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(54) Benævnelse: **KOLD-AKTIV BETA-GALACTOSIDASE, FREMGANGSMÅDE TIL FREMSTILLING HERAF OG ANVENDELSE AF ET SÅDANT ENZYM**

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DESCRIPTION

FIELD OF THE INVENTION

[0001] The present invention relates to a novel cold-active beta-galactosidase enzyme specific for lactose. The enzyme is thus useful in e.g. the food industry for catalyzing at low temperatures the hydrolysis of lactose disaccharide into its constituent monosaccharides, glucose and galactose.

BACKGROUND OF THE INVENTION

[0002] Beta-galactosidase (beta-D-galactoside galactohydrolase, EC 3.2.1.23) is an enzyme capable of hydrolyzing the disaccharide lactose to its monosaccharide constituents, D-glucose and D-galactose. Beta-galactosidases are found in a large variety of organisms, like mammals, plants, fungi, yeasts, and bacteria. In Nature, beta-galactosidases hydrolyze lactose and other D-galactose-containing carbohydrates. In the industry, beta-galactosidases have been used primarily within the food industry. Beta-galactosidase hydrolysis of lactose and lactose-containing dairy products are used throughout in the dairy industry in the preparation of lactose-free or low-lactose products, which may be consumed by humans suffering from lactose intolerance. Hydrolysis of lactose by beta-galactosidases may also be used in applications where the removal of lactose is required, i.e. prevention of crystallisation of lactose in food and removal of D-galactose moieties in glycosylated proteins. Other applications of beta-galactosidases comprise hydrolysis of lactose into D-galactose and D-glucose with the subsequent modification of the monosaccharides to high value products, like the sweetener D-tagatose (Jørgensen et al. 2004).

[0003] Application of beta-galactosidases could be used to produce lactose-free and low-lactose dairy products for lactose intolerant humans.

[0004] The major applications for lactose hydrolysis are listed below.

1. a) Liquid milk. Lactose hydrolysis in liquid milk improves digestibility for lactose intolerant consumers. In flavoured milks, lactose hydrolysis increases sweetness and enhances flavours.
2. b) Milk powders. Lactose hydrolysed milk powders for dietetic uses, especially for infants with temporary beta-galactosidase deficiency.
3. c) Fermented milk products. In some cases, lactose hydrolysis in milk used for the manufacture of cheese and yoghurt can increase the rate of acid development and thus reduce processing time.
4. d) Concentrated milk products. Lactose hydrolysis in concentrated milk products (e.g. sweetened condensed milk, ice cream) prevents crystallisation of lactose.
5. e) Whey for animal feed. Lactose hydrolysis in whey enables more whey solids to be fed to pigs and cattle and also prevents crystallisation in whey concentrate.
6. f) Whey. Lactose hydrolysed whey is concentrated to produce a syrup containing 70-75 per cent solids. This syrup provides a source of functional whey protein and sweet carbohydrate and is used as a food ingredient in ice cream, bakery and confectionery products.

[0005] The conventional approach in food processing is to carry out the hydrolysis of lactose at 40 °C during approximately four hours.

[0006] However, milk or lactose solution as a raw material is a preferable nutrition source for bacteria. As the result, the putrefaction owing to the saprophyte contamination during the treatment is a serious problem in the food production. Thus, the fact is that the conventional beta-galactosidase is of limited use.

[0007] Most beta-galactosidases in practical use are active only at temperatures above 20-30 °C, temperatures where food spoiling bacteria thrive at best.

[0008] Attempts to use thermophilic beta-galactosidases have been used but the products have suffered from off flavours and reduced organoleptic properties due to the heat treatment, and the processes have demanded high energetic costs.

[0009] A number of cold-active beta-galactosidases have been described from *Arthrobacter* (Coker, et al. 2003; Karasova-Lipovová, et al. 2003; Nakagawa et al. 2003; Nakagawa et al. 2006), from *Carnobacterium piscicola* (Coombs and Brenchley, 1999) and from *Pseudoalteromonas* (Cieslinski, et al. 2005; Fernandes, et al. 2002; Hoyoux, et al. 2001; Turkiewicz, et al. 2003). Furthermore, Nakagawa et al. (2006) described a cold-active beta-galactosidase from the yeast *Guehomyces pullulans*. However, the activity of cold-active beta-galactosidases described so far is low at the low temperatures, which is wanted by the dairy industry. The beta-galactosidase from the yeast *Guehomyces pullulans* had approximately 17% at 0 °C (Nakagawa et al. 2005), the beta-galactosidases from *Carnobacterium piscicola* BA showed approximately 24% activity at 10 °C (Coombs and Brenchley, 1999) and the enzymes from *Pseudoalteromonas* isolates showed 39% activity (Fernandes et al. 2002), 22% activity (Cieslinski et al. 2005), and 12% activity (Hoyoux et al. 2001) at 10 °C. So far, the beta-galactosidases with highest activity at low temperatures have been isolated from Antarctic *Arthrobacter* isolates. Karasová-Lipovová, et al. (2003) showed that a psychrotolerant *Arthrobacter* sp. C2-2 isolate produced beta-galactosidase, which displayed 19% of its maximal activity at 10 °C, Coker et al. (2003) described an enzyme from an Antarctic *Arthrobacter* isolate with approximately 50% at 0 °C, and Nakagawa et al. (2003, 2006) described a beta-galactosidase from A. *psychrolactophilus* F2, which had its temperature optimum at 10 °C.

[0010] However, the cold-active beta-galactosidase from the Antarctic *Arthrobacter* was produced in low amounts in native cells and attempts to produce the enzyme recombinantly in *E. coli* were unsuccessful since about 90% of the enzyme was located in insoluble inclusion bodies (Coker et al. 2003). The cold-active beta-galactosidase from A. *psychrolactophilus* F2 could be produced heterologously, but had lower activity than the other *Arthrobacter* beta-galactosidases (Nakagawa et al. 2006).

[0011] Therefore, in order to develop a low-temperature process for hydrolysis of lactose there is a need for a novel cold-active beta-galactosidase.

SUMMARY OF THE INVENTION

[0012] The present invention has, by using recombinant DNA technology, for the first time made it possible to provide a cold-active beta-galactosidase with high specific activity in industrially appropriate quantities for the manufacturing of food products and pharmaceuticals.

[0013] Accordingly, the present invention provides a purified cold-active beta-galactosidase, specific for lactose, having a stable enzymatic activity at temperatures less than 8°C, and specifically at 4°C, which corresponds to refrigerating conservation temperature for dairy products. The enzyme of the present invention is consequently able to hydrolyse lactose in dairy products and milk processing at such a low temperature that saprophytes are hindered to proliferate. The hydrolysis of lactose can be carried out in these refrigeration conditions with no need of a particular treatment to the dairy product concerned.

[0014] Specifically, the present invention provides a cold active beta-galactosidase having the sequence as defined in SEQ ID NO. 1, or one having at least 80% homology to the amino acid sequence as defined in SEQ ID NO. 1, the amino acid sequence being selected so that the enzyme has a stable enzymatic activity at temperatures less than 8 °C. Preferably the amino acid sequence has at least 90%, and more preferably 95%, homology to the amino acid sequence as defined in SEQ ID NO. 1.

[0015] In order to obtain the cold-active beta-galactosidase of the present invention there is further provided a DNA sequence, which

1. a) encodes a protein with an amino acid sequence as given in SEQ ID NO. 1, or
2. b) hybridises under stringent or very stringent conditions to the sequence of a), or
3. c) is degenerative of the sequence of a) or b)

[0016] Preferably the DNA sequence is derived from the genus *Alkalilactibacillus,* such as the species *Alkalilactibacillus ikkense,* and has a nucleotide sequence as given in SEQ ID NO. 2.

[0017] In a further embodiment, the present invention provides a recombinant vector comprising a DNA sequence that encodes a protein with an amino acid sequence as given in SEQ ID NO. 1, or one having at least 80% homology to the amino acid sequence as defined in SEQ ID NO. 1, the amino acid sequence being selected so that the enzyme has a stable enzymatic activity at temperatures less than 8 °C. Preferably the amino acid sequence has at least 90%, and more preferably 95%, homology to the amino acid sequence as defined in SEQ ID NO. 1.

[0018] Another object of the present invention is a strain of an isolated Alkalilactibacillus bacterium capable of producing a cold-active beta-galactosidase according to the present invention. A preferable strain is Alkalilactibacillus ikkense deposited on the 3rd of March, 2009, under the Budapest Treaty at the BCCM/LMG-Collection (Belgian Coordinated Collections of Microorganisms) with the Accession No LMG P-24866 and variants and mutants derived therefrom.

[0019] To purify the cold-active beta-galactosidase according to the present invention, a bacterium living in the Greenland area was isolated and characterised in order to study how its enzymes, and particularly, the beta-galactosidase was adapted to cold. These studies led to the purification of the beta-galactosidase, meaning that this protein was obtained substantially free of other proteins using protein purification steps known in the art.

[0020] Thus, another object of the present invention is a strain of an isolated Alkalilactibacillus bacterium capable of producing a cold-active beta-galactosidase according to the present invention. A preferable strain is alkalilactibacillus ikkense.

[0021] Another object of the invention is a recombinant plasmid or vector suited for transformation of a host, capable of directing the expression of a DNA sequence according to the invention in such a manner that the host expresses the cold-active beta-galactosidase of the present invention in recoverable form.

[0022] According to the invention, another object is the so transformed host. A variety of host-expression systems may be conceived to express the cold-active beta-galactosidase coding sequence, for example bacteria, yeast, insect cells, plant cells, mammalian cells, etc. Particularly, in yeast and in bacteria, a number of vectors containing constitutive or inducible promoters may be used.

[0023] An appropriate host organism is preferably selected from the group consisting of *Escherichia, Bacillus, Bifidobacterium, Lactococcus, Lactobacillus, Streptomyces, Leuconostoc, Streptomyces, Saccharomyces*, *Kluyveromyces*, *Candida*, *Torula, Torulopsis* and *Aspergillus.*

[0024] In a further aspect, the invention relates to a recombinant DNA molecule comprising a DNA fragment encoding a polypeptide having cold-active beta-galactosidase activity and to a microbial cell comprising such recombinant DNA molecule.

BRIEF DESCRIPTION OF THE FIGURES

[0025]

Figure 1 discloses a phylogenetic tree of 16S rRNA gene sequences from *Alkalilactibacillus ikkense* strain 517 and its closest relatives within the rRNA group 1 in the phyletic assemblage classically defined as the genus *Bacillus.* Bootstrap (n=100) values are shown. Bar, 0.015 substitutions per nucleotide position.

Figure 2 shows temperature dependence of native Ikka-beta-galactosidase. Y-axis is the relative activity in percent of the maximal activity. X-axis is the incubation temperature.

Figure 3 shows temperature stability of native Ikka-beta-galactosidase. Y-axis is the residual activity left after incubation at the time indicated on the X-axis. ◆, ●, ▲, ■, ◊, ○, indicate incubation temperatures at 0, 10, 20, 30, 40, and 50 °C, respectively.

Figure 4 discloses pH dependence of native Ikka-beta-galactosidase. Y-axis is the relative activity in percent of the maximal activity. X-axis is pH in the beta-galactosidase assay.

Figure 5 shows SDS-PAGE of extracts from native A. *ikkense* cells induced with 1 mM IPTG (lanes 2 and 3) and from uninduced, native A. *ikkense* (lanes 4 and 5). Arrows in lanes 2 and 3 indicate the 120 kDa beta-galactosidase band. Lane 1 is molecular weight marker. The marker at 120 kDa is beta-galactosidase from *E. coli.*

Figure 6 shows SDS-PAGE of three extract dilutions from recombinant *E. coli* cells expressing Ikka-beta-galactosidase (lanes 4, 5, and 6). For comparison, enzyme extract dilutions with beta-galactosidase from *Kluyveromyces lactis* (lanes 7, 8, and 9) and a molecular weight marker with native *E. coli* beta-galactosidase (marker at 120 kDa) were co-electrophoresed with the recombinant Ikka-beta-galactosidase. Arrows show the position of the 120 kDa beta-galactosidase bands

Figure 7 discloses temperature dependence of recombinant Ikka-beta-galactosidase produced in *E. coli.* Y-axis is the relative activity in percent of the maximal activity. X-axis is the incubation temperature. ◆, indicate recombinant Ikka-beta-galactosidase; ○, indicate *Kluyveromyces lactis* beta-galactosidase.

Figure 8 shows specific activity of recombinant Ikka-beta-galactosidase produced in *E. coli.* Y-axis is specific activity in nmoles ONP released/hour/mg enzyme. X-axis is temperature in °C. ◆ is recombinant Ikka-beta-galactosidase, ○ is *Kluyveromyces lactis* beta-galactosidase.

Figure 9 discloses thermal stability of recombinant Ikka-beta-galactosidase enzyme. An equal amount of enzyme was incubated at the temperature indicated and samples were withdrawn at different time intervals. Y-axis is absorbancy at wavelength 415 nm. X-axis is time in minutes.

Figure 10 also discloses thermal stability of recombinant Ikka-beta-galactosidase enzyme, while the X-axis is time in hours.

Figure 11 shows pH dependence of recombinant Ikka-beta-galactosidase. Y-axis is absorbancy measured at wevelength 415. X-axis is pH in the beta-galactosidase assay.

Figure 12 shows hydrolysis of lactose by recombinant Ikka-beta-galactosidase produced in *E.coli.* The reaction mixture contained lactose in three concentrations, 1.25 mg/ml, 2.5 mg/ml and 5 mg/ml. Samples were withdrawn after incubation for 15, 150, and 1440 minutes. The reactions were incubated at 5 °C (A) or 20 °C (B) and analysed by thin layer chromatography (TLC). The TCL plates were sprayed with an orcinol reagent, and hydrolysis of lactose was estimated by the disappearance of the lactose spots on TLC plates and the concomitant appearance of glucose and galactose spots.

Figure 13 shows SDS polyacrylamide gel electrophoresis of crude extract of *E. coli* cells expressing A. *ikkense* beta-galactosidase, Lane 1. The beta-galactosidase in the crude extract was further purified on ion exchange chromatography, Lane 2, or on affinity chromatography, Lane 3. The arrow indicates the position of the *E. coli* 120 kDa beta-galactosidase band. Numbers to the left indicate the position of protein bands in the PageRuler™ Plus Prestained Protein Ladder (Fermentas).

Figure 14 shows temperature dependence (A) and pH dependence (B) of the purified, recombinant *A. ikkense* beta-galactosidase and of a crude extract containing recombinant enzyme. Y-axis is the relative activity in percent of maximal activity. X-axis is the incubation temperature (A) or pH (B). The relative activity of the purified, recombinant enzyme with ONPG as substrate is illustrated by black squares. Assays were performed in triplicates, and standard error was below 0.05.

Figure 15 shows thermal stability of the purified, recombinant *A. ikkense* beta-galactosidase enzyme. An equal amount of enzyme was incubated at the temperatures indicated and samples were withdrawn at different time intervals. Y-axis is the residual activity in percent of maximal activity. Enzyme samples were incubated at temperatures from 0 °C to 60 °C and at the time points indicated (X-axis), samples were withdrawn and assayed for active beta-galactosidase at 20 °C. Assays were performed in triplicates, and standard error was below 0.05.

Figure 16 shows benchmarking the A. *ikkense* beta-galactosidase (black squares) with commercially available Lactozyme® 3 3000L from *K. lactis* (open circles). An equal amount of enzyme (2 mg/ml) was incubated with ONPG as substrate at temperatures from 0 °C to 20 °C. Samples were withdrawn at different time intervals and hydrolyzed ONP was measured at A420 nm. Hydrolysis efficiency was calculated as increase in A420 nm per min per microgram active enzyme. Assays were performed in triplicates, and standard error was below 0.05.

Figure 17 shows thin-layer chromatography (TLC) of the hydrolysis of lactose by the A. *ikkense* beta-galactosidase.

Lanes 4-6: samples incubated at 5 °C for 2½ h (4: 1.25 mg/ml lactose, 5: 2.5 mg/ml lactose, 6: 5 mg/ml lactose).

Lanes 7-9: samples incubated at 20 °C for 2½ h (7: 1.25 mg/ml lactose, 8: 2.5 mg/ml lactose, 9: 5 mg/ml lactose).

Lanes 1-3: Controls, 0.0125 µg of each of the carbohydrates 13 lactose (lane 1), galactose (lane 2) and glucose (lane 3).

DETAILED DESCRIPTION OF THE INVENTION

[0026] "Beta-galactosidase" (beta-D-galactoside galactohydrolase, EC 3.2.1.23) is defined as an enzyme capable of hydrolysing lactose to the monosaccharides D-glucose and D-galactose.

[0027] "Cold-active" is defined as having activity at temperatures at 15 °C and below, preferably at 10 °C and below and most preferably at 5 °C and below.

[0028] A "host cell" is selected from a group of microorganisms comprising fungi, yeasts, and prokaryotes. The microorganism is more preferably a prokaryote and most preferably a bacterium.

[0029] Conditions of incubating beta-galactosidase with lactose are defined by performing incubation at a temperature between 0 °C and 20 °C, preferably between 5 °C and 15 °C.

[0030] The term "stringent condition" refers to a condition such that a hybridization is conducted in a solution containing 6xSSC (20xSSC represents 333 mM Sodium citrate, 333 mM NaCl), 0.5% SDS and 50% formamide at 42 °C, and then the hybridized products are washed in a solution of 0.1xSSC, 0.5% SDS at 68°C, or to conditions as described in Nakayama, et al., Bio-Jikken-Illustrated, vol. 2, "Idenshi-Kaiseki-No-Kiso (A Basis for Gene Analysis)", pp. 148-151, Shujunsha, 1995.

EXAMPLES

EXAMPLE 1

Isolation of bacteria producing cold-active beta-galactosidases

1.1 Sampling of bacteria

[0031] Ikaite material was collected by scuba divers in the Ikka Fjord, South West Greenland (61°11'N, 48°01'W) from a depth of approximately six to ten meters. The columns were between 36-70 cm in length and between 5 and 30 cm in diameter. The columns were kept cold during transportation to the field laboratory.

1.2 Screening bacteria for beta-galactosidase production

[0032] Approximately 3 cm^3 of ikaite material from a slice 15-18 cm from the top of an ikaite column was drilled out and suspended in 250 ml R2 broth (Schmidt et al. 2006) buffered to pH 10 with 0.2 M Na_2CO_3/$NaHCO_3$ buffers as described by Stoll and Blanchard (1990). After incubation at 5 °C for 2 months the culture was inoculated onto R2 medium, pH 10 without glucose but supplemented with lactose (1% w/v), 5-bromo-4-chloro-3-indolyl-β-D-galactopyranoside (X-gal) (40 µg/ml), 10 mM isopropyl-beta-D-thiogalactopyranoside (IPTG) and agar (1.5 %, w/v). The plates were incubated at 5 °C for one to two weeks. A total of 17 blue colonies indicating production of beta-galactosidase were detected. Since 16S rRNA gene analyses of the seventeen isolates showed identical sequences only one of the isolates, strain 517, was chosen for further characterization.

EXAMPLE 2

Taxonomical analysis of Isolate 517 and description of a new genus and species, *Alkalilactibacillus ikkense*

2.1 Phylogenetic analysis of 16S rRNA gene sequences

[0033] DNA for phylogenetic analysis was extracted from cells of isolate 517 using FastDNA SPIN Kit for Soil as described by the manufacturer (BIO 101, Irvine, CA). 16S rRNA gene amplification was carried out using the primers 27F and 1492R (Lane 1991), and DNA sequencing was carried out at MWG Biotech AG (Ebersberg, Germany) using the same two primers plus additional primers 519R, 532F, 907F, and 907R (Lane 1991). The near full length DNA sequence of the 16S rRNA gene from isolate 517 was submitted to GenBank/EMBL/DDBJ with the accession number EU281853. Related sequences were retrieved from public databases using BLASTn at the NCBI server (http://www.ncbi.nlm.nih.gov/blast/). The closest related 16S rRNA gene sequences were aligned using the Clustal W multiple alignment program MegAlign 5.03 (DNASTAR, Inc., Madison, WI). The Clustal W analysis showed that the closest relatives were *Natronobacillus azotifigens* (accession no. EU850815) (Sorokin et al. 2008), *Paraliobacillus ryukyuensis* (accession no. AB087828) (Ishikawa et al. 2002), *Halolactibacillus halophilus* (accession no.

AB196783) (Ishikawa et al. 2005), *Halolactibacillus miurensis* (accession no. AB196784) (Ishikawa et al. 2005), *Amphibacillus tropicus* (accession no. AF418602) (Zhilina et al. 2001, 2002), and *Gracilibacillus halotolerans* (accession no. AF036922) (Wainø et al. 1999). Isolate 517 was most closely related to N. *azotifigens, P. ryukyuensis* and A. *tropicus* with 95,9%, 94.4% and 93.9% sequence similarity, respectively. The sequence similarity between isolate 517 and both H. *halophilus, H. miurensis,* and G. *halotolerans* was 93.4%. Thus, the distance in 16S rRNA gene sequence similarity between isolate 517 and the closest related is below the 97% similarity, which is often used as a preliminary guideline for species separation. A phylogenetic tree was created by neighbour-joining analysis (bootstrap=100) using TREECON 1.3b software (Van de Peer and De Wachter, R. 1994) Fig. 1.

2.2 DNA-DNA hybridization and base composition analysis of genomic DNA

[0034] DNA-DNA hybridization and DNA base composition (G+C content) was carried out at DSMZ (Braunschweig, Germany). DNA was isolated using a French pressure cell (Thermo Spectronic) and was purified by chromatography on hydroxyapatite as described by Cashion et al. (1977). DNA-DNA hybridization was carried out as described by De Ley et al. (1970) under consideration of the modifications described by Huss et al. (1983) using a model Cary 100 Bio UV/VIS-spectrophotometer equipped with a Peltier-thermostatted 6x6 multicell changer and a temperature controller with *in-situ* temperature probe (Varian). DNA-DNA hybridization between isolate 517 and the closest related based on 16S rRNA sequence similarity *P. ryukyuensis* was 28.8%, and between isolate 517 and *H. miurensis* it was 24.7%.

[0035] For determination of GC content, the DNA was hydrolyzed with P1 nuclease and the nucleotides dephosphorylated with bovine alkaline phosphatase (Mesbah et al. 1989) . The resulting deoxyribonucleotides were analyzed by HPLC. The DNA G+C content of isolate 517 was 38.4 mol%, which is fairly similar to the closest related species. The G+C content of *N. azotifigens* is 36.1-38.5 mol% (Sorokin et al. 2008), *H. halophilus* and *H. miurensis* is reported to be 38.5-40.7 mol% (Ishikawa et al. 2005), for *P. ryukyuensis* it is 35.6 mol% (Ishikawa et al. 2002), and for *G. halotolerans* it is reported to be 38 mol% (Wainø et al. 1999).

[0036] The phylogenetic results and data on GC content indicate that isolate 517 represent a new species within a new genus, since the threshold value for DNA-DNA hybridization to separate two species is 70 % (Wayne et al., 1987). Thus, we propose that isolate 517 represents a new genus *Alkalilactibacillus* gen. nov. comprising the species *Alkalilactibacillus ikkense* sp. nov.

EXAMPLE 3

Characterization of native Ikka-beta-galactosidase from *Alkalilactobacillus ikkense*

3.1 Beta-galactosidase assay.

[0037] Beta-galactosidase activity was assayed by hydrolysis of *o*-nitrophenyl-beta-D-galactopyranoside (ONPG) and measuring the absorbancy of the released o-nitrophenyl (ONP) compound in a spectrophotometer at 415 nm. In the assay, the release of ONP from 1 mM ONPG by the recombinant beta-galactosidase activity was measured at 415 nm at 20 °C and pH 7.0 (0.1 M NaH_2PO_4/Na_2HPO_4). The reactions were stopped by adding 300 µl 0.6 M Na_2CO_3. Assays were performed at 0, 5, 10, 20, 30, 40, 50, and 60 °C for 30 minutes. The sodium phosphate buffers were pre-heated to the respective temperatures prior to assay start.

[0038] Thermostability analysis of the enzyme was carried out by placing an aliquot of enzyme at temperatures 0, 10, 20, 30, 40, and 50 °C and taking samples at t = 0 to t = 24 hours. Immediately after taking the samples, they were cooled and assayed at 20 °C as described above.

[0039] The pH activity profile was studied using a mixed pH buffer (250 mM Tris, 250 mM MES, 250 mM acetic acid) adjusted from pH 4 to pH 10 with HCl or NaOH. The samples were incubated at 20 °C for 1 hour and assayed as described above.

3.2 Production of native Ikka-beta-galactosidase.

[0040] *Alkalilactibacillus ikkense* cells were cultivated in liquid R2 medium supplemented with lactose and IPTG at 15 °C for 3

days on a rotary shaker. Cells were harvested by centrifugation in a Sigma® 3-18M centrifuge at 4,700 rpm and the pellet was resuspended in 2 ml of 0.1 M NaH_2PO_4/Na_2HPO_4, pH 7. Cells were lysed by bead beating in a FastPrep FP120 instrument (Bio101/Savant) for 3 times 25 sec at speed 5.5. The supernatant was then removed from the glass beads and centrifuged for 15 min at 10,000*g at 4 °C. The cell free supernatant was then used for assaying.

3.3 Characterization of temperature optimum of native Ikka-beta-galactosidase.

[0041] The native Ikka-beta-galactosidase displayed maximal activity at 20 °C, 40% of the maximal activity was obtained at 0 °C, and more than 60% of the maximal activity was observed at 10 °C (Fig. 2). Above 30 °C the enzyme was only moderately active and virtually no activity was observed at 60 °C.

[0042] The temperature stability of the Ikka-beta-galactosidase was investigated. Fig. 3 shows that almost 100% of the residual activity was observed after 24 hours incubation at 0 °C and that more than 80% activity was left after 24 hours at 20 °C. At temperatures above 20 °C, the native Ikka-beta-galactosidase rapidly lost activity (Fig. 3). The inactivation at high temperatures was shown to be irreversible.

[0043] The pH dependence of the native Ikka-beta-galactosidase was investigated. Fig. 4 shows that maximal activity of the native Ikka-beta-galactosidase was observed at pH 7, and that the enzyme displayed approximately 70% of the maximal activity at pH 6 and at pH 8. At pH 9, approximately 25% of maximal activity was observed. The enzyme showed no activity at pH 5 and below or at pH 10 and above (Fig. 4).

3.4 SDS-PAGE of native Ikka-beta-galactosidase.

[0044] Extracts from cells of A. *ikkense* induced with 1 mM IPTG and uninduced were analysed in SDS-PAGE (sodium dodecyl sulfate - polyacrylamide gel electrophoresis) (Fig. 5). Intracellular extracts were prepared by lysing the cells using bead-beating as described above in 3.2. The extracts (0.5-5 µl) were mixed with 12.5 µl 4*LDS sample buffer, 5 µl 10*DTT and 0.1 M NaH_2PO_4/Na_2HPO_4 to a final volume of 50 µl. The samples were heated to 70 °C for 10 minutes and 30 µl was loaded onto a 4-12% SDS gel. The gel was run in a XCell *SureLock*™ Mini-Cell (Invitrogen, CA, USA) at 150 V for 1 hour at room temperature. After electrophoresis, the gel was stained using Coomassie Brilliant Blue R-250 (0.1% Coomassie Brilliant Blue R-250 (Serva, Heidelberg, Germany) in 40% EtOH and 10% acetic acid). Fig. 5 shows that a strong 120 kDa band was observed in the lanes with extracts from A. *ikkense* cells induced with IPTG and that this band was missing in lanes with extracts from non-induced cells. Thus, the 120 kDa band was assumed to be the native Ikka-beta-galactosidase.

EXAMPLE 4

Isolation and characterization of the Ikka-beta-galactosidase gene from *Alkalilactobacillus ikkense*

4.1 Isolation of the Ikka-beta-galactosidase gene

[0045] DNA from *A. ikkense* was isolated from a culture of 50 ml. The cells were harvested by centrifugation and the chromosomal DNA was isolated using conventional phenolchloroform extraction methods (Maniatis et al., 1982). The DNA was partially digested using *Sau3AI* (New England Biolabs, MA, USA), and fragments with the lengths between 3 kb and 10 kb were purified from an agarose gel using the QIAquick Gel Extraction Kit (Qiagen, Hilden, Germany) as described by the producer.

[0046] The vector for cloning chromosomal DNA from A. *ikkense* was a modified pUC18 plasmid (Stratagene, CA, USA). Plasmid pUC18 was restricted by NdeI and *Hind*III endonucleases (New England Biolabs). The sticky ends were refilled using Klenow fragment of DNA polymerase (New England Biolabs) and the blunt ends were ligated using T4 DNA ligase (New England Biolabs). DNA sequencing of the modified pUC18 plasmids, denoted pUC18dlacZ, at GATC Biotech AG (Konstanz, Germany) and analysis of the DNA sequence in the CLC Workbench 4 software (CLC bio, Aarhus, Denmark) confirmed that the α-subunit sequence of pUC18 was deleted in plasmid pUC18dlacZ. Thus, plasmid pUC18dlacZ was not able to mediate α-complementation when introduced into *E. coli* cells harbouring the beta-galactosidase ΔZ15 mutation. *Sau3A*I restricted and gel-purified chromosomal

DNA from A. *ikkense* was ligated into plasmid pUC18dlacZ treated with the restriction endonuclease *BamH*I and Antarctic Phosphatase (New England Biolabs). The ligation mixture was transformed into chemically competent *E. coli* TOP10 cells. Transformed cells were plated onto LB agar (10 g/l Bacto tryptone, 5 g/l yeast extract, 10 g/l NaCl) containing 20 µg/ml X-gal, 0.1 mM IPTG, and 100 µg/ml ampicillin and incubated over night at 37 °C. After a 16 hours over night incubation, the plates were transferred to 20 °C and incubated for another 20 hours. A total of 580 colonies were screened and 1 blue colony was detected. The colony that turned blue during incubation at 20 °C was selected and transferred to 10 ml LB broth and grown at 37 °C over night. Recombinant *E. coli* cells from over night cultures were harvested by centrifugation, and plasmid DNA was purified using the QIAprep Spin Miniprep Kit (Qiagen). Plasmid DNA was analysed for inserts by digestion with restriction endonucleases *EcoRI* and PstI (New England Biolabs). The insert in the plasmid, denoted pUClkka-bgal, was sequenced at GATC Biotech AG (Konstanz, Germany) using primer walking with the primer M13 reverse and custom made primers specific for the insert in pUClkka-bgal (SEQ ID 3: 5'CCGTCATCCATATCACC3'; SEQ ID 4: 5'CCTTTGCCCAAGAGCCAACC3'; SEQ ID 5: 5'GCTATTATCAGACTTGGCACC3'; SEQ ID 6: 5'GTAATTCAAT GTTCCAACGTG3'; Seq ID 7: 5'CGCTTATGGTGTGAAG3') and a sequence just downstream of the multiple cloning site in pUC18dlacZ, (SEQ ID 8: 5'GGGCTGGCTTAACTATGCGG3'). The lkka-beta-galactosidase gene sequence harboured by the DNA insert is shown as SEQ ID NO 2.

4.2 Characterization of the lkka-beta-galactosidase gene sequence

[0047] Analysis of the DNA sequence, SEQ ID NO 2, using the CLC Workbench 4 software (CLC bio, Aarhus, Denmark) showed an open reading frame with the coding capacity of 1,041 amino acids, SEQ ID NO 1. The NCBI search tool Blastp was used to search for related sequences in databases. The closest related sequences were beta-galactosidases from *Bacillus megaterium* (accession no ABN13675) 56.7% identity, *Paenibacillus* sp. JDR-2 (accession no ZP_02849115) 55.3% identity, and *Geobacillus* sp. Y412MC10 (accession no ZP_03036811) 54% identity, all of which belong to the Glycosyl Hydrolase Family 2. Thus, it is concluded that the lkka-beta-galactosidase belongs to this family. The calculated subunit molecular weight and pI of the lkka-beta-galactosidase was 119 kDa and pI 5.0, respectively (ExPASy ProtParam tool). The calculated subunit molecular weight was confirmed by SDS-PAGE, Figs. 5 and 6. Alignment of the lkka-beta-galactosidase with structurally resolved enzymes showed that the conserved active site region in *E. coli* (ILCEYAHAMGN) (pos. 534-544) (Gebler *et al.* 1992) is well conserved in the lkka-beta-galactosidase (ILCEFSHAMGN) (pos. 547-557), and the active site nucleophile Glu-537 is probably found as Glu-550.

EXAMPLE 5

Production of recombinant lkka-beta-galactosidase in *Escherichia coli.*

[0048] Native *Alkalilactibacillus ikkense* was shown to produce only moderate amounts of lkka-beta-galactosidase. Therefore, in order to produce larger amounts of the beta-galactosidase, subcloning of the lkka-beta-galactosidase gene into expression plasmids was carried out.

5.1 Construction of a vector for the expression of recombinant lkka-beta-galactosidase in *Escherichia coli.*

[0049] The lkka-beta-galactosidase gene was subcloned further using chromosomal DNA from *A. ikkense* as template and the PCR primers bGal5': 5'CTGAATTCGCATATGGCAAAAAAATTAAAAAAATTC3' (*Eco*RI restriction site underlined) (SEQ ID 9), and bGal3': 5'CCAAGCTTATCTGTTTAAACTATTCAACATG3' (*Hind*III site double underlined) (SEQ ID 10). The polymerase used was the proofreading polymerase Phusion® High-Fidelity DNA Polymerase (New England BioLabs). The PCR reaction was analyzed by gel electrophoresis on a 0.8% agarose gel (Seakem GTG) and the 3.9 kb fragment was ligated into pJET1.2/blunt cloning vector (Fermentas, Helsingborg, Sweden) and transformed into *E. coli* TOP10 cells. *E. coli* transformants containing pJET1.2/blunt were isolated on ampicillin containing LB agar plates and plasmid DNA was prepared as described above. Plasmid DNA was restricted with the enzymes *EcoRI* and *Hind*III and analysed on 0.8% (w/v) agarose gels as described. The 3.9 kb DNA fragment was purified from the gel using the QIAquick Gel Extraction Kit as described by the producer. The purified DNA fragment was ligated into plasmid pUC18dlacZ similar restricted with the enzymes *EcoRI* and *Hind*III and gel purified as described above. The ligation mixture was transformed into *E. coli* TOP10 cells and recombinant cells harbouring the plasmid pUC18dlacZ with the lkka-beta-galactosidase gene were selected as blue colonies on LA plates containing 100 µg/ml ampicillin, 1 mM IPTG, and 40 µg/ml X-gal. Transformants were selected and analysed for plasmids and inserts. Plasmid DNA was prepared from a 10 ml culture and the DNA was sent for sequencing at GATC biotech (Konstanz, Germany) using the primers described above in 4.1. The

entire Ikka-beta-galactosidase gene was sequenced on both strands in order to ensure that no mutations were introduced during PCR. One of the recombinant clones, which contained plasmid pUC18dlacZ with the Ikka-beta-galactosidase gene, denoted plasmid pUCIkka-bgal_exp, was selected for further expression studies.

5.2 Expression of recombinant Ikka-beta-galactosidase in *Escherichia coli.*

[0050] *E. coli* TOP10 cells harbouring plasmid pUCIkka-bgal and pUCIkka-bgal_exp were cultivated in 30 ml LB broth containing 100 µg/ml ampicillin over night at 37 °C. After an over night incubation, the cells were supplemented with 0.1 mM IPTG and incubated at 20 °C for further 20 hours. Cells were harvested by centrifugation for 30 min at 4,700 rpm at 10 °C and resuspended in 1 ml 0.1 M NaH_2PO_4/Na_2HPO_4 pH 7. The cells were lysed by bead beating in a Fast Prep instrument (Fast Prep FP120, Bio101/Savant Instruments Inc., Holbrook, NY) at speed 5.5 for 3 times 25 sek. The samples were cooled on ice in between the beating/shaking. The lysate was centrifuged at 10,000 g for 15 min at 4 °C, and the supernatant containing the Ikka-beta-galactosidase enzyme was transferred to a clean tube. This crude extract was used for subsequent analyses.

5.3 Properties of recombinant Ikka-beta-galactosidase produced in *Escherichia coli.*

5.3.1. SDS-PAGE of recombinant Ikka-beta-galactosidase and determination of yield produced.

[0051] Intracellular extracts from recombinant *E. coli* cells harbouring plasmid pUCIkka-bgal were analysed on SDS-PAGE (SDS gel 4-12%, PAGEgel, CA, USA) as described above in 3.4. Cultures induced with 1 mM IPTG and control cultures not induced were analysed.

[0052] The protein bands in extracts from cultures grown with and without IPTG were identical apart from a band of approximately 120 kDa in cultures induced with IPTG (arrow in Fig. 6). Thus, as the calculated molecular mass of the Ikka-beta-galactosidase is 119 kDa, and since the strong band at 120 kDa was observed only in cultures induced with IPTG, it is assumed that the 120 kDa band represent the Ikka-beta-galactosidase.

[0053] Extracts from cultures of plasmid pUCIkka-bgal harbouring *E. coli* were prepared as described above and diluted before electrophoresis on SDS-PAGE. Beta-galactosidases,from *E. coli* (Sigma-Aldrich, MO, USA) and *K. lactis* (Novozymes, Bagsvaerd, Denmark) with known molecular mass and in defined concentrations were co-electrophorezed on the same gel for comparison (arrows in Fig. 6). By comparing the migration and Coomassie Brilliant Blue staining of the known beta-galactosidases with that of the Ikka-beta-galactosidase an estimate of the amount of the Ikka-beta-galactosidase was obtained. The extract, which was used for the subsequent analyses was estimated to have a concentration of Ikka-beta-galactosidase of 2 mg/ml.

5.3.2 Temperature dependence of recombinant Ikka-beta-galactosidase.

[0054] The temperature optimum of recombinant Ikka-beta-galactosidase was determined as described above for the native enzyme using ONPG as a substrate. The temperature profile was determined at 0, 5, 10, 20, 30, 40, 50, and 60 °C for 30, 60 and 120 minutes for the recombinant Ikka-beta-galactosidase and, as controls, for the beta-galactosidases from *E. coli* and K. *lactis*. The optimal temperature for activity of the recombinant Ikka-beta-galactosidase was determined to be 20-30 °C (Fig. 7). However, the Ikka-beta-galactosidase also showed high activity at low temperatures with more than 40% activity at 0 °C, approximately 80% activity at 5 °C and more than 90% at 10 °C. Compared to the *K. lactis* beta-galactosidase, the specific activity of the Ikka-beta-galactosidase was almost twice as high at temperatures between 0 °C and 30 °C (Fig. 8). Both enzymes showed close to zero activity at 40 °C and above (Fig. 8).

[0055] The thermostability of the Ikka-beta-galactosidase was determined at 0, 10, 20, 30, 40, 50, and 60 °C. Samples were taken from t = 0 hours to t = 123 hours with increasing intervals from 5 minutes during the first hour to several hours at the end of the experiment (Fig. 9). Figs. 9 and 10 show that the Ikka-beta-galactosidase showed high stability at 0 °C and 5 °C, at 10 °C the enzyme was stable for approximately 100 hours and at temperatures above 20 °C the Ikka-beta-galactosidase was rather unstable. Treatment at 40 °C for 40 minutes resulted in complete inactivation. The inactivation of the Ikka-enzyme was irreversible.

5.3.3 pH dependence of recombinant Ikka-beta-galactosidase.

[0056] The pH activity profile was studied using a mixed pH buffer (250 mM Tris, 250 mM MES, 250 mM acetic acid) adjusted from pH 4 to pH 10 with HCl or NaOH. The samples were incubated at 20 °C for 2 hours. The optimal pH value for the Ikka-beta-galactosidase was shown to be approximately pH 7.0. At pH 6.0 the enzyme showed 60% of maximal activity and at pH 8.0 the Ikka-enzyme displayed 90% activity. At pH 9.0 15% activity was observed whereas no activity could be detected at pH 5.0 or below or at pH 10 and above.

5.3.3 pH dependence of recombinant Ikka-beta-galactosidase.

[0057] The substrate specificity of the Ikka-beta-galactosidase was determined in assays performed at pH 7.0 and 20 °C for 20 minutes using nine different chromogenic substrates, *o*-nitrophenyl-beta-D-galactopyranoside, p-nitrophenyl-alpha-D-galactopyranoside, *o*-nitrophenyl-beta-D-glucopyranoside, p-nitrophenyl-beta-D-glucopyranoside, p-nitrophenyl-beta-D-arabinopyranoside, p-nitrophenyl-beta-D-cellobioside, p-nitrophenyl-beta-D-fucopyranoside, p-nitrophenyl-beta-D-lactopyranoside, and p-nitrophenyl-beta-D-mannopyranoside. Each substrate was used at a concentration of 10 mM. The assays showed that the Ikka-beta-galactosidase was only able to hydrolyze o-nitrophenyl-beta-D-galactopyranoside (ONPG) and p-nitrophenyl-beta-D-fucopyranoside (4 % of the relative activity compared to hydrolysis of ONPG). The utilization of the remaining substrates was below detection.

[0058] Hydrolysis of lactose was determined in a solution of lactose in water. Three different lactose concentrations were tested: 1.25 mg/ml, 2.5 mg/ml, and 5 mg/ml. Total reaction volume was 0.2 ml and each reaction contained 0.2 mg/ml of recombinant Ikka-beta-galactosidase enzyme. The enzyme reactions were incubated at 5 °C and 20 °C, and samples were collected after 15 minutes, 2½ hour, and 24 hours. After incubation, the reactions were stopped by heating at 95 °C for 20 minutes. Visualization of the products was carried out by thin-layer chromatography (TLC) on a TLC Silica gel 60 (Merck, Darmstadt, Germany) in a solvent containing 1-butanol, 2-propanol, and water (3:12:4). Volumes containing 0.005 mg lactose were run on the TLC. Controls were 0.5 µl lactose (2.5 %), 0.5 µl galactose (2.5 %) and 0.5 µl glucose (2.5 %). After being dried, the sugars were visualized by spraying with an orcinol reagent followed by incubation for 5-10 min at 100 °C.

[0059] Hydrolysis of lactose was observed both at 5 °C and at 20 °C. At 5 °C, approximately 75-85% lactose was hydrolysed in the 1.25 mg/ml reaction after 15 minutes and 100% was hydrolysed within 2½ hours, (Fig. 12A). Similar hydrolysis efficiency was observed in the 1.25 mg/ml reaction incubated at 20 °C (Fig. 12B). Hydrolysis effectiveness in the 2.5 mg/ml lactose reaction showed approximately 90-95% hydrolysis within 2½ hours at both temperatures. After 24 hours, 100% hydrolysis was observed for all three lactose concentrations at both temperatures (Fig. 12).

5.3.4 Purification of recombinant Ikka-beta-galactosidase.

[0060] Beta-galactosidase was purified from crude extracts by ion exchange chromatography. Portions of two ml were subjected to chromatography on a 1 ml High Q cartridge on a BioLogic LP System (Bio-Rad).

[0061] The column was washed with 10 ml of 50 mM phosphate buffer (pH 7) and eluted by a gradient from 0 to 1 M of NaCl in 50 mM phosphate buffer (pH 7) at a flow rate of 0.5 ml/min. Fractions of 1 ml were collected. Crude extracts were also subjected to affinity chromatography on a 2 ml column of agarose coupled with p-aminobenzyl-1-thio-beta-D-galactopyranoside (PABTG-agarose, Sigma). The column was washed with 10 ml of 50 mM phosphate buffer (pH 7) before it was eluted by 100 mM NaCl in 50 mM phosphate buffer (pH 7) at a flow rate of 0.5 ml/min. Fractions of 1 ml were collected.

[0062] The fractions were analysed for the presence of proteins using a BCA Protein Assay Kit (Pierce), and beta-galactosidase was measured in o-nitrophenyl (ONP)-beta-D-galactopyranoside assays as described above. Fractions containing beta-galactosidase activity were analysed by SDS polyacrylamide gel electrophoresis (4-20 %, PAGEgel, CA, USA). Purified beta-galactosidase was used for subsequentstability and activity experiments.

[0063] Fig. 13 shows that the crude extract from *E. coli* contained recombinant beta-galactosidase with a monomeric molecular weight of approximately 115-120 kDa. Ion exchange chromatography resulted in pure beta-galactosidase (Fig. 13, lane 2), whereas affinity chromatography (Fig. 13, lane 3) only resulted in partially purified recombinant enzyme. Thus, for the subsequent

analyses, pure beta-galactosidase from ion exchange was used, unless otherwise specified.

5.3.5 Characterization of native and recombinant A. *ikkense* beta-galactosidase.

[0064] The molecular weight of the A. *ikkense* beta-galactosidase was determined to be approximately 115-120 kDa when analyzed on SDS-PAGE using known beta-galactosidases from *E. coli* and *K. lactis* as references. This result is in agreement with the calculated molecular weight as determined from the DNA sequence (119 kDa). The crude extract from *E. coli* was estimated to contain 10 mg/ml A. *ikkense* beta-galactosidase.

[0065] The specific activity, calculated on the basis of purified beta-galactosidase from ion exchange chromatography was 8.4 micromoles/min/mg protein at 20 °C with ONPG as substrate (Table 1).

Table 1

Purification	Volume (ml)	Protein (mg)	Specific activity (U mg^{-1})	Total activity (U)	Purification (fold)	Recovery (%)
Cell extract	10	30	1.6	48.0	1	100
Ion exchange	10	2.5	8.4	21.1	12	44
A culture of 200 ml *E. coli* cells expressing recombinant A. *ikkense* beta-galactosidase was harvested resulting in a 280 mg wet weight cell pellet. The cells were lyzed in a Fast Prep apparatus, and the extract was subjected to ion exchange chromatography. One U is 1 micromole per min at 20 °C with ONPG as substrate.b						

[0066] Extract of *E. coli* expressing recombinant A. *ikkense* beta-galactosidase was analyzed for beta-galactosidase activity. The thermal stability of the purified, recombinant A. *ikkense* beta-galactosidase enzyme was tested as shown in Figure 15. An equal amount of enzyme was incubated at the temperatures indicated and samples were withdrawn at different time intervals. Y-axis is the residual activity in percent of maximal activity. Enzyme samples were incubated at temperatures from 0 °C to 60 °C and at the time points indicated (X-axis), samples were withdrawn and assayed for active beta-galactosidase at 20 °C. Assays were performed in triplicates, and standard error was below 0.05.

[0067] At 0 °C the enzyme displayed more than 60 % of the maximal activity, and at 10 °C, more than 70 % of the maximal activity was observed both for purified recombinant enzyme. Analysis of enzyme stability showed that purified, recombinant beta-galactosidase was 100% stable at 0 °C to 20 °C for at least 5 hours (Fig. 15), and that the residual activity after 5 days storage at 0 °C to 20 °C was 50-60 % (data not shown). At 30 °C, the purified beta-galactosidase lost more than 80 % of its activity within 5 hours. Complete, irreversible inactivation was achieved within 5 minutes at 50 °C and within 10 minutes at 40 °C (Fig. 15).

[0068] The Temperature dependence (A) and pH dependence (B) of the purified, recombinant A. *ikkense* beta-galactosidase was tested (Figure 14). Y-axis is the relative activity in percent of maximal activity. X-axis is the incubation temperature (A) or pH (B). The relative activity of the purified, recombinant enzyme with ONPG as substrate is illustrated by black squares. Assays were performed in triplicates, and standard error was below 0.05.

[0069] Maximal activity of the purified, recombinant enzyme was observed at pH 8 (Fig. 14B). About 60 % of the maximal activity was maintained at pH 7, and at pH 9 approximately 90 % activity was observed (Fig. 14B).

[0070] The recombinant A. *ikkense* beta-galactosidase was benchmarked with Lactozyme 3000® 13 from *K. lactis*. At temperatures between 0 °C and 20 °C the A. *ikkense* beta-galactosidase showed a twofold increase in conversion rate, when compared to the *K. lactis*-beta-galactosidase (Fig. 16); *A. ikkense* beta-galactosidase (black squares) compared with commercially available Lactozyme® 3000L from *K. lactis* (open circles). Specically the experiment was conducted in the following way: An equal amount of enzyme (2 mg/ml) was incubated with ONPG as substrate at temperatures from 0 °C to 20 °C. Samples were withdrawn at different time intervals and hydrolyzed ONP was measured at A420 nm. Hydrolysis efficiency was calculated as increase in A420 nm per min per microgram active enzyme. Assays were performed in triplicates, and standard error was below 0.05.

[0071] Investigation of substrate specificity of the *A. ikkense* β-galactosidase was carried out using nine different chromogenic substrates. Hydrolysis was only observed with ONPG and with p-nitrophenyl-beta-D-fucopyranoside (4 % of the relative activity compared to hydrolysis of ONPG). Thin-layer chromatography (TLC) was used to demonstrate the hydrolysis of lactose by the A. *ikkense* beta-galactosidase (Figure 17). Lanes 4-6: samples incubated at 5 °C for 2½ h (4: 1.25 mg/ml lactose, 5: 2.5 mg/ml

lactose, 6: 5 mg/ml lactose). Lanes 7-9: samples incubated at 20 °C for 2½ h (7: 1.25 mg/ml lactose, 8: 2.5 mg/ml lactose, 9: 5 mg/ml lactose). Lanes 1-3: Controls, 0.0125 mg of each of the carbohydrates lactose (lane 1), galactose (lane 2) and glucose (lane 3).

[0072] Hydrolysis of the remaining substrates was below detection limit. Hydrolysis of lactose was observed at both 5 °C and at 20 °C (Fig. 17). At 5 °C, approximately 75-85 % of the lactose was hydrolyzed in the 1.25 mg/ml reaction after 15 minutes, and 100 % of the lactose was hydrolyzed within 2½ hours (Fig. 17, lane 4). Similar hydrolysis efficiency was observed in the 1.25 mg/ml reaction incubated at 20 °C for 2½ hours (Fig. 17, lane 7). Hydrolysis effectiveness in the 2.5 mg/ml lactose reaction showed approximately 90-95 % hydrolysis within 2½ hours at both temperatures (Fig. 17, lanes 5 and 8). After 24 hours, 100 % hydrolysis was observed for all three lactose concentrations at both temperatures (not shown). At the highest lactose concentration (27 mg/ml), the TLC gel indicated the formation of oligosaccharides (Fig. 17, lanes 6 and 9).

EXAMPLE 6

Production of recombinant Ikka-beta-galactosidase in *Bacillus subtilis*.

[0073] The Ikka-beta-galactosidase was further subcloned in a *Bacillus subtilis* expression vector, pAL10 (MoBiTech, GmbH). PCR was performed using chromosomal DNA from *Alkalilactibacillus ikkense* as template and the PCR primers Bs_pAL_bGal5' : 5'GGCCATGGATCCATGGCAAAAAAATTAAAAAAATTC3' (*BamHI* restriction site underlined) (SEQ ID NO 11) and Bs_pAL_bGal3': 5'GGCCATCCCGGGTTATCTGTTTAAACTATTCAACATG3' *(XmaI* restriction site double underlined) (SEQ ID NO 12). PCR, subsequent isolation of the fragment encoding the Ikka-beta-galactosidase, ligation into pUC18dLacZ and transformation of *E. coli* was as described above in 5.1. Plasmid pUC18dLacZ carrying the Ikka-beta-galactosidase gene was prepared and sequenced, before the plasmid DNA was restricted with restriction endonucleases *BamH*I and *Xma*I. The 3.1 kb fragment encoding Ikka-beta-galactosidase was purified, inserted into plasmid pAL10 similarly restricted with *BamH*I and *Xma*I, and transformed into *E. coli* as described in 5.1. Recombinant *E. coli* harbouring pAL10 containing the Ikka-beta-galactosidase gene was isolated on LB agar plates containing 100 µg/ml ampicillin. Plasmid pAL10_Ikka-bGal was purified and transformed into B. *subtilis* cells using an electroporation protocol for *B. subtilis* from Eppendorf (Germany) (Protocol No. 4308 915.504 - 08/2003). Recombinant cells harbouring plasmid pAL10 were selected on LB agar containing 5 µg/ml chloramphenicol.

[0074] Production of recombinant Ikka-beta-galactosidase in *B. subtilis* was carried out by growing *B. subtilis* pAL10_Ikka-bGal cells in LB containing 5 µg/ml of chloramphenicol at 37 °C for 16 hours. Induction of Ikka-beta-galactosidase synthesis was carried out by changing the temperature to 20 °C. *B. subtilis* pAL10_Ikka-bGal cells were cultivated at 20 °C for another 5 hours after which, the cells were harvested, and intracellular enzyme was isolated by Fast Prep as described in 5.2.

[0075] A crude, intracellular extract from *B. subtilis* pAL10_Ikka-bGal cells was analysed in ONPG assays as described in 5.3. ONPG assays showed the presence of a cold-active Ikka-beta-galactosidase with an activity similar to the native enzyme, to the recombinant enzyme in crude *E. coli* extracts, and to the pure enzyme produced in *E. coli* cells.

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[0076]

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SEQ ID NO: 1

1. (1) GENERAL INFORMATION:
    1. (i) APPLICANT:
        1. (A) NAME: University of Copenhagen
        2. (B) STREET:
        3. (C) CITY:
        4. (D) COUNTRY:
        5. (E) POSTAL CODE:
    2. (ii) TITLE OF INVENTION: Cold-active beta-galactosidase, a method of producing same and use of such enzyme
    3. (iii) NUMBER OF SEQUENCES: 10
2. (2) INFORMATION FOR SEQ ID NO: 1

    (i) SEQUENCE CHARACTERISTICS:

    1. (A) LENGTH: 1,041 amino acids
    2. (B) TYPE: amino acid
    3. (C) STRANDEDNESS: Not relevant
    4. (D) TOPOLOGY:linear

    (ii) MOLECULE TYPE: protein

    (xi) SEQUENCE DESCRIPTION SEQ ID NO: 1

```
 1                                  10
Met Ala Lys Lys Leu Lys Lys Phe Asn Tyr Leu Pro Pro Lys Asn

                20                                  30
Gly Tyr Pro Glu Trp Asn Asn Asn Pro Glu Ile Phe Gln Leu Asn

                                    40
Arg Arg Glu Ala His Ala Thr Leu Val Pro Tyr Ser Asn Leu Glu

                50                                  60
Leu Ala Leu Lys Gly Glu Arg Thr Ala Ser Ser Phe Tyr Gln Ser

                                    70
Leu Asn Gly Ser Trp Gln Phe Ala Phe Ala Gln Glu Pro Thr Lys

                80                                  90
Arg Val Ile Asp Phe Tyr Arg Lys Asp Phe Asp His Arg Asp Trp

                                   100
Asp Ser Ile Lys Val Pro Ser His Trp Gln Leu Glu Gly Tyr Asp

               110                                 120
Tyr Pro Gln Tyr Thr Asn Thr Thr Tyr Pro Trp Val Glu Lys Glu

                                   130
Thr Ile Lys Pro Pro Phe Ala Pro Thr Asn Tyr Asn Pro Val Gly

               140                                 150
Gln Tyr Val Arg Thr Phe Glu Leu Pro Thr Asp Trp Asn Gly Ala

                                   160
Pro Val Tyr Leu Asn Phe Gln Gly Val Glu Ser Ala Phe Tyr Val

               170                                 180
Trp Ile Asn Gly Asp Leu Val Gly Tyr Ser Glu Asp Thr Phe Thr

                                   190
Pro Ala Glu Phe Asp Ile Thr Pro Tyr Leu Ile Glu Gly Glu Asn

               200                                 210
Lys Leu Ala Val Glu Val Tyr Arg Trp Ser Asp Ala Ser Trp Leu

                                   220
Glu Asp Gln Asp Phe Trp Arg Leu Ser Gly Ile Phe Arg Asp Val
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                230                                         240
Tyr Leu Tyr Ala Thr Pro Ala Gln His Ile Asp Asp Phe Phe Val

                                    250
Thr His Glu Leu Asp Ala Asp Tyr Arg Asn Ala Thr Leu Lys Ile

                260                                         270
Asp Met Lys Val Arg Asp Tyr Phe Glu Ile Gly Glu Pro Val Thr

                                    280
Val Asn Ala Met Leu Phe Asp Leu Asn Gly Asn Pro Val Leu Lys

                290                                         300
Gln Pro Leu Leu Ser Ala Val Asp Phe Ser Gly Lys Glu Val Ala

                                    310
Asp Val Ser Val Ile Thr Thr Ile Asp Asn Pro Leu Lys Trp Ser

                320                                         330
Ala Glu Asp Pro Asn Leu Tyr Thr Leu Val Leu Ser Leu Val Asp


                                    340
Gln Asn Gly Lys Leu Leu Glu Thr Glu Ser Cys Arg Val Gly Phe

                350                                         360
Arg Lys Phe Glu Arg Lys Asp Gly Leu Met Gln Ile Asn Gly Lys

                                    370
Arg Ile Val Phe Lys Gly Thr Asn Arg His Glu Phe Ala Ser Asp

                380                                         390
Lys Gly Arg Ala Ile Thr Ile Asp Asp Met Val Asn Asp Ile Gln

                                    400
Leu Met Lys Gln His Asn Ile Asn Ala Val Arg Thr Ser His Tyr

                410                                         420
Pro Asn His Pro Leu Trp Tyr Glu Leu Cys Asp Thr Tyr Gly Leu

                                    430
Tyr Val Ile Asp Glu Thr Asn Leu Glu Thr His Gly Thr Trp Val

                440                                         450
Tyr Gly Gln Lys Gly Leu Ala Glu Thr Ile Pro Gly Ser Leu Pro

                                    460
Lys Trp Thr Glu Asn Val Leu Asp Arg Cys Asn Ser Met Phe Gln
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                470                                             480
Arg Asp Lys Asn His Pro Ser Ile Leu Asp Trp Ser Leu Gly Asn

                                    490
Glu Ser Phe Gly Gly Asp Asn Phe Leu Lys Met His Asp Phe Phe

                500                                             510
Thr Glu Gln Asp Pro Ala Arg Leu Val His Tyr Glu Gly Ile Phe

                                    520
His Tyr Arg Glu Ser Glu Arg Ala Ser Asp Met Glu Ser Thr Met

                530                                             540
Tyr Ile Ser Pro Glu Gly Ile Glu Asp Tyr Ala Lys Lys Ala Thr

                                    550
Lys Glu Thr Lys Pro Tyr Ile Leu Cys Glu Phe Ser His Ala Met

                560                                             570
Gly Asn Ser Leu Gly Asn Phe Tyr Lys Tyr Thr Glu Leu Phe Asp


                                    580
Gln Tyr Pro Ile Leu Gln Gly Gly Phe Ile Trp Asp Trp Lys Asp

                590                                             600
Gln Ser Leu Leu Thr Lys Thr Ala Gly Gly Thr Pro Tyr Leu Ala

                                    610
Tyr Gly Gly Asp Phe Gly Glu Ser Pro His Asp Gly Asn Phe Ala

                620                                             630
Gly Asn Gly Leu Ile Phe Gly Asp Gly Lys Val Ser Pro Lys Ile

                                    640
Phe Glu Val Lys Arg Cys Tyr Gln Asn Val Asp Phe Lys Ala Ile

                650                                             660
Asp Leu Val His Gly Gln Ile Glu Leu Thr Asn Lys Tyr Leu Phe

                                    670
Thr Asn Leu Ala Asp Tyr Gln Leu Asn Trp Val Ile Thr Arg Asn

                680                                             690
Gly Asp Ala Ile Glu Ser Gly Ala Thr Asn Ile Asn Val Leu Pro

                                    700
Gly Glu Lys Arg Glu Val Ile Leu Asp Tyr Thr Phe Pro Thr Gly
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                710                                             720
Val Cys Met Thr Asp Glu Tyr Ile Leu Thr Leu Arg Phe Ser Glu

                                        730
Lys Gly Asp Arg Leu Trp Cys Glu Ala Gly His Glu Val Ala Phe

                740                                             750
Asn Gln Phe Val Leu Pro Thr Lys Val Thr Lys Leu Arg Glu Lys

                                        760
Thr Gln Asp Thr Lys Thr Leu Ser Val Glu Val Met Gln Asp Arg

                770                                             780
Leu Val Thr Ser Gly Ala Gly Phe Ser Val Gly Phe Asp Thr Lys

                                        790
Ser Gly Met Leu Val Ser Tyr Gln Val Gly Gly Asn Glu Leu Val

                800                                             810
Lys Glu Ala Leu Val Pro Asn Phe Trp Arg Ala Met Thr Asp Asn


                                        820
Asp Arg Gly Asn Gly Leu Asp Gln Arg Ser Gln Ile Trp Arg Asp

                830                                             840
Ala Asn Glu Val Arg Glu Leu Val Ser Phe Gln Tyr Glu Val Leu

                                        850
Thr Asn Arg Val Ser Ile Ser Thr Val Phe Leu Tyr Glu Asp Leu

                860                                             870
Asn His Ser Arg Val Glu Leu Asn Phe Leu Ile Thr Gly Thr Gly

                                        880
Glu Ile Lys Val Asp Tyr Val Leu Lys Pro Gly Glu Asp Leu Pro

                890                                             900
Glu Ile Pro Glu Ile Gly Leu Met Leu Thr Met Pro Lys Ser Phe

                                        910
Asp Gln Leu Ser Trp Tyr Gly Lys Gly Pro His Glu Ser Tyr Trp

                920                                             930
Asp Lys Gln Lys Gly Ala Lys Ile Gly Leu Tyr Gln Gly Phe Val

                                        940
Gly Asp Gln Tyr Val Pro Tyr Leu Lys Pro Gln Glu Cys Gly Asn

                950                                             960
Lys Val Gly Val Arg Ser Ala Glu Leu Val Asn Asp Val Gly Val

                                        970
Gly Leu Ile Ile Ser Gly Leu Pro Thr Leu Glu Leu Asn Val Leu

                980                                             990
Pro Tyr Thr Pro Val Gln Leu Glu Ser Ala Asp His Ser Tyr Gln

                                       1000
Leu Pro Glu Thr Asp Gln Thr Val Val Arg Ile Asn Leu Gly Gln

               1010                                            1020
Met Gly Val Gly Gly Asp Asp Ser Trp Gly Gln Arg Thr His Gln

                                       1030
Asp Phe Thr Leu Phe Ala Asn Lys Thr Tyr His Tyr Ser Phe Met

               1040
Leu Asn Ser Leu Asn Arg
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SEQ ID NO: 2

(2) INFORMATION FOR SEQ ID NO: 2

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 3,123nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: genomic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 2

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ATGGCAAAAA AATTAAAAAA ATTCAACTAC CTCCCACCAA AAAACGGGTA   50

CCCAGAGTGG AATAATAATC CGGAAATTTT TCAACTTAAT CGAAGAGAGG  100

CGCATGCAAC ATTGGTGCCA TATTCTAATT TGGAATTGGC ACTTAAAGGG  150

GAGCGGACAG CATCATCATT TTATCAATCT TTAAATGGTA GTTGGCAGTT  200

TGCCTTTGCC CAAGAGCCAA CCAAGCGAGT GATAGATTTT TATCGGAAAG  250

ATTTTGATCA TCGCGATTGG GATTCGATTA AAGTACCAAG TCATTGGCAG  300

TTAGAAGGCT ATGACTACCC GCAATACACC AACACAACGT ACCCATGGGT  350

AGAAAAAGAA ACGATTAAAC CTCCATTTGC ACCAACAAAT TATAATCCAG  400

TCGGACAATA TGTTCGCACG TTTGAATTAC CGACTGATTG GAATGGAGCT  450

CCCGTTTATC TGAATTTCCA AGGTGTTGAA TCAGCTTTCT ACGTCTGGAT  500

AAATGGTGAT TTGGTCGGAT ACAGTGAGGA CACTTTCACA CCAGCTGAAT  550

TTGATATAAC TCCCTATTTA ATAGAGGGTG AAAATAAGCT AGCGGTAGAA  600

GTCTATCGTT GGAGTGATGC GAGCTGGCTT GAAGACCAGG ATTTCTGGAG  650

GTTAAGCGGG ATTTTTCGTG ACGTCTATCT ATATGCAACA CCAGCTCAGC  700

ACATTGATGA TTTCTTTGTC ACACACGAAC TTGATGCAGA CTATCGAAAT  750

GCAACGTTGA AGATTGATAT GAAAGTGCGC GATTATTTCG AGATTGGCGA  800

GCCTGTCACA GTTAATGCGA TGCTCTTTGA TCTTAATGGG AATCCGGTTC  850

TCAAGCAACC GCTTTTATCG GCAGTAGATT TTTCAGGTAA AGAAGTTGCT  900

GATGTGAGCG TAATAACAAC AATTGATAAT CCATTGAAAT GGAGTGCGGA  950

AGATCCCAAT CTGTACACTT TGGTTTTAAG TTTAGTTGAT CAGAATGGCA 1000

AGTTGCTTGA AACAGAAAGC TGTCGCGTTG GATTTCGTAA ATTTGAACGC 1050

AAGGACGGAT TGATGCAAAT TAATGGAAAG CGGATTGTCT TTAAAGGGAC 1100

AAATCGTCAC GAATTCGCTT CTGATAAAGG TCGGGCGATA ACGATAGATG 1150

ATATGGTTAA TGATATTCAG CTGATGAAGC AGCATAACAT TAATGCCGTT 1200

CGAACCTCAC ATTATCCGAA TCATCCGCTT TGGTATGAGT TGTGTGATAC 1250

GTATGGGTTA TATGTGATTG ACGAGACAAA CTTAGAGACG CACGGGACAT 1300

GGGTTTATGG TCAAAAAGGA TTGGCTGAGA CAATACCAGG TAGTCTACCA 1350

AAGTGGACTG AAAACGTCTT GGATCGTTGT AATTCAATGT TCCAACGTGA 1400

TAAAAACCAC CCATCGATTC TGGATTGGTC ACTTGGTAAT GAATCTTTTG 1450

GTGGCGATAA CTTCTTGAAG ATGCATGACT TCTTTACGGA ACAAGATCCA 1500
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GCTCGTCTGG TGCACTATGA GGGGATTTTT CATTATCGTG AATCTGAACG 1550

GGCATCTGAT ATGGAGAGTA CCATGTATAT TTCGCCAGAA GGCATTGAGG 1600

ACTATGCAAA GAAAGCGACC AAGGAGACGA AACCATATAT TTTATGCGAA 1650

TTCAGCCATG CGATGGGCAA CTCGCTAGGA AACTTTTATA AGTATACCGA 1700

GCTATTTGAT CAATATCCGA TCTTACAAGG AGGCTTCATT TGGGATTGGA 1750

AGGATCAATC GCTGCTAACG AAGACAGCAG GAGGCACACC GTATCTTGCT 1800

TATGGTGGTG ATTTTGGTGA ATCGCCACAC GACGGCAACT TTGCTGGTAA 1850

TGGTTTGATT TTTGGAGATG GCAAGGTTAG CCCGAAGATT TTTGAAGTGA 1900

AGCGTTGTTA CCAAAATGTT GATTTCAAAG CAATAGACTT AGTGCACGGA 1950

CAAATCGAAT TGACCAATAA ATACTTGTTC ACCAATCTCG CTGACTACCA 2000

ACTAAATTGG GTTATCACTC GAAACGGTGA TGCAATAGAG TCGGGTGCTA 2050

CTAACATCAA TGTCTTACCA GGTGAAAAAA GAGAGGTTAT ACTTGACTAC 2100

ACGTTCCCAA CAGGCGTTTG CATGACGGAT GAATATATTT TGACCCTTCG 2150

TTTTTCTGAG AAAGGTGATC GCTTATGGTG TGAAGCGGGA CATGAAGTTG 2200

CATTTAATCA GTTTGTTTTA CCAACAAAAG TTACGAAATT ACGTGAGAAG 2250

ACACAAGATA CCAAGACGCT TTCAGTTGAA GTAATGCAAG ATCGACTTGT 2300

TACATCTGGT GCTGGATTTA GCGTCGGATT TGACACTAAA TCGGGTATGC 2350

TTGTTTCTTA CCAAGTTGGA GGTAATGAAT TGGTGAAAGA GGCACTTGTG 2400

CCAAACTTCT GGCGTGCAAT GACTGATAAT GATCGCGGGA ACGGACTCGA 2450

TCAACGGAGT CAGATTTGGC GTGATGCAAA TGAGGTACGT GAATTGGTTT 2500

CATTTCAGTA TGAAGTGTTG ACCAATAGAG TAAGCATATC AACGGTTTTC 2550

TTATATGAAG ACCTCAACCA TTCACGCGTT GAACTTAACT TTTTGATTAC 2600

TGGAACTGGT GAAATAAAGG TGGATTATGT ACTGAAACCG GGAGAAGATT 2650

TACCAGAAAT ACCAGAGATA GGTTTGATGT TAACGATGCC TAAGTCGTTT 2700

GATCAGTTAA GTTGGTATGG AAAAGGCCCA CATGAATCGT ATTGGGATAA 2750

ACAAAAAGGC GCGAAAATAG GTCTTTATCA AGGATTTGTC GGCGATCAGT 2800

ATGTGCCGTA TTTGAAACCA CAAGAATGTG GCAACAAAGT AGGAGTTCGT 2850

TCAGCAGAAT TGGTTAATGA TGTTGGTGTT GGTTTGATTA TAAGTGGACT 2900

TCCAACGCTG GAGTTAAATG TCTTACCATA CACACCAGTG CAACTGGAAT 2950

CAGCTGATCA TAGCTATCAA TTACCAGAAA CAGATCAGAC TGTTGTGCGT 3000

ATTAATTTAG GACAAATGGG AGTTGGTGGT GATGATAGTT GGGGACAGCG 3050

AACACACCAA GACTTTACCT TATTTGCAAA TAAAACCTAT CACTATAGCT 3100

TCATGTTGAA TAGTTTAAAC AGA 3123

SEQ ID NO: 3

(2) INFORMATION FOR SEQ ID NO: 3

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 17 nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: synthetic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 3
CCGTCATCCA TATCACC

SEQ ID NO: 4

(2) INFORMATION FOR SEQ ID NO: 4

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 20 nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: synthetic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 4
CCTTTGCCCA AGAGCCAACC

SEQ ID NO: 5

(2) INFORMATION FOR SEQ ID NO: 5

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 21 nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: synthetic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 5
GCTATTATCA GACTTGGCAC C

SEQ ID NO: 6

(2) INFORMATION FOR SEQ ID NO: 6

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 21 nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: synthetic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 6
GTAATTCAAT GTTCCAACGT G

SEQ ID NO: 7

(2) INFORMATION FOR SEQ ID NO: 7

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 16 nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: synthetic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 7
CGCTTATGGT GTGAAG

SEQ ID NO: 8

(2) INFORMATION FOR SEQ ID NO: 8

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 20 nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: synthetic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 8
GGGCTGGCTT AACTATGCGG

SEQ ID NO: 9

(2) INFORMATION FOR SEQ ID NO: 9

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 36 nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: synthetic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 9
CTGAATTCGC ATATGGCAAA AAAATTAAAA AAATTC

SEQ ID NO: 10

(2) INFORMATION FOR SEQ ID NO: 10

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 31 nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: synthetic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 10
CCAAGCTTAT CTGTTTAAAC TATTCAACAT G

SEQ ID NO: 11

(2) INFORMATION FOR SEQ ID NO: 11

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 36 nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: synthetic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 11
GGCCATGGAT CCATGGCAAA AAAATTAAAA AAATTC

SEQ ID NO: 12

(2) INFORMATION FOR SEQ ID NO: 12

(i) SEQUENCE CHARACTERISTICS:

1. (A) LENGTH: 37 nucleotides
2. (B) TYPE: nucleic acid
3. (C) STRANDEDNESS: single
4. (D) TOPOLOGY:linear

(ii) MOLECULE TYPE: synthetic DNA

(xi) SEQUENCE DESCRIPTION SEQ ID NO: 12
GGCCATCCCG GGTTATCTGT TTAAACTATT CAACATG

REFERENCES CITED IN THE DESCRIPTION

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.

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KOLD-AKTIV BETA-GALACTOSIDASE, FREMGANGSMÅDE TIL FREMSTILLING HERAF OG ANVENDELSE AF ET SÅDANT ENZYM

Patentkrav

1. Oprenset kold-aktiv beta-galactosidase, der har aminosyresekvensen som defineret i SEQ ID NO: 1 eller har én mindst 80 % homologi med aminosyresekvensen som defineret i SEQ ID NO: 1, hvilken aminosyresekvens er udvalgt, således at enzymet har en stabil enzymatisk aktivitet ved temperaturer på mindre end 8 °C.

2. Beta-galactosidase ifølge krav 1, hvor aminosyresekvensen har mindst 90 %, fortrinsvis 95 %, homologi med aminosyresekvensen som defineret i SEQ ID NO: 1.

3. Beta-galactosidase ifølge krav 1 eller 2, hvor den er fremstillet af en stamme af en Alkalilactibacillus.

4. Beta-galactosidase ifølge krav 1, 2 eller 3, hvor den er fremstillet af Alkalilactibacillus ikkense.

5. Isoleret DNA-sekvens omfattende et gen, der koder for beta-galactosidasen ifølge et hvilket som helst af kravene 1 til 4.

6. Isoleret DNA-sekvens, der

a) koder for et protein med en aminosyresekvens som i SEQ ID NO: 2, eller

b) hybridiserer under stringente betingelser til sekvensen af a), eller

c) er degenerativ af sekvensen af a) eller b).

7. DNA-sekvens ifølge krav 6, hvor sekvensen er som i SEQ ID NO: 2.

8. Rekombinant vektor omfattende en DNA-sekvens ifølge et hvilket som helst af kravene 5 til 7.

9. Vektor ifølge krav 8, hvor vektoren er en ekspressionsvektor.

10. Værtscelle transformeret med en vektor ifølge krav 8 eller 9.

11. Celle ifølge krav 10, hvor cellen er udvalgt fra gruppen bestående af Escherichia, Bacillus, Bifidobacterium, Lactococcus, Lactobacillus, Streptomyces, Leuconostoc, Streptomyces, Saccharomyces, Kluyveromyces, Candida, Torula, Torulopsis og Aspergillus.

12. Celle ifølge krav 11, hvor cellen er udvalgt fra gruppen bestående af Bifidobacterium breve, Bifidobacterium longum, Bifidobacterium infantis, Bifidobacterium bifidum, Bifidobacterium animalis og Lactococcus lactis.

13. Anvendelse af en celle ifølge et hvilket som helst af kravene 10 til 12 til fremstilling af et produkt udvalgt fra gruppen bestående af lactosefri mælk, lavlactosemælk, yoghurt, ost, fermenterede mælkeprodukter, kosttilskud og probiotisk spiseligt produkt.

14. Anvendelse ifølge krav 13 til fremstilling af et mejeriprodukt med en lactosekoncentration på 1 % eller lavere.

15. Anvendelse ifølge krav 14, hvor lactosekoncentrationen er 0,1 % eller lavere.

16. Psykrofil bakterie Alkalilactibacillus ikkense BCCM-accessionsnr. LMG P-24866 og varianter og mutanter afledt deraf, der er i stand til at fremstille en kold-aktiv beta-galactosidase defineret i krav 1 eller 2.

DRAWINGS

Figure 1

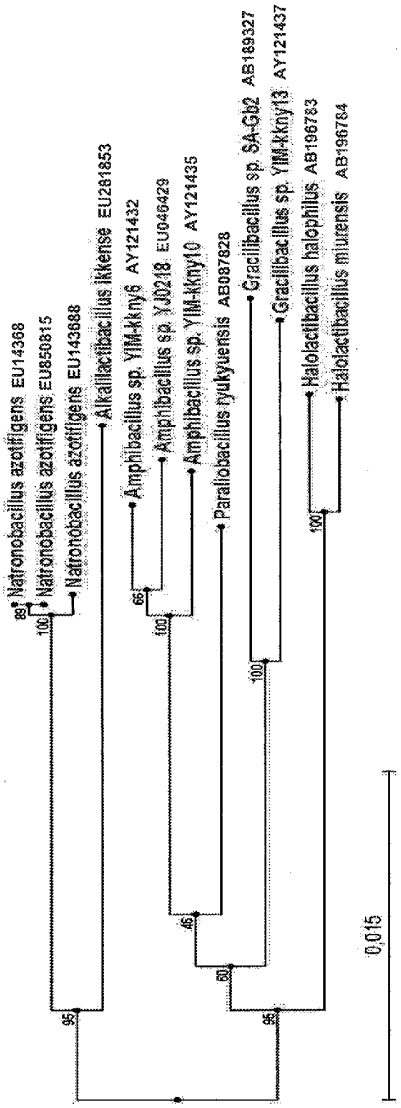


Figure 2

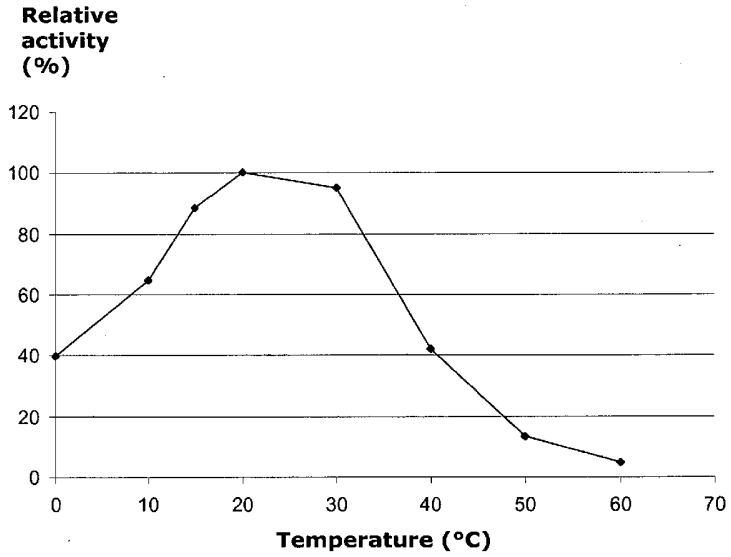

Relative activity (%)
120
100
80
60
40
20
0
0
10
20
30
40
50
60
70
Temperature (°C)

Figure 3

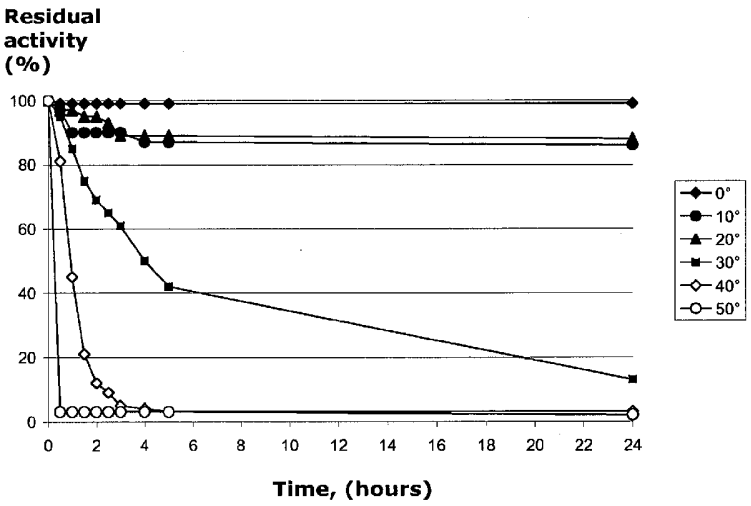

Residual activity (%)
100
80
60
40
20
0
0
2
4
6
8
10
12
14
16
18
20
22
24
Time, (hours)
0°
10°
20°
30°
40°
50°

Figure 4

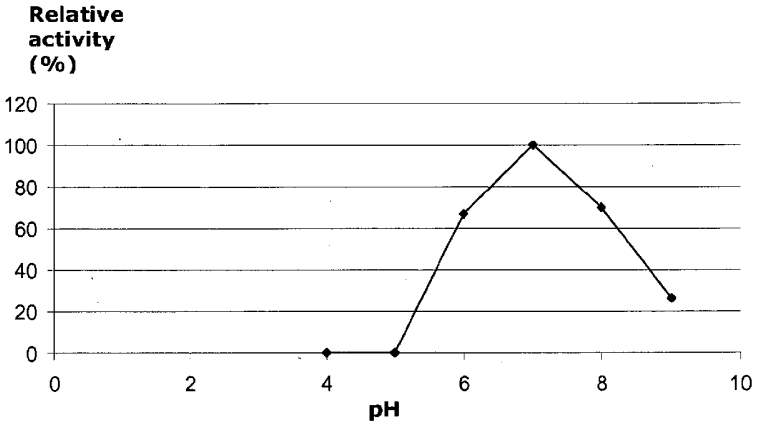


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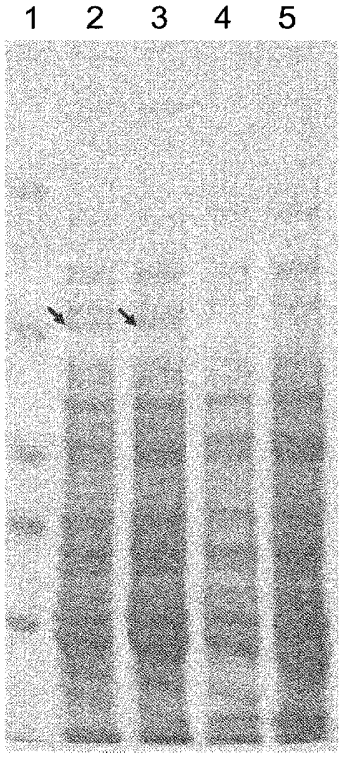


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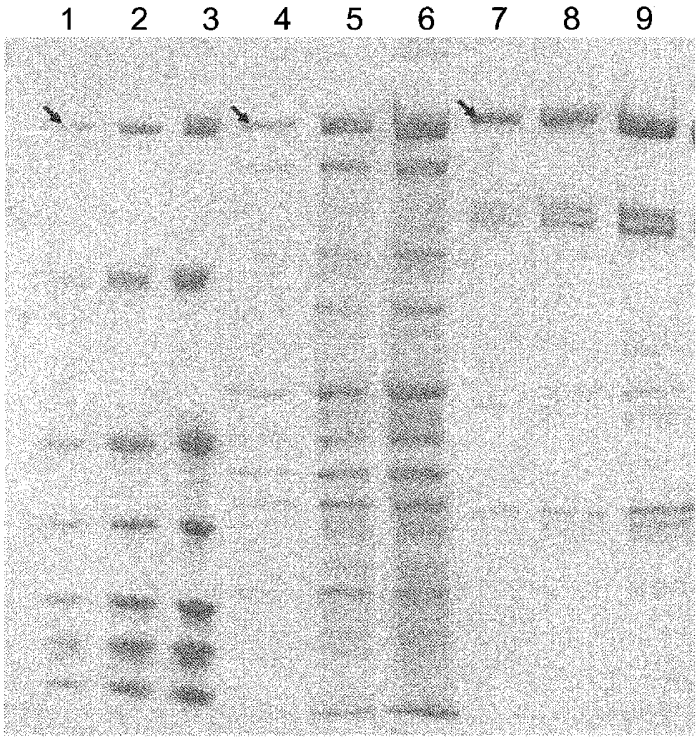


Figure 7

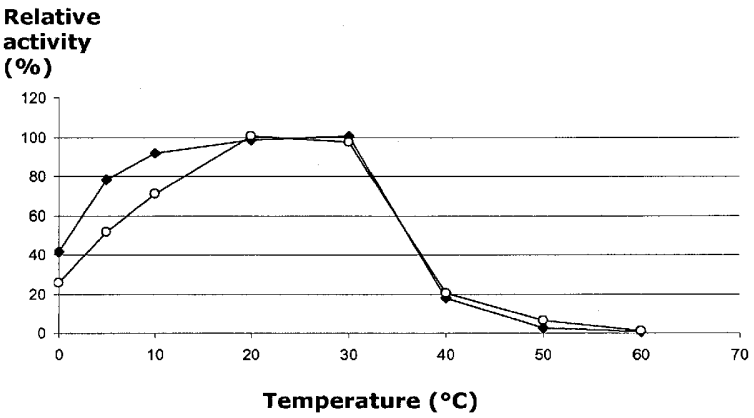

Relative activity (%)
120
100
80
60
40
20
0
0
10
20
30
40
50
60
70
Temperature (°C)

Figure 8

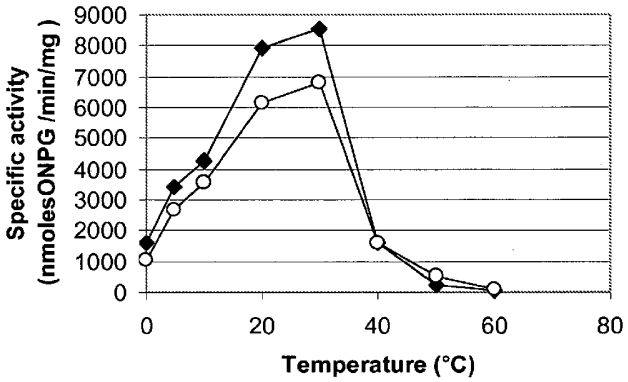

Specific activity
(nmolesONPG /min/mg)
9000
8000
7000
6000
5000
4000
3000
2000
1000
0
0
20
40
60
80
Temperature (°C)

Figure 9

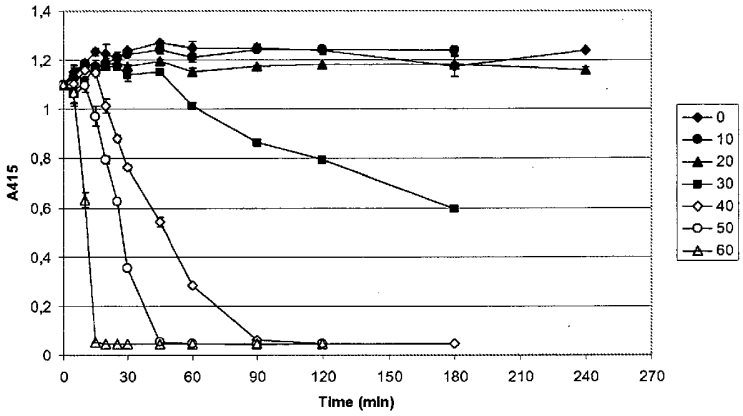

1,4
1,2
1
0,8
0,6
0,4
0,2
0
A415
0
30
60
90
120
150
180
210
240
270
Time (min)
0
10
20
30
40
50
60

Figure 10

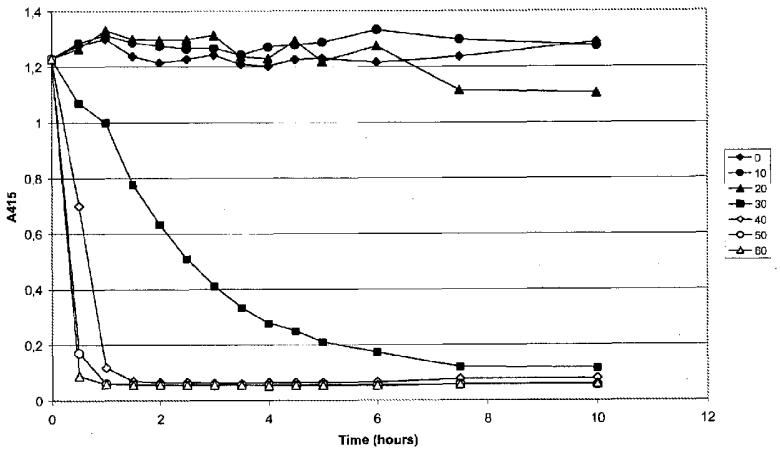


Figure 11

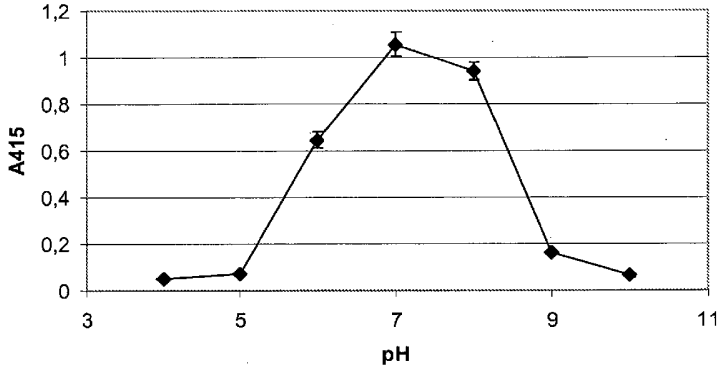
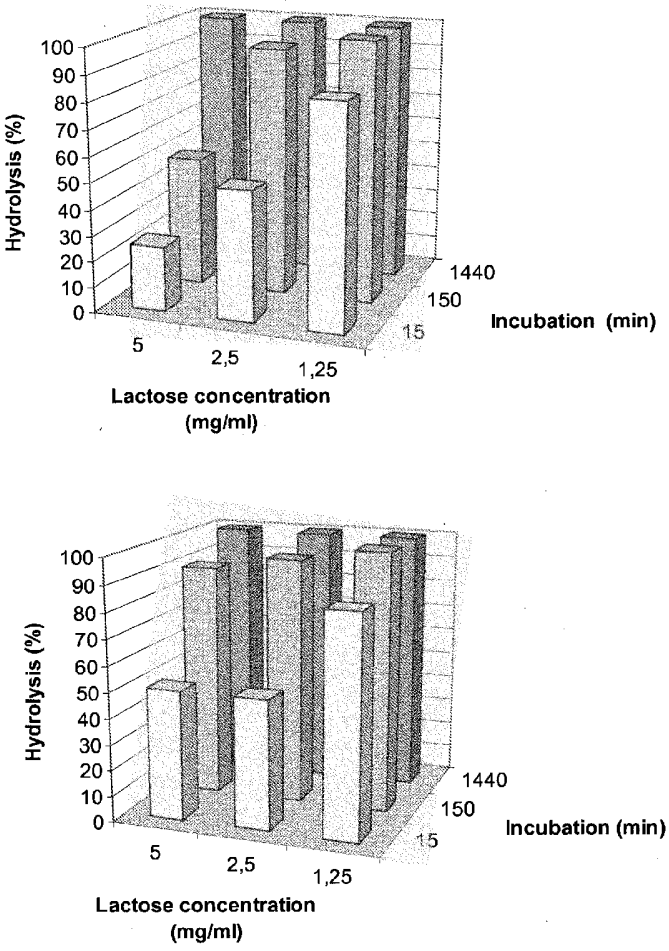


Figure 12


Hydrolysis (%)
100
90
80
70
60
50
40
30
20
10
0
5
2,5
1,25
Lactose concentration (mg/ml)
1440
150
15
Incubation (min)

Hydrolysis (%)
100
90
80
70
60
50
40
30
20
10
0
5
2,5
1,25
Lactose concentration (mg/ml)
1440
150
15
Incubation (min)

Figure 13

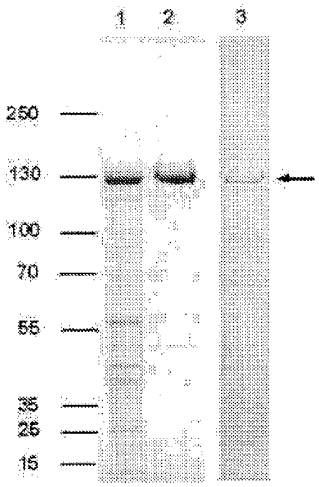


Figure 14

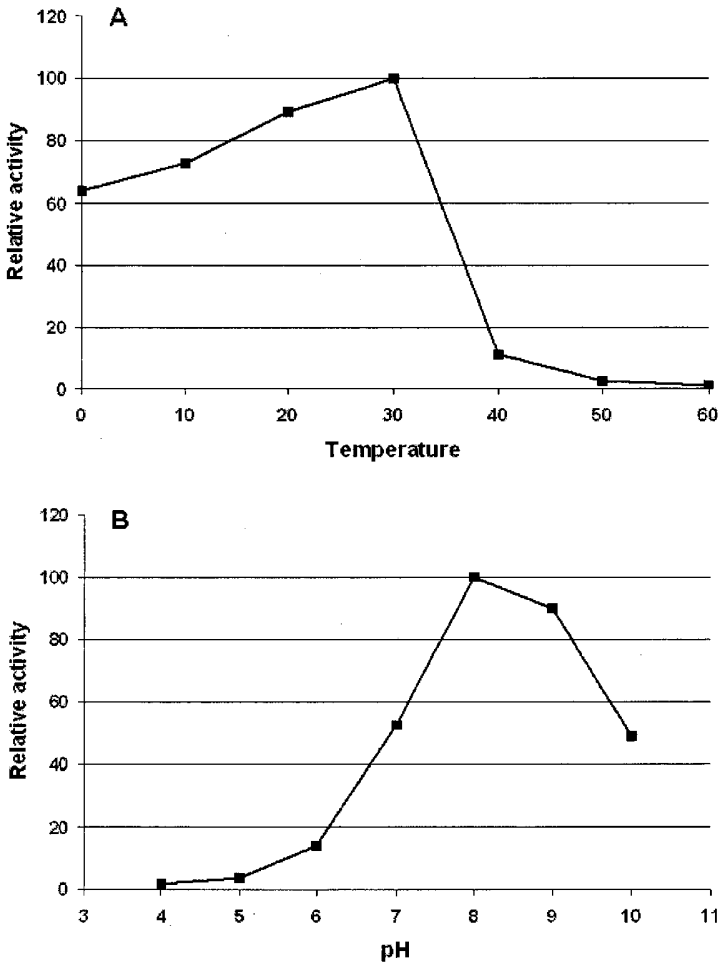

A
120
100
80
60
40
20
0
Relative activity
0
10
20
30
40
50
60
Temperature
B
120
100
80
60
40
20
0
Relative activity
3
4
5
6
7
8
9
10
11
pH

Figure 15

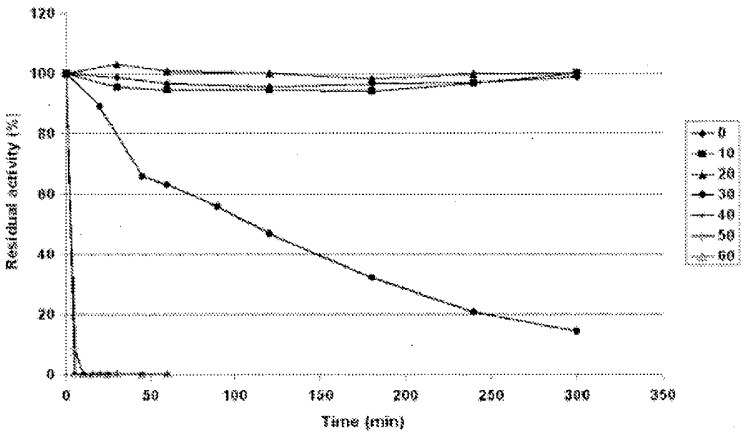

120
100
80
60
40
20
0
Residual activity (%)
0
50
100
150
200
250
300
350
Time (min)
0
10
20
30
40
50
60

Figure 16

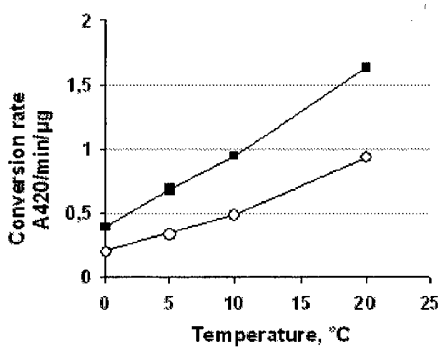

Conversion rate
A420/min/µg
2
1,5
1
0,5
0
0
5
10
15
20
25
Temperature, °C

Figure 17

