Bifidogenic effect of salep powder

Salep tozunun bifidojenik etkisi

Buse USTA GORGUN 1, Lutfiye YILMAZ ERSAN 2

1 Uludag University, Faculty of Agriculture, Institute of Natural Sciences, Bursa-Turkey
2 Uludag University, Faculty of Agriculture, Department of Food Engineering, Bursa-Turkey

To cite this article:

Address for Correspondence:
Lutfiye YILMAZ ERSAN
e-mail: lutfiyey@uludag.edu.tr

Introduction

There is growing interest in functional foods having beneficial effects on the human health. Probiotics, prebiotics and synbiotics have gained an important to modulate the gut microbiota with the aim of improving host health. Probiotics have been defined by WHO and FAO as “viable non-pathogenic microorganisms which, when administered in adequate amounts, demonstrate beneficial effect on the human health”. The beneficial effects of the probiotics are inhibition the growth of the intestinal pathogens, improved lactose digestion, enhancing the immune system,
lowering the risk of cancer, treatment and prevention of obesity, diabetes, cardiovascular diseases and some allergic reactions. Although there are hundreds of probiotic species such as *Bifidobacterium*, *Lactobacillus*, *Lactococcus*, *Saccharomyces*, and *Enterococcus*. *Bifidobacterium* and *Lactobacillus* species are most commonly used probiotic species in food products (Bernardeau et al., 2008; Pan et al., 2009; Shokryazdan et al., 2017; Kerry et al., 2018; Zoumpopoulou et al., 2018).

*Bifidobacterium* species are gram-positive, high guanine+cytosine, catalase negative, rod-shaped, non-immobile, non-sporeulate, obligate anaerobes with the exceptions of a few species, highly sensitive to pH above 8.0 or below 4.5 with optimum growth between 37 to 41°C. *B. breve*, *B. bifidum*, *B. adolescentis*, *B. animalis*, *B. infantis*, *B. lactis* and *B. longum* are the most recognized species of Bifidobacteria used as probiotics. *Bifidobacterium* species consumed at adequate levels have demonstrated beneficial effects on treatment and/or prevention of some diseases namely concerning immune system, gastrointestinal infections, hypertension, cholesterol and cancer. Bifidobacteria constitute 95% of the total intestinal microbiota in the colons of breast-fed infants and decline with age and changes in eating habits (Champagne et al., 2005). Many attempts have been made to stimulate the growth of bifidobacteria and to increase their populations in the intestines of humans and/or animals and a food by supplying biologically active components called as “bifidogenic factor” or “bifidus factor” (Wang et al., 2010; González-Rodríguez et al., 2013). Bifidogenic factor was described by Modler (1994) and Gomes and Malcata (1999) as a carbohydrate that survive direct metabolism by the host and reach the colon where they are preferentially metabolised by Bifidobacteria as source of energy. Recently, this factor is called as a prebiotic. Prebiotics are non-digested, not-hydrolyzed and non-absorbed food ingredients in stomach and small intestine. They have been fermented by the gut microbiota that causes specific changes, both in the composition and/or activity in the gastrointestinal microbiota. As a result of fermentation, a) potentially health promoting metabolites such as lactic acid, short chain fatty acids, bacteriocins and B vitamins are formed, b) the colonic flora by exerting bifidogenic effect (the proliferation of bifidobacteria) have been modulated and c) the growth of undesirable bacteria has been inhibited. Thus, prebiotics have demonstrated health benefits including detoxification of toxic substances, reduction of blood ammonia levels and cholesterol, stimulation of mineral uptake and the treatment/prevention of cancer and infectious diseases (Bindels et al., 2015; Shigwedha et al., 2016; Carlson et al., 2018).

Cereals, legume crops, vegetables (chicory, onions, leek, artichoke etc.), fruits (dragon fruit, jack fruit etc.) and milk components (oligosaccharides) are natural sources for prebiotics. However, inulin, fructooligosaccharides, oligofructose, (trans-)galactooligosaccharides (TOS or GOS) or lactulose derived through biochemical and/or enzymatic techniques from these foods are the most used prebiotics at production of industrial products (Gibson and Rastall, 2006; Wang et al., 2010; Rios-Covian et al., 2013; Sousa et al., 2015; Underwood et al., 2015).

Synbiotic has been called as combining probiotic bacteria with prebiotics. In a synbiotic composition, prebiotics should be fermented selectively by the probiotic strain. Fermentation of the prebiotic results in a) increasing of beneficial microbiota (e.g., bifidobacteria and lactobacilli), b) decreasing the populations of potentially pathogenic microbiota (e.g., *Clostridium perfringens* and *E. coli*), c) lowering of the pH and d) production of short-chain fatty acids (SCFAs), mainly acetic, propionic and butyric acids. SCFAs might serve health promoting properties such as promoting the absorption of water and minerals including calcium, magnesium and iron, inhibiting cholesterol synthesis and the prevention/treatment of the metabolic syndrome, bowel disorders and certain types of cancer (Morrison and Preston, 2016;...
Orchids are mostly cultivated as beautiful flowers which have immense economic importance and medicinal value. The salep powder is obtained from grinding dried tubers of *Orchis mascula*, *Orchis militaris* and related species of orchids growing naturally in Turkey. Salep is used as a very nutritive ingredient in a delicious hot drink known by the same name, traditional Kahramanmaras-type of ice cream, soft drinks and confectionary industries. In traditional medicine, salep is used as an aphrodisiac in sexual activity and erectile dysfunction, as a blood sugar manager to relieve pancreas stress, in the prevention of chronic disorders, in treatment of bowel disorders, tuberculosis, diarrhea, Parkinson’s, cancer, fever and weight management (Jagdale et al., 2009; Altundag et al., 2012; Pourahmad, 2015; Jahromi et al., 2018). Although the components of salep vary according to the season of collection and orchid species, generally it contains mucilage (48%), moisture (12%), sugar (1%), starch (3%), nitrogenous substance (5%), ash (2%) and glucomannan (16%–60%). Glucomannan is a natural neutral water-soluble fiber which is important in normalizing blood sugar, relieving stress on the pancreas and preventing blood sugar abnormalities, such as hypoglycemia (Staiano et al., 2000; Keithley and Swanson, 2005; Onakpoya et al., 2014; Tester and Al-Ghazzewi, 2017).

Recently, researchers have studied to find innovative or alternative sources which are commercially available prebiotics. In this context, prebiotic properties of glucomannan have been reported by many researchers (Connolly et al., 2010; Muller et al., 2012; Harmayani et al., 2014). Salep powder may be metabolized by bifidobacteria and demonstrate bifidogenic as well prebiotic effect, probably due to its glucomannan content.

In order to develop candidate prebiotic substrates, firstly the bacterial fermentation properties of a substrate can be initially determined using *in vitro* models and its prebiotic activity can later be confirmed by controlled human clinical trials. The objective of this research was to determine the fermentability of salep powder by some bifidobacteria species. Glucose, a non-prebiotic simple carbon source, is able to metabolized by probiotic fastly and reach maximum growth more quickly, thus it was chosen as a positive control. The comparison was made in both basal media with salep powder and glucose as well as glucose. The growth of bifidobacteria species was tested by determining the optical density (OD) and pH values. In order to examine the biochemical activities of the species, high performance liquid chromatography (HPLC) analysis of lactic acid and short chain fatty acids (acetic, propionic and butyric acids) concentrations was also carried out.

**Materials and Methods**

**Bacterial strains**

The bacteria used in this study were *Bifidobacterium longum* subsp. *infantis*, *Bifidobacterium animalis* subsp. *lactis*, *Bifidobacterium longum* subsp. *longum* and *Bifidobacterium bifidum*. These strains were obtained from DSMZ (Deutsche Sammlung von Mikroorganismen und Zellkulturen GmbH, Braunschweig, Germany). These strains were activated according to the method suggested by DSMZ and incubated at 37°C using an anaerobic atmosphere generation system (Anaerocult A, Merck, Darmstadt, Germany).

**Salep sample**

Salep from orchids grown different parts of Turkey region was provided by Kadem Sahlepcilik (Istanbul, Turkey). Stock solutions of salep were prepared in distilled water and filter-sterilized using Millipore-Stericup-GP 0.45 µm (Kaplan and Hutkins, 2000; Mumcu and Temiz, 2014).

**Growth conditions**

Tryptone Peptone Yeast Extract (TPY) was used as the basal medium containing peptone...
(5.00 g l⁻¹), yeast extract (2.50 g l⁻¹), glucose (5.00 g l⁻¹), tween 80 (1.00 g l⁻¹), K₂HPO₄·3H₂O (2.00 g l⁻¹), MgCl₂ (0.50 g l⁻¹), ZnSO₄·7H₂O (0.20 g l⁻¹), CaCl₂ (0.15 g l⁻¹), FeCl₃·6H₂O (0.003 g l⁻¹) and L-cysteine HCl (0.50 g l⁻¹). The medium was sterilised at 121°C for 15 min. Sterile salep solutions were added into the basal TPY medium to obtain final concentrations of 0.5% (w/v) as positive control TPY with 0.5% (w/v) glucose was used. The assay was performed by adding 2% (w/v) of an overnight culture of Bifidobacterium subsp.

**pH measurement**

The pH of each sample was determined during fermentation using a pH-meter (pH 315i / SET; WTW, Germany).

**Growth measurement**

The optical density as cell density of bacteria was determined at 600 nm (OD₆₀₀) with a spectrophotometer (Shimadzu UV 1800, Kyoto, Japan) during fermentation. The corresponding sterile TPY solutions without bacteria were used as blanks for the absorbance measurements.

**Lactic acid and SCFA analyses**

To evaluate the efficiency of the fermentation of salep by the Bifidobacterium subsp., HPLC was performed. Lactic, acetic, butyric and propionic acids can be detected in the growth medium and quantified by HPLC (Shimadzu marka LC-20 AD, Japan, [17]). The HPLC equipment consisted of Transgenomics ORH-801 column and Refractive Index Detectors (RID) (Shimadzu, Kyoto, Japan) connected to a recorder. Samples were filtered through a 0.45 μm syringe filter prior to injection into the HPLC column. The injection volume was 20μl, 0.0025 N H₂SO₄ was used as the mobile phase, under a flow rate of 0.6 mL min⁻¹ at 65 °C (Anonymous, 2012).

**Statistical analysis**

All results were expressed as mean ± SD. Data was analyzed by two-way analysis of variance (ANOVA). Duncan’s Multiple Range Test was applied to determine the significant difference between the fermentation times and substrates used as carbohydrate source at P ≤ 0.01.

**Results and Discussion**

In order to determine the carbohydrate preferences of Bifidobacterium species, the optical density (OD) and pH were determined. Table 1 shows OD and pH values after the five fermentation times (0, 12, 24, 36 and 48 hours) in anaerobic culture following supplementation with glucose as a positive control and salep powder (test substrate).

The results were presented as the mean value of each fermentation time (FT), regardless of the Bifidobacterium species (BS), and the mean value of each strain, regardless of the fermentation time. For all parameters, the interaction among both factors (BSxFT) could be determined, and for all parameters and every time, a significant interaction was detected (P<0.01). Regarding the fermentation time, the highest value of optical density was recorded for media with glucose, while it showed pH value. In glucose samples, the pH decreased significantly (P<0.01) during 48 h of fermentation. It was found that pH value for salep powder samples showed a temporal increase and decrease between 36 h and 48 h. B. infantis has the lowest pH values in both media with glucose (4.52) and salep powder (5.25).

As shown in Table 1, Bifidobacterium species used salep powder as a source of carbon and energy. All strains grew well on glucose, the highest OD values were found for the strains B. lactis in media containing glucose (1.114) and B. infantis in media containing salep powder (0.952). Optical density value when salep powder for B. infantis was used as a single carbon source was able to generate similar growth as that promoted by glucose (the traditional carbon source). Wang et al. (2010) reported that B. adolescentis displayed the highest growth on xylooligosaccharides with the maximum OD of 1.68 and the lowest pH value of 5.1, followed by B. longum and B. bifidum, however, B. breve did
not show any growth on XOS. García-Cayuela et al. (2014) determined that the OD values after 48 h growth of B. lactis BB-12, B. breve 26M2 and B. bifidum HDD541 with the six carbohydrates (glucose, lactulose, lactosucrose, kojibiose, lactulosucrose and 40-galactosyl-kojibiose) used as carbon sources ranged from 0 to 1.21. On the other hand, the maximum OD for glucose values were obtained at 48 h of fermentation, while at 36 h of fermentation for salep powder. Voragen (1998) reported that chemical structure, degrees of polymerization (DP), composition of monomer units and water solubility of substrate affect their utilisation by probiotic microorganisms.

Table 1 The mean values of OD and pH values obtained by four Bifidobacterium species grown on glucose and salep powder for 48 h

<table>
<thead>
<tr>
<th>Bifidobacterium strains (Bifidobacterium türleri)</th>
<th>N</th>
<th>OD (Optik yoğunluk)</th>
<th>pH</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>Glucose (Glikoz)</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>Salep powder (Salep tozu)</td>
<td></td>
</tr>
<tr>
<td>B. bifidum</td>
<td>10</td>
<td>0.901&lt;sup&gt;Aa&lt;/sup&gt;</td>
<td>4.70&lt;sup&gt;Ab&lt;/sup&gt;</td>
</tr>
<tr>
<td></td>
<td></td>
<td>0.682&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>5.52&lt;sup&gt;As&lt;/sup&gt;</td>
</tr>
<tr>
<td>B. infantis</td>
<td>10</td>
<td>0.951&lt;sup&gt;As&lt;/sup&gt;</td>
<td>4.52&lt;sup&gt;Cb&lt;/sup&gt;</td>
</tr>
<tr>
<td></td>
<td></td>
<td>0.952&lt;sup&gt;As&lt;/sup&gt;</td>
<td>5.25&lt;sup&gt;Cs&lt;/sup&gt;</td>
</tr>
<tr>
<td>B. lactis</td>
<td>10</td>
<td>1.114&lt;sup&gt;Ad&lt;/sup&gt;</td>
<td>4.89&lt;sup&gt;Ab&lt;/sup&gt;</td>
</tr>
<tr>
<td></td>
<td></td>
<td>0.645&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>5.59&lt;sup&gt;Es&lt;/sup&gt;</td>
</tr>
<tr>
<td>B. longum</td>
<td>10</td>
<td>0.703&lt;sup&gt;Ba&lt;/sup&gt;</td>
<td>3.98&lt;sup&gt;As&lt;/sup&gt;</td>
</tr>
<tr>
<td></td>
<td></td>
<td>0.619&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>5.90&lt;sup&gt;Es&lt;/sup&gt;</td>
</tr>
<tr>
<td>Fermentation time (hours)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>0</td>
<td>8</td>
<td>0.077&lt;sup&gt;Ba&lt;/sup&gt;</td>
<td>3.72&lt;sup&gt;Ad&lt;/sup&gt;</td>
</tr>
<tr>
<td></td>
<td></td>
<td>0.113&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>6.32&lt;sup&gt;Es&lt;/sup&gt;</td>
</tr>
<tr>
<td>12</td>
<td>8</td>
<td>0.831&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>5.12&lt;sup&gt;As&lt;/sup&gt;</td>
</tr>
<tr>
<td></td>
<td></td>
<td>0.608&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>5.69&lt;sup&gt;Cs&lt;/sup&gt;</td>
</tr>
<tr>
<td>24</td>
<td>8</td>
<td>1.152&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>4.46&lt;sup&gt;Ab&lt;/sup&gt;</td>
</tr>
<tr>
<td></td>
<td></td>
<td>0.874&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>5.35&lt;sup&gt;Cs&lt;/sup&gt;</td>
</tr>
<tr>
<td>36</td>
<td>8</td>
<td>1.194&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>4.20&lt;sup&gt;Ad&lt;/sup&gt;</td>
</tr>
<tr>
<td></td>
<td></td>
<td>1.092&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>5.17&lt;sup&gt;Ds&lt;/sup&gt;</td>
</tr>
<tr>
<td>48</td>
<td>8</td>
<td>1.334&lt;sup&gt;Bb&lt;/sup&gt;</td>
<td>4.16&lt;sup&gt;Cb&lt;/sup&gt;</td>
</tr>
<tr>
<td></td>
<td></td>
<td>0.935&lt;sup&gt;Ca&lt;/sup&gt;</td>
<td>5.25&lt;sup&gt;Ds&lt;/sup&gt;</td>
</tr>
<tr>
<td>ANOVA</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Bifidobacterium strains (BS)</td>
<td>**</td>
<td>**</td>
<td>**</td>
</tr>
<tr>
<td>Fermentation time (FT)</td>
<td>**</td>
<td>**</td>
<td>**</td>
</tr>
<tr>
<td>BSKFT</td>
<td>**</td>
<td>**</td>
<td>**</td>
</tr>
</tbody>
</table>

<sup>A-D</sup>Different uppercase superscripts in the same column depict the significant difference between means for fermentation time and Bifidobacterium strains (<P<0.01)

<sup>a-d</sup>Different lowercase superscripts in the same row depict the significant difference between glucose and salep powder (<P<0.01)

**Lactic acid**

Figure 1. Changes of lactic acid values of samples after the 48 h-fermentation

Şekil 1. 48 saatlik fermantasyon sonunda örneklerdeki laktik asit miktarındaki değişim

<sup>a-d</sup>Different lowercase superscripts depict the significant difference among samples (<P<0.01);<br/>

<sup>A-D</sup>Different uppercase superscripts depict the significant difference between Bifidobacterium subsp. (<P<0.01)
Lactic acid is the main end-product of the glycolytic fermentation produced by lactic acid bacteria. For all the tested strains, lactic acid was the most abundant produced metabolite, this finding is in agreement with LAB metabolism. Figure 1 showed the lactic acid concentrations of Bifidobacterium species in media including glucose and salep powder. The concentration of lactic acid ranged from 0.19 g l⁻¹ of B. lactis in media with salep powder to the 1.37 g l⁻¹ of B. infantis in media with glucose, depending on the strain and type of substrate. More lactic acid was produced from glucose than salep powder. B. infantis produced the higher lactic acid in media with salep powder than other species. Barczynska et al. (2012) reported that lactic acid content determined 109.3 mg 100 mL⁻¹ for B. bifidum Bb12 and 108.8 mg 100 mL⁻¹ for B. animalis DN–173 010 in the broth containing the tartaric acid-dextrin as the only one source of carbon after 24-h incubation.

Short chain fatty acids

Short-chain fatty acids (SCFAs) such as acetic, propionic and butyric acids are produced as the end products of the microbial fermentation of dietary carbohydrates via Bifidobacteria and Lactobacilli. The type number and enzymatic capabilities of microorganisms and substrate sources have affected amount and type of these metabolites’ in vitro media (Wong et al., 2006). Also, in human body endogenous colonic microbiota like probiotic bacteria and gut transit time have affected production (Barczyńska et al., 2015; Pessone et al., 2015).

Acetic acid

![Graph showing acetic acid values](image)

Figure 2. Changes of acetic acid values of samples after the 48 h-fermentation

Acetic acid, the most abundant SCFA produced by the human colonic microbiota, can be used as energetic substrate in muscle tissue, controlling inflammation and counteracting pathogen invasion (Fukuda et al., 2011). The acetic acid concentrations of Bifidobacterium species were shown in Figure 2. Its concentration depended on the strain and substrate type. Acetic acid, it ranged from 0.19 g l⁻¹ of B. longum to 1.20 g l⁻¹ of B. infantis in media with glucose. The higher acetic acid values were observed for B. bifidum and B. longum in media with salep powder than glucose. Generally, during the whole fermentation progress, samples including salep powder exhibited a lower acetic acid value than glucose. It was found that acetic acid content determined 19.1 mg 100 mL⁻¹ for B. bifidum Bb12 and 16.9 mg 100 mL⁻¹ for B. animalis DN–173 010 in the broth containing the tartaric acid-dextrin as the only one source of carbon after 24-h incubation (Barczynska et al., 2012).

Propionic acid produced by certain colon bacteria can be transported to many parts of the
body and utilized in liver cells (Hongpattarakere et al., 2012). It could be observed (Figure 3) that the fermentation time might influence the propionic acid values of Bifidobacterium species fermented in media including glucose and salep powder. Propionic acid contents ranged from 0.01 g L\(^{-1}\) of B. longum to 0.13 g L\(^{-1}\) of B. lactis in media with salep powder. For Bifidobacterium species, the maximum propionic acid value of B. lactis fermented in media with salep powder was observed. Barczynska et al. (2012) reported that propionic acid content determined 4.5 mg 100 mL\(^{-1}\) for B. bifidum Bb12 and 4.6 mg 100 mL\(^{-1}\) for B. animalis DN–173 010 in the broth containing the tartaric acid-dextrin as the only one source of carbon after 24-h incubation.

![Propionic acid](image)

Figure 3. Changes of propionic acid values of samples after the 48 h-fermentation

Different lowercase superscripts depict the significant difference among samples (\(P<0.01\));

Different uppercase superscripts depict the significant difference between Bifidobacterium subsp. (\(P<0.01\))

Butyric acid among SCFAs displays beneficial health effects such as the prevention and treatment of colonic diseases (Fung et al., 2012; Pessione et al., 2015). The Bifidobacterium species and type of substrate affected the butyric acid values. Generally, higher butyric acid values were present in media with salep powder. A maximum butyric acid value of B. lactis fermented in media with salep powder was observed, while B. bifidum and B. lactis fermented in media with
glucose had a lower butyric acid value.

Interestingly the trends of production of propionic and butyric acids were similar for the tested strains, the highest producers of the both SCFAs were for B. lactis in media with salep powder, while and the lowest producers were B. longum in media with salep powder. These results agreed with Pessione et al. (2015) who studied characterization of potentially probiotic lactic acid bacteria isolated from olives and evaluation of short chain fatty acids production.

Conclusion

In this study, for the evaluation of bifidogenic effect by growth and metabolic activity achievements, growth curves were monitored by optical density assessment, and bacterial metabolism was assessed by pH measurement and as well as determination of short chain fatty acids production. Based on the results reported herein, salep powder may stimulate the growth of Bifidobacterium species. Enhancement of Bifidobacterium species growth and metabolism by the increase of lactic acid and short chain fatty acids production that it may be stated that salep powder has potential as “bifidus factor”

Acknowledgements

This study was funded in part by the Commission of Scientific Research Projects of Uludag University, Bursa, Turkey (QUAP (Z) 2013/50). This study is a part of Buse Usta’s M.Sc. thesis submitted to the Uludag University, Institute of Natural Sciences.

References


