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β-Galactosidase from *Lactobacillus pentosus*: Purification, characterization and formation of galacto-oligosaccharides

THOMAS MAISCHBERGER^{1,2}, ELISABETH LEITNER¹, SUNEE NITISINPRASERT³, ONLADDA JUAJUN^{1,4}, MONTAROP YAMABHAI⁴, THU-HA NGUYEN^{1,2,*} AND DIETMAR HALTRICH¹

¹ Food Biotechnology Laboratory, BOKU University of Natural Resources and Applied Sciences,

Vienna, Austria

² Research Centre Applied Biocatalysis, Graz, Austria

³ Department of Biotechnology, Kasetsart University, Bangkok, Thailand

⁴ School of Biotechnology, Suranaree University of Technology, Nakhon Ratchasima, Thailand

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* **Correspondence**: Dr. Nguyen Thu-Ha (e-mail: thu-ha.nguyen@boku.ac.at). Address: Lebensmittelbiotechnologie, Department für Lebensmittelwissenschaften und technologie, Universität für Bodenkultur Wien, Muthgasse 18, A-1190 Vienna, Austria. Phone: 43-1-47654 6146. Fax: 43-1-47654 6251.

Abbreviations: β -Gal, β -galactosidase; CE, capillary electrophoresis; GOS, galactooligosaccharides; HPAEC-PAD, high-performance anion exchange chromatography with pulsed amperometric detection; MUG, 4-methylumbelliferyl- β -D-galactopyranoside; *o*NPG, *o*nitrophenyl- β -D-galactopyranoside, *o*NP, *o*-nitrophenol

Abstract

A novel heterodimeric β-galactosidase with a molecular mass of 105 kDa was purified from the crude cell extracts of the soil isolate Lactobacillus pentosus KUB-ST10-1 using ammonium sulphate fractionation followed by hydrophobic interaction and affinity chromatography. The electrophoretically homogenous enzyme has a specific activity of 97 U_{oNPG}/mg protein. The K_m, k_{cat} and k_{cat}/K_m values for lactose and *o*NPG were 38 mM, 20 s⁻¹, 530 M⁻¹·s⁻¹ and 1.67 mM, 540 s⁻¹, 325,000 M⁻¹·s⁻¹, respectively. The temperature optimum of β -galactosidase activity was 60–65 °C for a 10-min assay, which is considerably higher than the values reported for other lactobacillal β -galactosidases. Mg²⁺ ions enhanced both activity and stability significantly. L. pentosus β-galactosidase was used for the production of prebiotic galacto-oligosaccharides (GOS) from lactose. A maximum yield of 31% GOS of total sugars was obtained at 78% lactose conversion. The enzyme showed a strong preference for the formation of β -(1 \rightarrow 3) and β -(1 \rightarrow 6) linkages, and the main transgalactosylation products identified were the disaccharides β -D-Galp-(1 \rightarrow 6)-D-Glc, β -D-Galp-(1 \rightarrow 3)-D-Glc, β -D-Galp-(1 \rightarrow 6)-D-Gal, β -D-Galp-(1 \rightarrow 3)-D-Gal, and the trisaccharides β -D-Galp-(1 \rightarrow 3)-D-Lac, β -D-Galp-(1 \rightarrow 6)-D-Lac.

1 Introduction

The lactic acid bacterium *L. pentosus* is frequently used as starter culture for silage fermentations and various food fermentation processes, including sauerkraut and raw sausage production, or olive and tea leave fermentations, to name a few. Most of *L. pentosus* subspecies are plant isolates, generally recognised as safe (GRAS), and capable of metabolising different pentoses such as ribose, arabinose and xylose [1]. Yet, some subspecies of *L. pentosus* were isolated from e.g. human faeces [2] or from raw milk [3], which were proposed as potentially probiotic strains. Furthermore, Fred *et al.* previously described very effective growth of *L. pentosus* on lactose, which could imply high β-galactosidase activity [4], but this property has not been studied to date.

Lactic acid bacteria (LAB) and β -galactosidases have been studied extensively in the past, both from a fundamental and an applied point of view. β-Galactosidases from e.g. Bifidobacterium bifidum, B. infantis [5], Carnobacterium piscicola [6], L. acidophilus [7], L. bulgaricus [8], L. reuteri [9], and Leuconostoc lactis [10] were cloned and characterised to a varying detail. Despite of the importance of LAB and in particular Lactobacillus spp. for food technology and dairy applications and despite of numerous studies on the gene clusters involved in lactose utilisation, only few lactobacillal β-galactosidases were characterised in detail pertaining to their biochemical properties or were investigated for their ability to produce galactooligosaccharides (GOS) in biocatalytic processes [11-13]. These carbohydrate-based food ingredients are of interest as novel and possibly improved prebiotics, or for the development of synbiotic functional food that could introduce new dimensions of applications [14].

GOS belong to the prebiotics, which are defined as a "selectively fermented ingredient that allows specific changes, both in the composition and/or activity in the

gastrointestinal microbiota that confers benefits upon host well-being and health" [15]. In practice, the beneficial bacteria that serve as targets for prebiotics and GOS are almost exclusively bifidobacteria and lactobacilli. An advantage of the concept of prebiotics to modify gut function is that the target bacteria are already commensal to the large intestine, whereas with probiotics allochthonous micro-organisms are introduced and have to compete against established colonic communities [16]. GOS have been classified as one of the few proven prebiotics fulfilling the three criteria (i) resistance to gastric acidity, hydrolysis by mammalian enzymes and gastrointestinal absorption; (ii) fermentation by intestinal microflora; and (iii) selective stimulation of the growth and/or activity of intestinal bacteria associated with health and well-being [15]. GOS have been widely investigated for their prebiotic properties and physiological effects using in vitro, animal, human, and infant studies, and functional effects of GOS on human health are summarised in [16], [20] and [21].

GOS are produced from lactose using β-galactosidases (E.C. 3.2.1.23), which in addition to their hydrolytic activity catalyse glycosyl transfer reactions. In this reaction galactosyl residues are transferred onto suitable acceptors such as another sugar
(lactose, the primary reaction products glucose and galactose, or other oligosaccharides formed), thus building up complex series of higher oligosaccharides [17-19]. The spectrum of the oligosaccharides making up these mixtures strongly depends on the source of the β-galactosidase used for the biocatalytic reaction, and on the conversion conditions used in their production. Since these differences in GOS spectrum and yields are a result of structural and/or mechanistic differences between β-galactosidases from various strains can be of significant interest [18]. Rabiu *et al.* [22] and Tzortzis *et al.* [23] synthesised a range of GOS mixtures from lactose using β-galactosidases from different bifidobacteria. Subsequently it was

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shown that these different mixtures typically resulted in better growth of that strain that had served as the source of the enzyme for GOS production. This concept may serve as the basis for a new generation of functionally enhanced, targeted oligosaccharides, and has increased the interest in β -galactosidases from beneficial probiotic organisms.

The objective of this work was to study the β-galactosidase from *L. pentosus* in detail and to compare it to other lactobacillal β-galactosidases isolated from organisms of typical animal origin. Recent data are indicating that *L. pentosus* might be of interest as a beneficial LAB making it attractive as a probiotic strain, e.g. based on immunomodulating properties [24] or the reduction of pathogens in animals [25].

2 Material and Methods

2.1 Strain and culture conditions

Lactobacillus pentosus KUB-ST10-1 was from the culture collection of Kasetsart University (Bangkok, Thailand). It was isolated from soil of a dairy farm in Thailand and identified by using both the API 50 CH carbohydrate fermentation stripes test (bioMérieux, Inc., Marcy l'Etoile, France) and sequence analysis of the amplified chromosomal 16S rDNA. It was stored at −70 °C in MRS broth medium (Merck, Darmstadt, Germany) containing 15% glycerol. To prepare fresh inocula, it was transferred twice after an 18-h cultivation period to fresh MRS broth medium supplemented with lactose (final concentration of 2%).

L. pentosus KUB-ST10-1 was grown anaerobically in 10 L of MRS broth medium containing 2% lactose at 37 °C with slight agitation and without pH control. A preculture, grown overnight in MRS medium (2% lactose (w/v)), was used to inoculate the fresh medium to a final concentration of 1% (v/v) inoculum. Cells were

95 harvested at an optical density (OD₆₀₀) of 13 by centrifugation at 8800 x g for 15 min at 4 °C.

2.2 Enzyme purification

β-Galactosidase was purified from the crude extract using a previously described protocol [9]. Wet biomass (100 g) was resuspended in 200 mL of sodium phosphate buffer (50 mM, pH 6.5) and homogenized (APV-2000; Silkeborg, Denmark) to disrupt the cells. Cell debris was removed by ultracentrifugation (36000 x g, 30 min, 4 °C) to obtain the crude enzyme extract.

Ammonium sulphate precipitation (ASP). Ammonium sulphate was added in small portions to the crude extract to a final concentration of 50% saturation under continuous stirring at 4 °C. The precipitated protein was collected by centrifugation (6200 x g, 15 min, 4 °C), and the pellet was re-dissolved in 50 mM sodium phosphate buffer (pH 6.5) containing 1 M ammonium sulphate (buffer A).

Hydrophobic Interaction Chromatography (HIC). The dissolved pellet was loaded onto a 20-mL phenyl-sepharose fast flow column (50 mm x 200 mm; Amersham,
Uppsala, Sweden) that had been pre-equilibrated with buffer A. Proteins were eluted by a linear gradient of 0% to 100% buffer B (50 mM sodium phosphate buffer, pH 6.5) over 10 column volumes (CV) and at a flow rate of 1.5 mL/min. Fractions with high β-galactosidase activity were pooled and concentrated by ultrafiltration (30-kDa cut-off; Amicon, Beverly, MA).

115 Affinity Chromatography (AC). The concentrated protein solution was applied to a 5mL column containing crosslinked (4%) beaded agarose immobilized with *p*aminobenzyl-1-thio-β-D-galactopyranoside (Sigma, St. Louis, MO) that was preequilibrated with 50 mM phosphate buffer pH 6.5. The enzyme was eluted using a flow rate of 0.5 mL/min and a linear gradient (0 to 1 M NaCl in the same buffer) over

10 CV. Active fractions were pooled, desalted and concentrated by ultrafiltration. The purified enzyme was stored in 50 mM sodium phosphate buffer (pH 6.5) at 4 °C.

2.3 Enzyme assays

β-Galactosidase activity was determined using both the artificial chromogenic substrate *o*-nitrophenyl-β-D-galactopyranoside (*o*NPG) and the natural substrate 125 lactose as described previously [9]. When *o*NPG was the substrate, the reaction was performed in 50 mM sodium phosphate buffer (pH 6.5) and using a 10-min incubation at 30 °C, after which the reaction was stopped. One unit of *o*NPG activity (U_{*o*NPG}) is defined as the amount of enzyme releasing 1 µmol of *o*NP per minute under the assay conditions described above. When lactose was used as the substrate, the 130 reaction was done in 50 mM sodium phosphate buffer (pH 6.5) for 10 min at 30 °C, after which the reaction was stopped. The release of D-glucose was determined colourimetrically using the glucose oxidase/peroxidase (GOD/POD) assay [26]. One unit of lactase activity (U_{Lac}) is defined as the amount of enzyme releasing 1 µmol of D-glucose per minute under the conditions listed above.

42 135 **2.4 Protein analysis and gel electrophoresis**

Protein concentrations were determined by the method of Bradford [27] using bovine serum albumin as the standard. The PhastSystem unit (Amersham) and precast polyacrylamide gels (Phastgel 8–25, Amersham) were used for performing native and SDS-PAGE. Samples were treated as described by Laemmli [28] with slight modifications. The enzyme was diluted to 1–2 mg of protein per mL and incubated with 2x Laemmli buffer at 60 °C for 5 min. Protein bands were stained using either Coomassie Blue staining or 4-methylumbelliferyl-β-D-galactopyranoside (MUG) as substrate for active staining. For SDS-PAGE and native PAGE molecular masses

were estimated using the Precision Plus Protein Dual Color Kit (Biorad) and the HighMolecular Weight calibration Kit (Amersham), respectively.

2.5 Steady-state kinetic measurement

Catalytic constants were determined at 30 °C and pH 6.5 for the two substrates *o*NPG and lactose with their concentrations in 50 mM phosphate buffer varying from 0.1–22 mM and 0.1–600 mM, respectively. The kinetic parameters were calculated by non-linear regression, and the obtained data were fit to the Henri-Michaelis-Menten equation (using SigmaPlot, SPSS Inc., Chicago, IL, USA).

2.6 Effects of temperature and pH on enzyme activity and stability

The temperature dependence of β-galactosidase activity (both *o*NPG and lactose) was determined by assaying the enzyme samples over the range of 20 to 70 °C for 10 min. To study thermal stability, enzyme samples were incubated in 50 mM sodium phosphate buffer, pH 6.5, at 4, 23, 30, 37, and 42 °C. At certain time intervals samples were taken and the residual *o*NPG activity was measured. For measuring the pH dependence of β-galactosidase activity three buffer systems (sodium citrate, 50 mM, pH 4.0–5.5; sodium phosphate, 50 mM, pH 5.5–7.5; glycine, 50 mM, pH 7.5– 9.0) were used. For the determination of the pH stability enzyme samples were incubated at various pH values and 37 °C for up to 72 h, and the remaining enzyme activity was measured at certain time intervals using *o*NPG as substrate under standard assay conditions.

2.7 Substrate specificity

To determine the substrate specificity of the *L. pentosus* β-galactosidase for various structurally related substrates *p*-nitrophenyl-β-D-mannopyranoside, *p*-nitrophenyl-β-D-

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xylopyranoside, *o*-nitrophenyl- β -D-glucopyranoside and *p*-nitrophenyl- β -Dcellobioside were used. Activities were determined in accordance to standard *o*NPG assay conditions using the respective substrate in concentrations of 22 mM.

2.8 Transgalactosylation activity

Purified β -galactosidase from *L. pentosus* (5 U_{oNPG} per mL reaction mixture) was used for the discontinuous formation of galacto-oligosaccharides (GOS) at 30 °C using 205 g/L of lactose in 50 mM sodium phosphate buffer (pH 6.5, 2 mM MgCl₂) as substrate. Samples were taken at various intervals and the composition of the GOS mixture was analyzed by capillary electrophoresis (CE) and high-performance anion exchange chromatography with pulsed amperometric detection (HPAEC-PAD) as described previously [13].

3 Results

3.1 Formation and purification of β-galactosidase from *L. pentosus*

L. pentosus KUB-ST10-1 was selected for detailed studies on its β-galactosidase since this strain grew very well on lactose, it showed high β-gal activity in a screening, in which several lactic acid bacteria were compared pertaining to their ability to form this enzyme activity (data not shown), and since its β-galactosidase had not been studied in any detail previously. Batch cultivations of *L. pentosus* were carried out in a 10-L laboratory fermenter at 37 °C without pH control using lactose-based MRS medium. The biomass was harvested after 18 h of growth when β-gal activity reached a maximum of 644 units (U_{oNPG}) per litre fermentation broth. β-Galactosidase was purified to apparent homogeneity from the crude cell extract using a two-step purification protocol based on hydrophobic interaction and affinity

chromatography. The enzyme was purified 16-fold to a specific activity of 97 U/mg of protein, using standard assay conditions with *o*NPG as the substrate.

3.2 Properties of β-galactosidase from *L. pentosus*

L. pentosus β -galactosidase is a heterodimer (M_r ~107 kDa) consisting of a small (~35 kDa) and large (~72 kDa) subunit as judged by SDS-PAGE (Figure 1A) and native PAGE (Figure 1B). The additional protein band (~80 kDa) in native PAGE when staining with Coomassie Blue presumably results from degradation of the intact ~107-kDa enzyme, as was previously shown for the related β -gal from *L. reuteri* by using mass spectrometry [9]. Active staining on native PAGE with MUG yielded only one band with β -galactosidase activity, corresponding to the larger band in Figure 1B. Active staining on the SDS-PAGE gel after treating the enzyme in SDS buffer at 60° for 5 min, thus separating the two subunits without unfolding, showed that one band corresponding to the large subunit exhibited activity with MUG (not shown). In contrast, the small subunit by itself did not show any activity. When the enzyme was treated in SDS buffer at ~100 °C for 5 min, no activity could be detected after active staining on the gel.

3.3 Substrate specificity

To study the substrate specificity of *L. pentosus* β -galactosidase various structurally related artificial substrates were tested under standard assay conditions and 210 compared to *o*NPG. No appreciable activity was detected when using *p*-nitrophenyl- β -D-xylopyranoside, *o*-nitrophenyl- β -D-glucopyranoside, *p*-nitrophenyl- β -Dmannopyranoside and *p*-nitrophenyl- β -D-cellobioside as substrates for the activity assay.

3.4 Enzyme kinetics

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Steady-state kinetic studies were performed with the preferred substrates lactose and *o*NPG and the kinetic constants K_m, v_{max}, and k_{cat} were determined. The k_{cat} values were calculated on the basis of theoretical v_{max} values obtained by nonlinear regression using SigmaPlot. Data are summarized in **Table 1**. The catalytic efficiencies (k_{cat}/K_m) for the two substrates, *o*NPG and lactose, indicate that *o*NPG is the preferred substrate in accordance with other microbial β-galactosidases.

3.5 Effects of pH and temperature

The pH optimum of *L. pentosus* β -galactosidase activity is pH 8.0 and pH 7.5 in glycine buffer when using lactose and oNPG as substrate, respectively (Figure 2). In general, the enzyme is more active in the alkaline range (pH 7.0-8.5), and it showed a pronounced effect of the buffer system used on activity. The effect of different buffers and pH values on stability was investigated subsequently in more detail. Catalytic or kinetic stability, *i.e.*, the length of time an enzyme remains active before undergoing irreversible inactivation, was measured for various pH values at a constant temperature of 37 °C, and the inactivation constants (k_{in}) and the half-life times of denaturation ($\tau_{1/2}$) were determined (**Table 2**). β -Galactosidase showed firstorder inactivation kinetics at all pH values studied when analysed in the In(residual activity) versus time plot (not shown). Highest $\tau_{1/2}$ values were calculated in glycine buffer (pH 8.0 and 8.5) where the stability was increased up to 3.2-fold compared to when using sodium phosphate buffer at pH 6.5. Highest inactivation constants and hence least stabilities were obtained in sodium acetate buffer (pH 4.0 and 5.0) and glycine buffer (pH 9.0), with $\tau_{1/2}$ values of less than 1.5 h.

The optimum temperature measured under standard assay conditions (10 min) for lactose and *o*NPG hydrolysis was 60–65 °C and 55 °C, respectively (**Figure 3**). The effect of temperature on stability of β -galactosidase from *L. pentosus* was

240 investigated at the pH value of milk, pH 6.5, both in the absence and presence of 2 mM MgCl₂. Inactivation constants and half-life times were calculated as described above. The addition of 2 mM MgCl₂ to the buffer improved stability significantly, increasing the half life times $\tau_{1/2}$ of the enzyme at 30, 37 and 42 °C 3.2-, 4.9- and 5.6fold, respectively (**Table 3; Figure 4**).

3.6 Transgalactosylation activity

 In order to assess the potential of the novel β-galactosidase from *L. pentoses* for the production of GOS, a discontinuous lactose conversion process was studied at 30 °C, using an initial lactose concentration of 205 g/L in 50 mM sodium phosphate buffer (+ 2 mM MgCl₂, pH 6.5) and 5 U_{oNPG}/mL of β-galactosidase activity to produce GOS. The maximum GOS yield (31%) was obtained at a lactose conversion of 78% (**Figure 5**). The newly formed sugar mixture contained 31.5% glucose, 15.9% galactose, 21.9% lactose, 11.4% non-lactose disaccharides, 18.7% trisaccharides and 0.6% tetrasaccharides as analysed by capillary electrophoresis and high-performance anion exchange chromatography with pulsed amperometric detection, and using authentic sugars and the standard addition technique for comparison. The main GOS products identified were non-lactose disaccharides mainly of the β-(1→3) and β-(1→6) linkage type as well as the two trisaccharides β-D-Gal*p*-(1→3)-lactose, and β-D-Gal*p*-(1→6)-lactose.

4 Discussion

The genus *Lactobacillus* is the largest among the *Lactobacteriaceae* and includes more than a hundred described species [29]. Lactobacilli are nutritionally fastidious, metabolize a varying range of different carbohydrates, and are associated with a large variety of animals and plants. They are extensively used for fermentations of Page 13 of 31

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dairy products, meat, fish and plant material such as vegetables, and thus have been investigated intensively for their industrial applications [30]. Lactose utilisation is a primary function of lactobacilli and other LAB used in the fermentation of milk. The mechanism by which lactose is transported into the cell determines largely the subsequent pathway for the hydrolysis of this disaccharide. In several Lactobacillus species lactose is transported via phosphotransferase systems, which results in phosphorylation of lactose concomitant with its uptake. The resulting lactose-6'phosphate is then hydrolysed by phospho- β -galactosidase. Alternatively, lactose is taken up by secondary transport systems, and lactose is further metabolised by β galactosidase within the cell [31, 32]. While the organisation of these lactose genes, which are often forming operons or operon-like structures with modular organisation, has been studied in much detail, the properties of some of the enzymes encoded by the genes organised in these operons, including β -galactosidases, have received significantly less attention. It is thus surprising that β -galactosidases from lactobacilli, which play an eminent role in the metabolism of the principal carbohydrate in milk, have only been characterised in detail pertaining to their physical and biochemical properties in a few instances, and little attention has been paid to the catalytic ability of these enzymes with respect to their transglycosylation reactions [12, 17, 33]. Recently, we reported on the comprehensive characterisation of the β -galactosidases from L. reuteri and L. acidophilus, and pointed out their applicability for the biocatalytic production of GOS [9, 13, 34, 35]. In this paper, we present the detailed biochemical characterisation and the transgalactosylation activity of a novel βgalactosidase from L. pentosus. The strain KUB-ST10-1 used throughout this research was isolated from a soil sample in Thailand, and was selected for further characterisation based on a screening and its high β-galactosidase activity when

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grown on lactose. It was of interest to determine whether a β -galactosidase from a

 source other than those traditionally used in the dairy industry may have unique properties or characteristics. In a recent phylogenetic analysis based on 16S rRNA genes the genus *Lactobacillus* was grouped into five major divisions [36]. In this grouping, *L. acidophilus* belongs to group A and *L. reuteri* to group B, while *L. pentosus* is a member of group C [36], hence might differ in its metabolic activities from the sources of β-galactosidases recently characterised in our group.

The structural features of β -galactosidases from lactobacilli are varying to a great extent. While heterodimeric structures are in general less frequently encountered among microbial β -galactosidases, these seem to prevail among Lactobacillus β -galactosidases. Lactobacillal heterodimeric *β*-galactosidases are encoded by two genes, lacL and lacM, and have been described in e.g., L. acidophilus [34, 37], L. coryniformis [38], L. helveticus [39], L. johnsonii [40], L. plantarum [41], L. reuteri [9] and *L. sakei* [42]. These β -galactosidases of the LacLM type have been classified as members of glycoside family GH2 in the CAZy (Carbohydrate-Active Enzymes) databank (http://www.cazy.org). In contrast to this, β-galactosidase from *L. salivarius* was reported to be a monomer of ca. 30 kDa [43], the enzyme from L. delbrueckii subsp. bulgaricus a homodimer of 220 kDa [44], β-galactosidase from L. helveticus a homotetramer of 257 kDa [45], and L. plantarum forms - in addition to the above mentioned heterodimeric enzyme - a homohexameric β-galactosidase of ca. 325 kDa [46]. Some of these later β -galactosidases are members of glycoside family GH42. β-Galactosidase from L. pentosus KUB-ST10-1 is a heterodimer consisting of two subunits of approximately 35 and 72 kDa, which is in agreement with the majority of lactobacillal β-galactosidases reported hitherto, and is thus also of the LacLM type. The steady-state kinetic constants for β -galactosidase from *L. pentosus* were determined for the artificial chromogenic substrate oNPG and the natural substrate lactose. The k_{cat}/K_m value for oNPG is significantly higher than that of lactose,

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indicating *o*NPG as the better substrate. The K_m value of 37.8 mM determined for lactose compares positively with the Michaelis constants reported for commercially employed β-galactosidases, e.g., from *Aspergillus niger* (54–100 mM), *A. oryzae*(36–180 mM), *Kluyveromyces fragilis* (15–52 mM) [47] and *Kluyveromyces lactis* (20 mM) [48]. This value is, however, somewhat higher than the values reported for other *Lactobacillus* β-galactosidases: 13 mM for β-gal from *L. reuteri* L103 [9], 1.27 mM for β-gal from *L. kefiranofaciens* [49], and 4.04 mM for β-gal from *L. acidophilus* [34]. The higher K_{m,lactose} value for *L. pentosus* β-gal as compared to other lactobacillal enzymes, especially those of milk origin, could indicate a lesser adaptation of this soil-isolated strain to lactose utilization. *L. pentosus* β-gal is rather specific for the β-(1→4) galactosyl linkage. When using a range of structurally related nitrophenyl substrates, essentially no activity was measured for the β-(1→4) linked D-glucoside, D-mannoside, and D-xyloside substrates.

L. pentosus β-galactosidase is optimally active at pH 7.5 (glycine buffer) and 55 °C when using oNPG as substrate, and pH 8.0 (glycine buffer) and 60–65°C when lactose was used as substrate under otherwise standard activity assay conditions (10-min assay). This pH optimum is slightly more alkaline than reported for most other lactobacillal β -galactosidases, which show an optimum of 6.5–7.0 [9, 34]. however, we observed a pronounced effect of the buffer system used on the optimal pH, and that certainly affects the actual value determined. The temperature optimum of *L. pentosus* β-gal of 60–65 °C is considerably higher than reported for most other β-galactosidases from lactobacilli, which typically are optimally active in the range of 45–50 °C [9, 34, 43, 50]. L. pentosus KUB-ST10-1 can grow at temperatures of up to 45℃. This is higher than typical growth temperatures observed for many Lactobacillus spp. The increased temperature optimum and stability of the L. pentosus β-galactosidase might reflect an adaptation to these higher growth

temperatures. Because of this increased temperature optimum we investigated the thermostability of *L. pentosus* β-gal in more detail, determining the inactivation kinetics at different conditions. With respect to pH, highest half-life times $\tau_{1/2}$ were obtained in glycine buffer at pH 8.0-8.5 and at 37 °C (43-47 h). Again, the buffer system had a pronounced effect on stability, since the inactivation constants k_{in} determined were approximately twice as high at pH 7.5 when using sodium phosphate instead of glycine buffer; however the use of glycine buffer might be impractical for technical applications. When using the former buffer, highest stability was observed at pH 6.5–7.0 with $\tau_{1\!\!/_2}$ of 14.6 h. Addition of 2 mM Mg^{2+} increased stability and hence $\tau_{1/2}$ of β -gal activity significantly by a factor of up to 5.6-fold. For example, $\tau_{1/2}$ values of e.g. ~75 h were obtained at pH 6.5, 37 °C and when using phosphate buffer containing 2 mM of Mg^{2+} . This increase in stability and activity of L. pentosus β -gal could be a considerable advantage when using technical substrates such as cheese whey or whey permeates for the production of GOS, since cheese whey contains approximately 1.5 mM of Mg²⁺. Thus, use of these technical substrates could improve the performance of L. pentosus β -gal without the need of adding exogenous Mg²⁺.

β-Galactosidases have recently attained significant interest in biocatalytic processes towards the synthesis of oligosaccharides or various galactosylated compounds; for this purpose various galactosides can serve as galactosyl donor. Especially galacto-oligosaccharides (GOS), which are formed from lactose as a result of transgalactosylation, have attracted renewed attention in the field of functional food because of their proven benefits on health [16, 17, 18, 20]. As GOS are only transiently formed and are in turn substrates for the β -galactosidase as well, a maximum GOS yield of 31% (w/v) was reached at 78% lactose conversion when using the β -gal of *L. pentosus*. The main GOS spectrum formed from lactose by this

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enzyme is very similar to those found for β-gal from *L. reuteri* [13] and *L. acidophilus*[34]. *L. pentosus* β-gal shows a high specificity for the formation of β-(1→6) and β-(1→3) linkages, and hence the main transgalactosylation products detected were the disaccharides β-D-Gal*p*-(1→6)-D-Glc, β-D-Gal*p*-(1→3)-D-Glc, β-D-Gal*p*-(1→6)-D-Gal, β-D-Gal*p*-(1→3)-D-Gal, and the trisaccharides β-D-Gal*p*-(1→3)-D-Lac, β-D-Gal*p*-(1→6)-D-Lac. In a recent study a strong prebiotic effect was attributed to some of these compounds [51], and hence it is likely that the GOS mixture formed by *L. pentosus* β-gal has a considerable potential as prebiotic.

In conclusion, this work presented the characterisation and transgalactosylation capacity of a novel β-galactosidase from *L. pentosus*. Although the β-galactosidase described here is not derived from a proven probiotic *Lactobacillus* strain, the physical and biochemical characteristics are quite similar to those derived from probiotic strains [9, 34] with the exception that it is considerably more stable at higher temperatures. Also the GOS spectrum and yields are comparable with the former studies, making this novel enzyme that is produced by a well-known food-grade organism interesting for potential applications in food industry.

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<u>Tables</u>

Table 1. Kinetic properties of β -galactosidase from *Lactobacillus pentosus* for the hydrolysis of lactose and *o*-nitrophenyl- β -D-galactopyranoside (*o*NPG)

| substrate | method for | kinetic parameter | L. pentosus |
|--------------|----------------------|--|---|
| | determination of | | β-galactosidase |
| | enzyme activity | | |
| lactose | release of D-glucose | v _{max} (µmol min ⁻¹ mg ⁻¹) | 11.3 ± 0.75 |
| | | K _m (mM) | 37.8 ± 9.41 |
| | | k_{cat} (s ⁻¹) | 20.1 |
| | | $k_{cat}/K_{m} \ (mM^{-1} \ s^{-1})$ | 0.532 |
| <i>o</i> NPG | release of oNP | ν _{max} (μmol min ⁻¹ mg ⁻¹) | 304 ± 24.6 |
| | | K _m (mM) | 1.67 ± 0.64 |
| | | k _{cat} (s⁻¹) | 543 |
| | | $k_{cat}/K_{m} (mM^{-1} s^{-1})$ | 325 |
| | | | |
| | substrate | substrate method for determination of enzyme activity lactose release of D-glucose oNPG release of oNP | substrate method for kinetic parameter determination of enzyme activity lactose release of D-glucose V_{max} (µmol min ⁻¹ mg ⁻¹) K_m (mM) k_{cat} (s ⁻¹) K_m (mM) K_{cat} (s ⁻¹) K_m (mM) K_{cat} (s ⁻¹) K_{cat} / K_m (mM ⁻¹ s ⁻¹) |

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Table 2. Kinetic stability of β -galactosidase from *Lactobacillus pentosus* at various pH values at 37 °C.

| Buffer | рН | Inactivation constant k_{in} (h ⁻¹) | Half-life $r_{1/2}$ (h) | |
|------------------|-----|---|-------------------------|--|
| sodium acetate | 4.0 | 3770 × 10 ⁻³ | 0.18 | |
| sodium acetate | 5.0 | 504×10^{-3} | 1.37 | |
| sodium acetate | 5.5 | 131×10^{-3} | 5.31 | |
| sodium acetate | 6.0 | 91.9×10^{-3} | 7.54 | |
| sodium phosphate | 6.0 | 63.3×10^{-3} | 10.9 | |
| sodium phosphate | 6.5 | 47.6×10^{-3} | 14.6 | |
| sodium phosphate | 7.0 | 47.6×10^{-3} | 14.6 | |
| sodium phosphate | 7.5 | 95.8 × 10 ⁻³ | 7.24 | |
| glycine | 7.5 | 43.6×10^{-3} | 15.9 | |
| glycine | 8.0 | 14.7×10^{-3} | 47.2 | |
| glycine | 8.5 | 16.0×10^{-3} | 43.3 | |
| glycine | 9.0 | 1045 × 10⁻³ | 0.66 | |

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 Table 3. Kinetic stability of β -galactosidase from *Lactobacillus pentosus* at various temperatures with and without MgCl₂ supplementation (2 mM).

| Sodium phosph | | ate buffer (pH 6.5) Sodium phosp | | nate buffer (pH 6.5, + Mg ²⁺) | |
|---------------|--------------------------------------|----------------------------------|--------------------------------------|---|--|
| Temperature | Inactivation | | Inactivation | | |
| (°°) | constant k_{in} (h ⁻¹) | Half-life r _{1/2} (n) | constant k_{in} (h ⁻¹) | Haif-life $r_{1/2}$ (n) | |
| 4 | 0.5×10^{-3} | 1386 | nd | nd | |
| 23 | 0.8×10^{-3} | 866 | nd | nd | |
| 30 | 9.0×10^{-3} | 77.0 | $2.8 	imes 10^{-3}$ | 248 | |
| 37 | 46.6×10^{-3} | 14.9 | $9.4 	imes 10^{-3}$ | 73.7 | |
| 42 | 212.7×10^{-3} | 3.26 | 37.7×10^{-3} | 18.4 | |

nd...not determined

Figure captions

Figure 1. SDS-PAGE (A) and native PAGE (B) of purified β-galactosidase from *L. pentosus*. A: lane 1, molecular weight marker (Precision Plus Protein Dual Color, Biorad), CBB staining; lane 2, enzyme sample after treatment in SDS buffer at 100 °C for 5 min, CBB staining. B: lane 1, molecular weight marker (High Molecular Weight Calibration Kit for electrophoresis, Amersham); lane 2, CBB staining.

> **Figure 2.** pH optimum of β -galactosidase from *L. pentosus* using lactose (A) and *o*NPG (B) as the substrate. Symbols: •, sodium acetate buffer; O, sodium phosphate buffer; $\mathbf{\nabla}$, glycine buffer. Activities are given relative to standard assay conditions (sodium phosphate buffer, pH 6.5).

Figure 3. Temperature optimum of β -galactosidase from *L. pentosus* when using lactose (\bullet) and *o*NPG (O) as the substrate. Activities are given relative to standard assay conditions at 30 °C.

Figure 4. Inactivation kinetics of β-galactosidase from *L. pentosus* in sodium phosphate buffer (A; 50 mM, pH 6.5) and sodium phosphate buffer supplemented with 1 mM MgSO₄ (B; 50 mM, pH 6.5) at different temperatures.

Symbols: (A): ●, 4°C; O, 23°C; ▼, 30°C; ▽, 37°C; ■, 42°C; (B): ●, 30°C; O, 37°C; ▼, 42°C.

Figure 5. Formation of galacto-oligosaccharides (GOS) by β -galactosidase from *L. pentosus* during the discontinuous conversion reaction of 205 g/L lactose in 50 mM sodium phosphate buffer at 30 °C. (A) Formation of disaccharides (\bullet), trisaccharides

(O), tetrasaccharides ($\mathbf{\nabla}$) and total GOS (∇) during lactose conversion. (B) Time 590 course of reaction for lactose conversion; symbols: lactose ($\mathbf{\nabla}$), glucose ($\mathbf{\Phi}$), galactose (O).



210x297mm (600 x 600 DPI)

B

669 kDa

440 kDa

- 232 kDa

140 kDa

67 kDa



210x297mm (600 x 600 DPI)