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Lactobacillus rhamnosus strains of oral and vaginal origin show strong antifungal activity in vitro

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**ABSTRACT**

**Background:** Intake of probiotic bacteria may prevent oral Candida infection.

**Objective:** To screen the antifungal activity of 14 Lactobacillus candidate strains of human origin, against six opportunistic C. albicans and non-albicans species. A second aim was to study the acid production of the four strains showing the strongest antifungal activity.

**Methods:** We used an agar overlay growth inhibition assay to assess the antifungal activity of the lactobacilli. The acid-producing capacity was measured with pH micro-sensors.

**Results:** All 14 Lactobacillus candidates inhibited the growth of the Candida spp. The four best-performing strains were L. rhamnosus DSM 32992 (oral origin), L. rhamnosus DSM 32991 (oral), L. jensenii 22842 (vaginal), and L. rhamnosus P801 (vaginal). The difference between L. rhamnosus DSM 32992 and the other three strains was statistically significant (p < 0.001). The Candida spp. differed in susceptibility: C. parapsilosis was highly inhibited, while C. krusei was not or slightly inhibited. The oral L. rhamnosus DSM 32992 and DSM 32991 strains showed the lowest pH-values.

**Conclusion:** Screening of probiotic lactobacilli showed significant strain-dependent variations in their antifungal capacity in a pH-dependent mode. Two strains of oral origin were most effective. A further characterization seems justified to elaborate on their probiotic properties.

**Introduction**

Candida species (spp.) are ubiquitous commensals taking part of a healthy oral microbiota. C. albicans is the predominant species and is believed to be present in its natural blastospore form in around 50% of the healthy population [1,2]. In healthy individuals, the oral microbiota appears to prevent Candida spp. to shift from harmless commensals to disease-causing pathogens. However, certain conditions may disrupt the homeostatic bacteriome–mycobiome relationship and predispose the Candida spp. to become pathogenic and cause disease in the oral mucosa, termed oral candidosis [3]. While C. albicans is the main contributor to infection, the incidence of oral infection due to non-albicans spp. have increased over the last decades, and the most frequently isolated strains associated with disease are C. glabrata, C. krusei, C. tropicalis, C. dubliniensis, and C. parapsilosis [4,5].

Currently, treatment options of oral candidosis include topical and systemic antifungtics. However, the American Centers for Disease Control and Prevention has reported that about 7% of all Candida bloodstream isolates are resistant to fluconazole [6]. Although C. albicans is the most common cause of severe Candida infections, resistance is most common in other species, particularly C. glabrata and C. parapsilosis [6]. According to the World Health Organization, Candida infection is a significant concern for human health in vulnerable populations due to the emergence of new resistance mechanisms in the microorganisms. In most countries where data are available, drug resistance appears to be higher among non-albicans species than among C. albicans [7]. Hence, the search of alternative antifungtic treatments to prevent and combat oral mucosal infections of both C. albicans and non-albicans spp. is highly relevant.

Probiotic bacteria have been proposed as a supplement to conventional therapies to prevent and combat oral Candida-infection [8–10]. Probiotic bacteria are defined as ‘live microorganism that, when administered in adequate amounts, confer a health benefit on the host’ [11]. The most investigated probiotics for anti-Candida activity belong to the genera Lactobacillus and Bifidobacterium, but also Streptococcus has been proposed [12]. Laboratory studies have shown that probiotic lactobacilli can inhibit C. albicans growth [13,14], interfere with the C. albicans biofilm formation on dentures [15], and to interfere in hyphae formation
In addition, Tan and coworkers [17] showed that sterile spent media of L. gasseri BF and L. rhamnosus BF could disrupt biofilm formation of C. tropicalis BF, C. krusei BF, and C. parapsilosis BF and reduce their metabolic activity. It has also been demonstrated that cell-free supernatants of the probiotic L. pentosus strain LAP1 could inhibit the growth of C. albicans, C. tropicalis, and C. krusei in vitro [18].

Several clinical trials have confirmed that regular probiotic administration can decrease the Candida cell counts in samples from the oral cavity, in particular among elderly individuals prone to infections [8,19–22] and be a possible supplement to conventional antymycotic treatment [23,24]. However, the genera of probiotics used in the trials are wide-spread, and the choice of an appropriate type of probiotic for this specific target is challenging. Recently, we showed significant differences in the antifungal properties of two L. reuteri strains (ATCC PTA 5289 and DSM 17938) [9]. This finding emphasizes that the probiotic strain-specificity cannot be disregarded, and that the search for appropriate probiotic strains with strong anti-Candida properties for oral application is warranted. The primary aim of this study was, therefore, to screen the antifungal activity of 14 selected probiotic Lactobacillus candidate strains, isolated from the human oral cavity or vagina, against six opportunistic Candida spp. in vitro. A secondary aim was to study the acid production abilities of the four best-performing lactobacilli in the growth inhibition assay with aid of pH microsensor measurements through Candida and Lactobacillus co-cultures.

Materials and methods

Strains and culture conditions

Fourteen selected Lactobacillus spp. (Deerland Probiotics & Enzymes A/S, Hundested, Denmark) were used in this study together with six clinical Candida spp. from the Department of Clinical Microbiology, Copenhagen University Hospital, Copenhagen, Denmark, and six control spp. from the Culture Collection, University of Gothenburg, Sweden (Table 1). The lactobacilli were initially cultured on de Man Rogosa Sharpe (MRS) agar (Oxoid Ltd., Basingstoke, Hampshire, UK) for 24 hours (h) in an anaerobic chamber at 37°C (10% H₂, 5% CO₂ and 85% N₂). The Candida strains were cultured on BD Difco™ Sabouraud Dextrose (SD) agar (Becton, Dickinson and Company, Sparks, MD, USA) for 24 h in ambient air at 37°C.

Growth inhibition assay

The growth inhibition assay was performed as described earlier [9]. In brief, one distinct colony of overnight cultured lactobacilli was transferred to 5 mL MRS broth (Oxoid Ltd., Basingstoke, Hampshire, UK) and incubated at 37°C for 24 h under anaerobic conditions. The following day the lactobacilli were harvested by centrifugation at 2,000 rpm for 10 min at room temperature. The supernatants of the Lactobacillus strains were obtained after centrifugation and destroyed. The pellets were washed three times in phosphate-buffered saline (PBS), and the OD was adjusted to 1.8 at 630 nm (Genesys™ 10S UV-Vis Spectrophotometer, Thermo Fisher Scientific, Waltham, MA, USA), corresponding to approximately 10⁸ cfu/mL. The cultures were then serially diluted in MRS broth in sixfold steps. One mL of the supernatants, undiluted suspensions and cell suspensions corresponding to approximately 10⁷ and 10⁶ CFU/mL were added to 24 mL sterilized molten MRS agar (−45°C) in Petri dishes and the agar could solidify. The plates were incubated overnight at 37°C under anaerobic conditions. One single colony of each of the overnight cultured Candida strains was added to 5 mL broth and aerobically incubated at 37°C for 24 h. The following day, one additional layer of 25 mL of molten sterile SD agar was poured onto top of the MRS agar with grown lactobacilli and could solidify and air-dry for 3 h in room temperature. The overnight cultured Candida strains were diluted in SD broth to a final OD of 0.2 at 500 nm. The Candida suspensions were stamped on the plates with a Steers steel pin replicator (CMI-Promex ICN, Pedricktown, NJ, USA) and left to dry for 2 h at room temperature.

Table 1. List of Lactobacillus spp. and Candida spp. used in the study.

<table>
<thead>
<tr>
<th>Genus</th>
<th>Origin</th>
<th>Species</th>
</tr>
</thead>
<tbody>
<tr>
<td>Lactobacillus</td>
<td>Oral cavity</td>
<td>L. rhamnosus NEU427</td>
</tr>
<tr>
<td></td>
<td></td>
<td>L. rhamnosus ERB18, DSM</td>
</tr>
<tr>
<td></td>
<td></td>
<td>L. rhamnosus ERB 36, DSM</td>
</tr>
<tr>
<td></td>
<td></td>
<td>L. reuteri 5289</td>
</tr>
<tr>
<td></td>
<td></td>
<td>L. reuteri 17938</td>
</tr>
<tr>
<td>Vaginal</td>
<td></td>
<td>L. acidophilus EB03</td>
</tr>
<tr>
<td></td>
<td></td>
<td>L. crispatus 23B33</td>
</tr>
<tr>
<td></td>
<td></td>
<td>L. crispatus NEU458 DSM15224</td>
</tr>
<tr>
<td></td>
<td></td>
<td>L. gasseri EB01</td>
</tr>
<tr>
<td></td>
<td></td>
<td>L. jensenii 12B1</td>
</tr>
<tr>
<td></td>
<td></td>
<td>L. jensenii 22B42</td>
</tr>
<tr>
<td></td>
<td></td>
<td>L. rhamnosus PB01</td>
</tr>
<tr>
<td>Candida</td>
<td>Control CCUG</td>
<td>C. albicans CCUG 46,390</td>
</tr>
<tr>
<td></td>
<td>strains</td>
<td>C. dublinsiis CCUG 48,722</td>
</tr>
<tr>
<td></td>
<td></td>
<td>C. glabrata CCUG 63,819</td>
</tr>
<tr>
<td></td>
<td></td>
<td>C. krusei CCUG 36,126</td>
</tr>
<tr>
<td></td>
<td></td>
<td>C. parapsilosis CCUG 56,136</td>
</tr>
<tr>
<td></td>
<td></td>
<td>C. tropicalis CCUG 47,037</td>
</tr>
<tr>
<td></td>
<td>Clinical isolates</td>
<td>C. albicans CBS 562 NT</td>
</tr>
<tr>
<td></td>
<td></td>
<td>C. dublinsiis 41_3 ZZMK</td>
</tr>
<tr>
<td></td>
<td></td>
<td>C. glabrata CBS 863</td>
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<tr>
<td></td>
<td></td>
<td>C. krusei RV 491</td>
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<tr>
<td></td>
<td></td>
<td>C. parapsilosis 36 PBS</td>
</tr>
<tr>
<td></td>
<td></td>
<td>C. tropicalis DSM 7524</td>
</tr>
</tbody>
</table>
The plates were subsequently aerobically incubated overnight at 37°C. As controls, the *Candida* strains were stamped on the top of plates with no lactobacilli within the bottom MRS agar layer.

The assays were carried out in duplicates and repeated three times on different occasions. The results were evaluated with a 4-step score, modified from Simark-Mattson et al. [25]: Score 0 = complete growth inhibition (no visible colonies), Score 1 = almost total growth inhibition (colonies slightly visible), Score 2 = slight growth inhibition (colonies are clearly visible but smaller than at the control plate), and Score 3 = no growth inhibition (colonies equal to those at the control plate). Two observers (MRJ and PTR) scored the plates independently, and in case of disagreement, consensus was reached through discussion.

**Microsensor measurement of pH**

Acid production (measured by pH) of the four best-performing lactobacilli in the growth inhibition assay and the effect of pH on *Candida* growth was measured with pH microelectrodes (pH-100 and ref-100; Unisense A/S, Århus, Denmark) with a linear response between pH 4–9 and a 90% response time of <10 s and a detection limit of 0.01 pH units. Measurements were performed using a modification of a previously described procedure [9] in selected plates from the growth inhibition test. In brief, the electrodes were mounted in a motorized PC-controlled profiling setup (MM33 and MC-232, Unisense A/S), and positioning and data acquisition were controlled by dedicated software (SensorTrace Profiling, Unisense A/S). The pH microsensor was calibrated against buffers of pH 4 and 7 at room temperature with a linear slope of −56.1 mV/pH unit. The pH was measured on the final day of the growth inhibition test. Before measuring, the pH and reference electrodes were placed approximately 2 mm above the surface of the *Candida* colonies. The pH was measured from the surface of the colonies in 100 μm increments until the depth of maximum 4.5 mm into the agar to make sure the sensors had reached the bottom agar layer containing the lactobacilli. Selection of the plates was based on the results from the interference test; the plates with the highest inhibition scores for each *Lactobacillus* spp. were selected regardless of concentration (CK and POJ). The pH was measured through dense, slightly inhibited colonies of *Candida* (*C. krusei* and *C. tropicalis*) and through vague, almost completely inhibited colonies (*C. albicans*, *C. glabrata*, *C. dubliniensis*, and *C. parapsilosis*) incubated on plates with *L. rhamnosus* DSM 32992, *L. rhamnosus* DSM 32991, *L. jensenii* 22B42, and *L. rhamnosus* PB01, respectively. In addition, pH was measured in control plates containing only the lactobacilli in the bottom agar layer, in a control plate with only *Candida* strains, and lastly in a control agar plate without lactobacilli or *Candida* strains.

**Statistical analyses**

All data were processed with SAS Enterprise guide software (version 7.1, SAS Institute Inc., Cary, NC, USA). A p-value <0.05 (two-sided) was considered statistically significant. For descriptive statistics, the growth inhibition scores for the lactobacilli at different doses are presented as the median score. The frequency of growth inhibition scores 0–3 for each *Lactobacillus* strain, for each dose, and for each *Candida* strain is presented in percentage distribution. A Poisson regression model was employed to test the influence of the *Lactobacillus* strains, *Candida* strains and dose variables on the growth inhibition score. Based on the results from the model, the four lactobacilli displaying the lowest growth inhibition scores were subjected to Chi-square tests.

**Results**

**Growth inhibition**

All the 14 *Lactobacillus* strains proved to inhibit the growth of the *Candida* spp., but some were more effective than others, with a substantial *Candida* spp. inhibition variation (Figure 1, Suppl. Table S1). The four best-performing lactobacilli were *L. rhamnosus* DSM 32992, which demonstrated the highest number of score 0 and 1, followed by *L. rhamnosus* DSM 32991, *L. jensenii* 22B42, and *L. rhamnosus* PB01. The difference between *L. rhamnosus* DSM 32992 and the other three strains was statistically significant (p < 0.001). In general, the growth inhibition was almost equal for the cell concentrations 10⁸ and 10⁷ CFU/mL, while the lowest concentration (10⁵ CFU/mL) appeared to be least effective (Suppl. Table S1, Suppl. Figure S1). The various *Candida* spp. displayed contrasting susceptibility; the growth of the clinical and control strains of *C. parapsilosis* was highly inhibited, while the two *C. krusei* strains presented no growth inhibition or only slight growth inhibition (Figure 2, Suppl. Table S1). The clinical *Candida* isolates and the control strains of the same species presented a similar susceptibility in the assays except for *C. tropicalis* for which the clinical isolate seemed to be more inhibited than the control strain.

**Microsensor pH measurements**

Microsensor measurements of pH were performed for the four above-mentioned best-performing *Lactobacillus* strains: *L. rhamnosus* DSM 32992,
L. rhamnosus DSM 32991, L. jensenii 22B42, and L. rhamnosus PB01, of which the two former were isolated from the oral cavity and the two latter had vaginal origin. The pH of the plain double-layered
agar was 5.8, and pH measurements from *L. rhamnosus* DSM 32992, *L. rhamnosus* DSM 32991, and *L. rhamnosus* PB01 generally showed lower pH values than *L. jensenii* 22B42 ranging from 3.5 to 3.8 (Suppl. Figure S2). In plates with only *Candida* spp. incubated, pH measured through dense colonies of *C. krusei* and *C. tropicalis* was approximately 6.8 at the surface of the colonies, reaching 5.8 around 700–1,000 µm down in the agar and remained stable. A similar picture was obtained when the pH-value was measured through *C. albicans* colonies, but the pH drop seemed to be more rapid, reaching 5.8 approximately 500 µm inside the agar. The measurements for *L. rhamnosus* DSM 32992 (10⁷ CFU/mL) through *C. albicans* and non-*C. albicans* colonies are presented in Figure 3. The surface pH was substantially lower compared with the *Candida* spp. incubated plates and the pH dropped rapidly through the first 400 µm. However, for the *C. krusei* and *C. tropicalis* reference strains, this drop was delayed and less dramatic (Figure 3). The pH profiles of the remaining three *Lactobacillus* strains inoculated with *Candida* spp. at the top agar layer showed very similar patterns (data not shown).

### Discussion

During recent years, the use of probiotic bacteria to control oral *Candida* colonization has been investigated as an adjunct therapeutic strategy to prevent infections in susceptible individuals [24,26,27]. In this study, we investigated the growth inhibition of 14 selected *Lactobacillus* strains isolated from the human oral cavity or vagina. We found a clear but strain-dependent effect on *Candida* growth which was in harmony with several previous *in vitro* experiments [9,17,26,28,29]. Interestingly, three of the best performing strains were *L. rhamnosus* strains, and the two best growth inhibitors, *L. rhamnosus* DSM 32992 and *L. rhamnosus* DSM 32991, were isolated from the oral cavity. *L. rhamnosus* is one of the most investigated probiotic *Lactobacillus* spp. and especially, the strain *L. rhamnosus* GG ATCC 53103 (LGG), isolated from a healthy human gastrointestinal tract, has shown remarkable probiotic properties [30,31]. Oral candidosis is a result of bacterial-fungal dysbiosis appearing when the microbial homeostasis is disrupted due to antibiotic treatment, local or systemic immune disturbances, smoking, hyposalivation, use of dentures, and poor oral hygiene [32]. To our

![Figure 3](image-url)

**Figure 3.** Micro-sensor measurement of pH with selected *Candida* strains with *L. rhamnosus* DSM 32992 (10⁷ CFU/mL) in the bottom agar layer of the co-cultured agar overlay assay. Zero on the vertical axis represents the first measurement from the sensor of either a *Candida* colony or agar. No changes in pH values appeared for any of the strains at deeper levels in the agar.
knowledge, only one randomized clinical trial (RCT) has investigated the antifungal effect of a single strain of L. rhamnosus in the oral cavity [22]. When added to milk, L. rhamnosus SP1 provided a significant reduction in the severity of denture stomatitis and significantly reduced Candida counts in the probiotic group after 6 months’ intervention [22]. Four RCT’s [19–21,33] have shown decreased Candida counts after intervention with different L. rhamnosus strains; however, in these trials the strains were combined with non-L. rhamnosus probiotic species, and the antifungal effect can therefore not completely be ascribed to one probiotic strain. The beneficial clinical effects are however supported by laboratory studies showing that L. rhamnosus strains can protect oral epithelial tissue from damage, caused by C. albicans, by impacting its major virulence factors, including adhesion, competition for nutrients, invasion and hyphae formation [26,30,34].

The probiotic mechanisms of action against C. albicans are not fully clear [27] but our present results suggest that some of the inhibiting effects are pH-driven. Similar observations have been made before [9,28]. It is thought that colonization of C. glabrata and C. albicans depends on their ability to cope with the presence of lactic and acetic acids produced by commensal microbiota [35]. Lactobacilli produce different weak organic acids (WOAs), mainly lactic and acetic acids, but the capability and rate of acid production are strain-specific [36]. In our study, L. rhamnosus DSM 32992 and L. rhamnosus DSM 32991 generally showed the lowest pH values indicating the highest acid production of the selected lactobacilli. The findings correspond well with the results from the growth inhibition test, in which these two strains exhibited the best inhibitory effect. This indicates that acid production plays an important role in the inhibition of Candida growth in agar overlay interference assay and indicates that the acids produced by the lactobacilli diffuse into the upper agar layer and interfere with Candida growth and survival. Cottier and co-workers [37] investigated the transcriptional stress response of C. albicans to WOAs in vitro and found the response to be a significant enrichment of genes involved in iron homeostasis, and down-regulation of RNA synthesis and ribosome biogenesis genes. The findings were mostly apparent upon chronic exposure to the acids and suggested that exposure of C. albicans to WOAs over time might shift the cells into a ‘starvation-like’ metabolic state with low transcription, translation and growth of the cell [37]. In a study by Köhler et al. [14], C. albicans growth was suppressed at low pH by the supernatant of the lactobacilli; however, pH neutralization of the culture filtrate completely abrogated the inhibitory effects of the supernatants. It is likely that low pH inhibits the transition between the blastospore form and the more invasive hyphae form of C. albicans and increases the intracellular concentration of protons in the yeasts. This may lead to an increased activity of an energy-consuming plasma membrane H+-ATPase that exhausts the available energy for growth and metabolism, leading to growth inhibition and, finally, cell death of the yeast [38]. Our experiments revealed that the tested C. krusei and C. tropicalis strains were able to resist and neutralize the acids produced by the lactobacilli. This may be understood in view of an in vitro study by Halm et al. [39] who found that lactic acid only induced a short-term (few seconds) pH change intracellularly in C. krusei. Fungal resistance to organic acids could be caused by extracellular production of ammonia [40,41], less permeable plasma membranes to lactic acid, higher buffer capacity inside the fungal cells, or higher H+-ATPase capacity [38].

Results from in vitro inference studies with single bacteria or multi-species biofilms must be considered with great caution taken the complex environment of the oral cavity into account. The agar overlay interference assay allows screening of multiple strains at a time on a single plate, with different Lactobacillus strains in the bottom agar laying at different cell concentrations. The assay is a relatively simple, yet a robust and well-proven method for screening purposes of probiotic bacteria against oral pathogens [9,13,25]. In our experiments, we modified the growth inhibition scoring system to allow for a more sensitive screening. Based on our previous experience [9], we separated the original score 1 (‘slight inhibition’) into score 1 ‘almost total growth inhibition (colonies slightly visible’) and score 2 ‘slight growth inhibition (colonies are clearly visible but smaller than at the control plate)’. The advantage of this modified system was that we were able to distinguish the growth inhibition capacity of the 14 probiotic candidates but the downside was that the results were harder to compare with the results found in comparable studies using the same inhibition assay [9,13,42].

There is evidence to support the hypothesis that the efficacy of probiotics is both strain-specific and disease-specific [43]. Therefore, our present the finding of two probiotic L. rhamnosus candidates (strain-specificity) with excellent antifungal capabilities (disease-specificity) isolated from the oral cavity, the environment where they are intended to exhibit their probiotic action (target site-specificity), was promising. Future studies on the characteristics of the four candidates, or combinations thereof, seems motivated to further unveil the probiotic and physiological properties including the risk of transmission of antibiotic resistance genes. In addition to clinically demonstrated health benefits and safety for human use, the persistence of cell viability and probiotic
activities throughout the processing, handling and storage needs to be investigated [31].

Conclusion

All 14 Lactobacillus strains employed in this in vitro screening demonstrated antifungal activity and inhibited the growth of six Candida spp. to varying degrees. The effect seemed pH-driven and two L. rhamnosus strains isolated from the oral cavity showed the strongest growth inhibition and acid production ability among the selected strains. The findings are promising for future clinical employment of these strains in the prevention of oral C. albicans and non-C. albicans infections; however, further investigations and characterization of the strains are needed to elaborate on their probiotic properties.

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Disclosure statement

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