strains inhibited the growth of both of the tested CD NAP1/027 strains. Notably, other tested strains of Lactobacilli did not inhibit the growth of *C. difficile* to the same extent.

**[0080] Interactions of lactobacilli and immune cells:** It has been shown that direct interaction of *C. difficile* with the intestinal epithelial cells initiates a cascade of inflammatory processes that contribute to intestinal diseases such as diarrhea and pseudomembranous colitis. An altered intestinal microbiota composition and inflammation disrupts the gut barrier function, allowing *C. difficile* to further multiply and colonize the gut. Foxp3 expressing CD4<sup>+</sup> T cells (Regulatory T cells; Tregs) are essential in the maintenance of immune tolerance in the gut suppressing harmful inflammatory responses. Based on previous experiences, we have isolated intestinal lymphoid tissues (Peyer's patches and mesenteric lymph nodes) from mice. They were pooled and single cells suspensions of immune cells were provided *in vitro*. The cells were then co-incubated with five different lactobacilli; *L. plantarum* SH1313 (LMG P-28884), *L. salivarius* CW30 (LMG P-28887), *L. brevis* SH111 (LMG P-28888), *L. plantarum* KS11 (LMG P-28885), and *L. plantarum* Y1An734 (LMG P-28886) in single or combinations together. A total of  $1 \times 10^6$  of immune cells was seeded/well in a 96-well plate (in the RPMI 1640 cell culture medium).  $1 \times 10^7$  CFU/ml (100µl) of bacteria (suspended in RPMI 1640) were added to the cells and the co-culture was incubated for 20 h (37°C, humidified atmosphere with 5% CO<sub>2</sub>). At the next day, the cells were washed and analysed for prevalence of Tregs using flow cytometry. Tregs were detected expressing the receptor CD4, transcription factor (forkhead box P3; Foxp3), and intracellular cytokine IL-10. The bacteria were used either in single form or in combinations of two, three, four of five strains. The combinations contained equal amounts of each strains (added at the same day of experiment), all to a final concentration of  $1 \times 10^7$  CFU/ml. Control samples contained cell culture medium only. The results are shown in Figure 1. The data represent the frequency of CD4<sup>+</sup>Foxp3<sup>+</sup> T cells in percent.

**[0081]** It was clearly evident from the results that only selected single strains of *L. plantarum* (SH1313, KS11, or Y1An734), have the potential to induce an anti inflammatory profile and

activate Tregs among the cells from freshly isolated intestinal lymphoid tissues. *L. salivarius* CW30, *L. brevis* SH111 can trigger Tregs only in combination with carefully selected strains of *L. plantarum*. The results also demonstrated that the *L. plantarum* SH1313, *L. plantarum* KS11, *L. plantarum* Y1 An734, *L. salivarius* CW30, and *L. brevis* SH111 work synergistically to activate and increase Tregs. The most significant and profound anti inflammatory effect achieves by using a combination of all five strains.

**[0082]** Using the same assay, we have further compared the potential of different lactobacilli to induce Tregs. We have used *L. plantarum* SH1313 (LMG P-28884), *L. salivarius* CW30 (LMG P-28887), *L. brevis* SH111 (LMG P-28888), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and four lactobacilli strains isolated from different commercially available food products; *L. paracasei* CNCM I-1518 (Actimel), *L. plantarum* DSM 9843 (ProViva), *L. acidophilus* DSM 13241 (Aria A-fil) and *L. rhamnosus* ATCC 53103 (Valio yoghurt).  $1 \times 10^7$  CFU/ml (100 $\mu$ l) of each strain were co-cultured with immune cells for 20 h at 37°C. The results are shown in Figure 2. The data represent the frequency of CD4<sup>+</sup>Foxp3<sup>+</sup> T cells in percent.

**[0083]** The results revealed that *L. plantarum* (SH1313, KS11, and Y1An734) have significantly stronger potential to induce Tregs than other strains including the *L. paracasei* CNCM I-1518, *L. plantarum* DSM 9843, *L. acidophilus* DSM 13241 and *L. rhamnosus* ATCC 53103, i.e. that these *L. plantarum* strains (*L. plantarum* (SH1313, KS11, and Y1An734)) have a pronounced anti-inflammatory effect.

**[0084]** Statistical evaluations were all performed using StatView and the data were evaluated using nonparametric Mann-Whitney test.

**[0085] *Inter-species inhibition of five different Lactobacillus strains:*** There are increasing evidence showing that certain mixtures of *Lactobacillus* species have more beneficial effects than their component strains. The various synergistic or additive effects *in vivo* has been suggested to be influenced by possible mutual inhibition by the component strains. We investigated whether the five different lactobacilli; *L. plantarum* SH1313 (LMG P-28884), *L. salivarius* CW30 (LMG P-28887), *L. brevis* SH111 (LMG P-28888), *L. plantarum* KS11 (LMG P-28885), and *L. plantarum* Y1An734 (LMG P-28886) may inhibit the growth of the others *in vitro*. A cross-streak assay was used for this purpose. Each strain was streaked onto MRS agar using a 1 ml loop (3 parallel lines). After the lines had dried, other strains were streaked perpendicular to these strains creating 3 potential zones of inhibition for each combination of strains. If the tester strain hindered the growth of the second-streaked strain, it implicated an inhibition. The results are shown in Table 2.

**[0086]** The results clearly revealed that *L. salivarius* CW30, and *L. brevis* SH111, have very little, and the *L. plantarum* SH1313, *L. plantarum* KS11, *L. plantarum* Y1An734, have no ability to inhibit the growth of the other selected bacterial strains. Despite the fact that the proposed multi-strain lactobacilli preparation (consisting of *L. salivarius* CW30, and *L. brevis* SH111, *L. plantarum* SH1313, *L. plantarum* KS11, and *L. plantarum* Y1An734) can maintain inhibition of a

range of pathogenic *C. difficile*, there is no inhibition between the component strains influencing the therapeutic efficacy of the bacterial mixture.

**[0087] Antibiotic susceptibility of *Lactobacillus* strains:** The antibiotic susceptibilities of lactobacilli have shown to be species- and strain-dependent. Therefore, the antibiotic susceptibility of the five different lactobacilli; *L. plantarum* SH1313 (LMG P-28884), *L. salivarius* CW30 (LMG P-28887), *L. brevis* SH111 (LMG P-28888), *L. plantarum* KS11 (LMG P-28885), and *L. plantarum* Y1An734 (LMG P-28886) were examined using disk diffusion test. Rogosa Agar plates were inoculated with bacteria from freshly prepared cultures of *Lactobacillus* strains. Disk diffusion tests were carried out for erythromycin (15 µg disk), vancomycin (30 µg disk), and metronidazole (5 µg disk). The results are shown in Table 3. "S", stands for sensitive and "R", stands for resistant.

**[0088]** The antibiotic profiling data reveals that *L. salivarius* CW30, and *L. brevis* SH111, *L. plantarum* SH1313, *L. plantarum* KS11, and *L. plantarum* Y1An734 are all sensitive to erythromycin, which is a type of medicine known as a macrolide antibiotic, and none of the strains is sensitive to vancomycin or metronidazole.

**[0089] Adhesion of lactobacilli strains to human enterocyte-like Caco-2 cells:** The adhesion of lactobacilli strains to the intestinal mucosa is one of the appropriate properties for their colonization in the intestinal tract, where they compete with other bacteria. The human intestinal cell line Caco-2 (from colonic adenocarcinoma) is a well characterized cellular model which develops characteristics of mature enterocytes with functional brush-border microvilli when fully differentiated. The Caco-2 cell line has been extensively used to study bacterial adhesion and invasion.

**[0090]** In the present study, the adhesion of the various *Lactobacillus* strains to Caco-2 cells was compared. Caco-2 cells were grown in 100-mm plastic Petri dishes. Cells were seeded at approximately  $5 \times 10^5$  in a 35-mm dish and used two weeks after confluence (fully differentiated cells). Bacterial cells ( $5 \times 10^8$ ) were added to washed cell monolayers and incubated for 1 h at 37°C. We have used *L. plantarum* SH1313 (LMG P-28884), *L. salivarius* CW30 (LMG P-28887), *L. brevis* SH111 (LMG P-28888), *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886) and four lactobacilli strains isolated from different commercially available food products; *L. paracasei* CNCM I-1518, *L. plantarum* DSM 9843, *L. acidophilus* DSM 13241, and *L. rhamnosus* ATCC 53103.

**[0091]** All monolayers were then washed 3 times with PBS to release unbound bacteria and stained with Giemsa stain solution. Dishes were then washed, dried for 1 h and examined microscopically (magnification  $\times 100$ ) under oil immersion. The *Lactobacillus* strains in 20 randomly selected microscopic fields were counted and averaged.

**[0092]** As indicated in Figure 6, there were significant variations in adherence of different lactobacilli strains to the Caco-2 cells, indicating that adhesive properties are not a universal

feature of lactobacilli. *L. plantarum* SH1313 (LMG P-28884), *L. brevis* SH111 (LMG P-28888), and *L. salivarius* CW30 (LMG P-28887) showed strongest efficiency of adhesion to Caco-2 cells. Although the adhesion properties of *L. plantarum* KS11 (LMG P-28885), *L. plantarum* Y1An734 (LMG P-28886), and *L. paracasei* CNCM 1-1518 are a bit weaker their adherence to Caco-2 cells still seems more prominent than the adhesion properties of *L. plantarum* DSM 9843, *L. acidophilus* DSM 13241, and *L. rhamnosus* ATCC 53103.

**[0093] Antagonistic activity of a combination of *Lactobacillus* strains against strains of**

***Salmonella* and *Escherichia coli*:** We have further evaluated the ability of the combination of five lactobacilli; *L. plantarum* SH1313 (LMG P-28884), *L. salivarius* CW30 (LMG P-28887), *L. brevis* SH111 (LMG P-28888), *L. plantarum* KS11 (LMG P-28885), and *L. plantarum* Y1An734 (LMG P-28886) to inhibit the *in vitro* growth of *Salmonella enteritidis*, *Salmonella typhimurium*, and *Escherichia coli*. *Lactobacillus* strains were cultured 48 h in Man-Rogosa-Sharpe (MRS) broth at 37°C. *Salmonella* and *E. coli* strains were cultured 24 h in brain heart infusion (BHI) broth at 37°C. All bacterial cultures were centrifuged at 4°C and the pellets were suspended in 0.9% NaCl solution to prepare the concentration equal to 0.5 on the McFarland turbidity scale. 100 µl cell suspensions of *Salmonella* or *E. coli* strains were inoculated and spread onto Wilkins-Chalgren blood agar plates. A mixture containing equal amounts of each five *Lactobacillus* strains (added at the same day of experiment), were prepared to a final concentration of  $1 \times 10^7$  CFU/ml. 10 µl of *Lactobacillus* suspension were streaked in different lines (about 2 cm). Plates were incubated for 48 h at 37°C (with 5% CO<sub>2</sub>). The growth inhibition zones were then measured.

**[0094]** As indicated in the Table 4, the results clearly demonstrate that a combination of *L. salivarius* CW30, and *L. brevis* SH111, *L. plantarum* SH1313, *L. plantarum* KS11, and *L. plantarum* Y1An734 is able to strongly inhibit growth of the *Salmonella enteritidis*, *Salmonella typhimurium*, and *Escherichia coli*. The data further demonstrate the antimicrobial potential of the lactobacilli mixture against gastrointestinal pathogenic bacteria.

### **Clinical cases**

**[0095]** The present invention is demonstrated by the following examples of clinical trials which were performed at the Department of Infectious Diseases, Skåne University Hospital in Malmö, Sweden, (during 2013-2014). The implicated examples, materials and procedures are to be understood broadly with the scope and spirit of the invention as set forth herein.

**[0096] Case 1:** *Clostridium difficile* infection is a complication commonly associated with antibiotic therapy in hospitalized patients. A 70 years old female patient with ongoing anal cancer and stoma surgery receives several courses of antibiotics due to wound problems. She showed recurrent *C. difficile* colitis during one year and were treated several times with metronidazole and then vancomycin with a de-escalation strategy under six weeks. Once more, she developed recurrent *C. difficile* infection (CDI). Method: A combination of five

lactobacilli; *L. plantarum* SH1313 (LMG P-28884), *L. salivarius* CW30 (LMG P-28887), *L. brevis* SH111 (LMG P-28888), *L. plantarum* KS11 (LMG P-28885), and *L. plantarum* Y1An734 (LMG P-28886) were provided (equal amounts of each strain).  $1 \times 10^{12}$  CFU were combined with a sterile fiber-rich gruel. The composition was then mixed to an enema solution and a colonic infusion was performed by a rectal tube. The treatment was repeated five times with an interval of 3-5 days.

**[0097] Result:** The patient was clinically and bacteriological cured after 3 months and showed no sign of recurrent infection up to one year after the last treatment.

**[0098] Case 2:** A 75 years old male patient with multiple diseases. He showed recurrent *C. difficile* colitis after long-term treatment with Dalacin, containing the antibiotic called clindamycin hydrochloride, the year before. He received two courses of metronidazole, followed by 10 days of vancomycin treatment. After six weeks under de-escalation strategy, he showed recurrent CDI. He received a fecal microbiota transplantation containing feces from healthy donors prepared by the clinical investigators at the hospital. After new course of the antibiotic treatment he developed recurrent CDI once more.

**[0099] Method:** A composition of five lactobacilli was provided as described above and the enema solution was infused. The treatment was repeated three times with an interval of 7 days.

**[0100] Result:** The patient was clinically cured after a few days and showed no sign of recurrent infection up to one year after the last treatment.

**[0101] Case 3:** A 55 years old female patient showed recurrent *C. difficile* colitis after long-term antibiotic treatments. She received two courses of metronidazole, followed by 10 days of vancomycin treatment. After six weeks under de-escalation strategy, she showed recurrent CDI.

**[0102] Method:** A composition of five lactobacilli was provided as described above and the enema solution was infused. The treatment was repeated three times with an interval of 7 days.

**[0103] Result:** The patient was clinical cured after a few days and showed no sign of recurrent infection up to one year after the last treatment.

**[0104] Case 4:** A young man, 20 years old, developed irritable bowel syndrome (IBS) after earlier gastroenteritis. He received fecal microbiota transplantation containing feces from healthy donors (prepared by the clinical investigators at the hospital) at two occasions with no sign of improvement of his condition.

**[0105] Method:** A composition of five lactobacilli was provided as described above and the enema solution was infused. The treatment was repeated three times with an interval of 7

days.

**[0106]** *Result:* The patient showed significant improvement in symptomatology of IBS. He has not shown any signs of clinical deterioration up to one year after the last treatment.

**[0107]** The results from the clinical investigations demonstrate that a bacterial composition consisting of *L. plantarum* SH1313 (LMG P-28884), *L. salivarius* CW30 (LMG P-28887), *L. brevis* SH111 (LMG P-28888), *L. plantarum* KS11 (LMG P-28885), and *L. plantarum* Y1An734 (LMG P-28886), is effective and relatively safe for treatment of patients suffering from severe gastrointestinal conditions

## REFERENCES CITED IN THE DESCRIPTION

### Cited references

This list of references cited by the applicant is for the reader's convenience only. It does not form part of the European patent document. Even though great care has been taken in compiling the references, errors or omissions cannot be excluded and the EPO disclaims all liability in this regard.

### Non-patent literature cited in the description

- HELL et al. Beneficial Microbes, 2013, vol. 4, 39-51 [\[0012\]](#)
- SCHOSTER et al. Anaerobe, 2013, vol. 20, 36-41 [\[0012\]](#)

## P A T E N T K R A V

1. Sammensætning til anvendelse i behandlingen og/eller forebyggelsen af en mave-tarm-tilstand, omfattende en kombination af *L. salivarius* CW30, deponeret som LMG P-28887, *L. plantarum* Y1An734, deponeret som LMG P-28886, *L. plantarum* SH1313, deponeret som LMG P-28884, *L. plantarum* KS11, deponeret som LMG P-28885, og *L. brevis* SH111, deponeret som LMG P-28888.
2. Sammensætning til anvendelse ifølge krav 1, hvor nævnte mave-tarm-tilstand indbefatter inflammation.
3. Sammensætning til anvendelse ifølge et hvilket som helst af de foregående krav, hvor mave-tarm-tilstanden er en diarré, antibiotika-associeret diarré (AAD), gastroenteritis, akut gastroenteritis og infektiøs diarre.
4. Sammensætning til anvendelse ifølge et hvilket som helst af de foregående krav, hvor mave-tarm-tilstanden er en bakterieinfektion med *Clostridium difficile*, *Salmonella* eller *Escherichia coli*.
5. Sammensætning til anvendelse ifølge et hvilket som helst af de foregående krav, hvor mave-tarm-tilstanden er en infektion med *Clostridium difficile* NAP1/027.
6. Sammensætning til anvendelse ifølge et hvilket som helst af krav 1 til 4, hvor mave-tarm-tilstanden er irritabel tarmsyndrom (IBS) eller inflammatorisk tarmsygdom (IBD).
7. Sammensætning til anvendelse ifølge et hvilket som helst af de foregående krav, hvor sammensætningen gives oralt eller som et klyasma eller som en infusion til den øvre mave-tarm-kanal.
8. Sammensætning til anvendelse ifølge et hvilket som helst af de foregående krav, hvor den samlede mængde af *Lactobacillus* er  $10^9$  til  $10^{13}$  CFU per behandling.
9. Sammensætning til anvendelse ifølge et hvilket som helst af de foregående krav, hvor sammensætningen er frysetørret.
10. Sammensætning til anvendelse ifølge krav 9, hvor sammensætningen yderligere omfatter en fiberkomponent.
11. Klyasma omfattende sammensætningen ifølge krav 1.

# DRAWINGS

Figure 1:

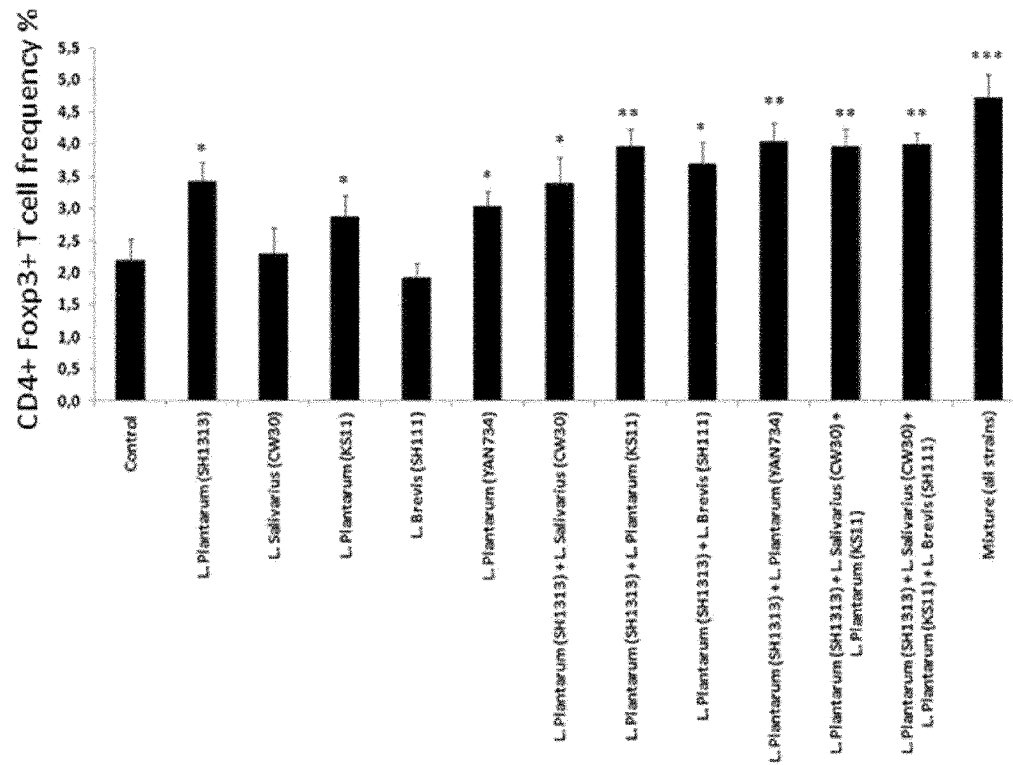


Figure 2:

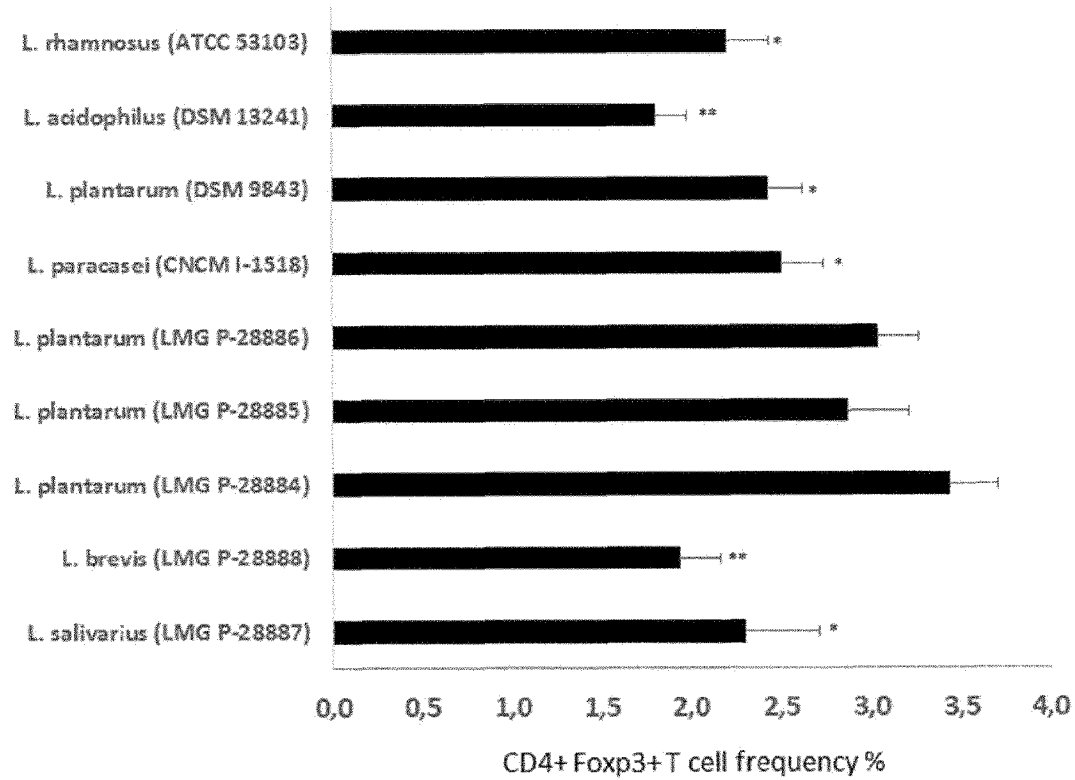


Figure 3:

Table 1

[illegible]

Figure 4:

Table 2

	L. Plantarum (SH1313)	L. Salivarius (CW30)	L. Brevis (SH111)	L. Plantarum (KS11)	L. Plantarum (VAN734)
L. Plantarum (SH1313)	-	-	+	-	-
L. Salivarius (CW30)	-	-	+	+	-
L. Brevis (SH111)	-	+	-	-	+
L. Plantarum (KS11)	-	-	-	-	-
L. Plantarum (VAN734)	-	+	-	-	-

Figure 5:

Table 3

	L. Plantarum (SH1313)	L. Salivarius (CW30)	L. Brevis (SH111)	L. Plantarum (KS11)	L. Plantarum (VAN734)
Erythromycin	S	S	S	S	S
Vancomycin	R	R	R	R	R
Metronidazole	R	R	R	R	R

Figure 6:

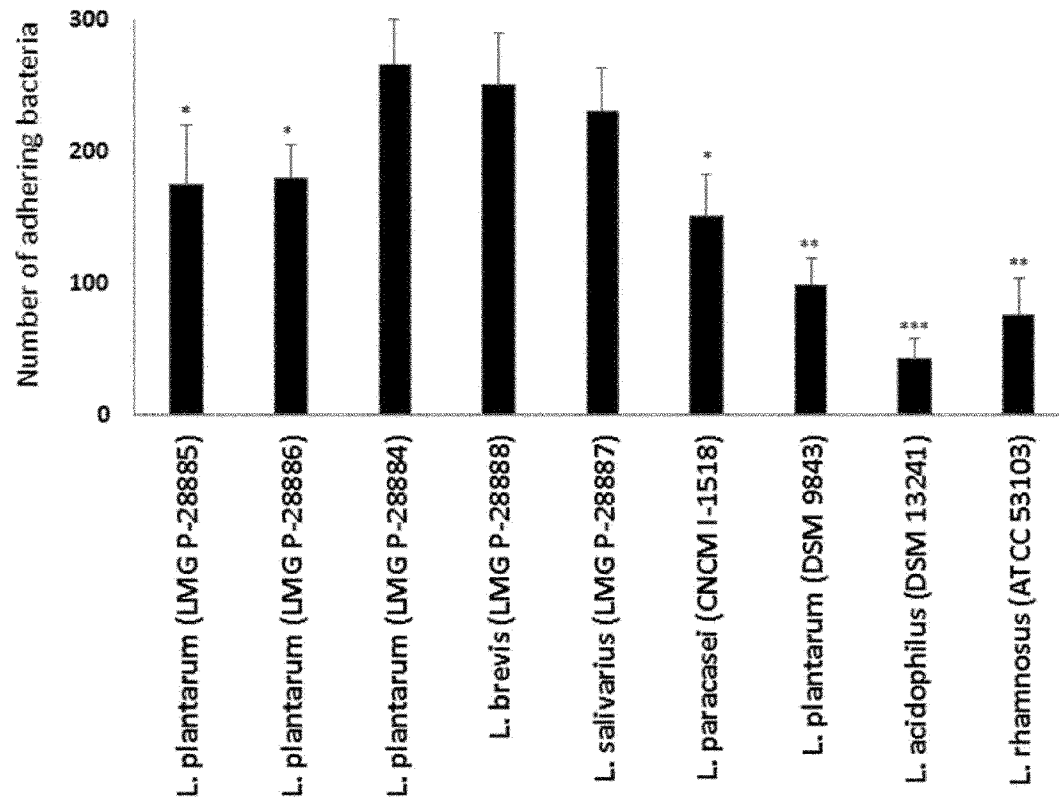


Figure 7:

Table 4

	<i>Salmonella enteritidis</i>	<i>Salmonella typhimurium</i>	<i>Escherichia coli</i>
Mixture of <i>L. Plantarum</i> (SH1313)			
+ <i>L. Salivarius</i> (CW30) + <i>L. Brevis</i> (SH111) + <i>L. Plantarum</i> (KS11)	+++	++	+++
+ <i>L. Plantarum</i> (YAN734)			

Figure 8:

	Glucose				Maltotriose				Maltose				Maltotetraose				Maltopentaose				Maltohexaose				Maltoseptosa				Maltooctaose				Maltononaose				Maltodecaose				Maltodecaose				Maltotridecaose				Maltotetradecaose				Maltopentadecaose				Maltohexadecaose				Maltoseptadecaose				Maltooctadecaose				Maltononaidecaose				Maltodecaidecaose				Maltotridecaidecaose				Maltotetradecaidecaose				Maltopentadecaidecaose				Maltohexadecaidecaose				Maltoseptadecaidecaose				Maltooctadecaidecaose				Maltononaidecaidecaose				Maltodecaidecaidecaose				Maltotridecaidecaidecaose				Maltotetradecaidecaidecaose				Maltopentadecaidecaidecaose				Maltohexadecaidecaidecaose				Maltoseptadecaidecaidecaose				Maltooctadecaidecaidecaose				Maltononaidecaidecaidecaose				Maltodecaidecaidecaidecaose				Maltotridecaidecaidecaidecaose				Maltotetradecaidecaidecaidecaose				Maltopentadecaidecaidecaidecaose				Maltohexadecaidecaidecaidecaose				Maltoseptadecaidecaidecaidecaose				Maltooctadecaidecaidecaidecaose				Maltononaidecaidecaidecaidecaose				Maltodecaidecaidecaidecaidecaose				Maltotridecaidecaidecaidecaidecaose				Maltotetradecaidecaidecaidecaidecaose				Maltopentadecaidecaidecaidecaidecaose				Maltohexadecaidecaidecaidecaidecaose				Maltoseptadecaidecaidecaidecaidecaose				Maltooctadecaidecaidecaidecaidecaose				Maltononaidecaidecaidecaidecaidecaose				Maltodecaidecaidecaidecaidecaidecaose				Maltotridecaidecaidecaidecaidecaidecaose				Maltotetradecaidecaidecaidecaidecaidecaose				Maltopentadecaidecaidecaidecaidecaidecaose				Maltohexadecaidecaidecaidecaidecaidecaose				Maltoseptadecaidecaidecaidecaidecaidecaose				Maltooctadecaidecaidecaidecaidecaidecaose				Maltononaidecaidecaidecaidecaidecaidecaose				Maltodecaidecaidecaidecaidecaidecaidecaose				Maltotridecaidecaidecaidecaidecaidecaidecaose				Maltotetradecaidecaidecaidecaidecaidecaidecaose				Maltopentadecaidecaidecaidecaidecaidecaidecaose				Maltohexadecaidecaidecaidecaidecaidecaidecaose				Maltoseptadecaidecaidecaidecaidecaidecaidecaose				Maltooctadecaidecaidecaidecaidecaidecaidecaose				Maltononaidecaidecaidecaidecaidecaidecaidecaose				Maltodecaidecaidecaidecaidecaidecaidecaidecaose				Maltotridecaidecaidecaidecaidecaidecaidecaidecaose				Maltotetradecaidecaidecaidecaidecaidecaidecaidecaose				Maltopentadecaidecaidecaidecaidecaidecaidecaidecaose				Maltohexadecaidecaidecaidecaidecaidecaidecaidecaose				Maltoseptadecaidecaidecaidecaidecaidecaidecaidecaose				Maltooctadecaidecaidecaidecaidecaidecaidecaidecaose				Maltononaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltodecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltotridecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltotetradecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltopentadecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltohexadecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltoseptadecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltooctadecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltononaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltodecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltotridecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltotetradecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltopentadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltohexadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltoseptadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltooctadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltononaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltodecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltotridecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltotetradecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltopentadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltohexadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltoseptadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltooctadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltononaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltodecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltotridecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltotetradecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltopentadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltohexadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltoseptadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltooctadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltononaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltodecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltotridecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltotetradecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaidecaose				Maltopentadecaidecaidecaidecaidecaidecaidecaidecaidecaidecaideca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