

UNIVERSITI PUTRA MALAYSIA

IMPROVEMENT OF CYCLODEXTRIN GLYCOSYLTRANSFERASE BIOSYNTHESIS BY RECOMBINANT Lactococcus lactis NZ:NSP:CGT

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FBSB 2014 13



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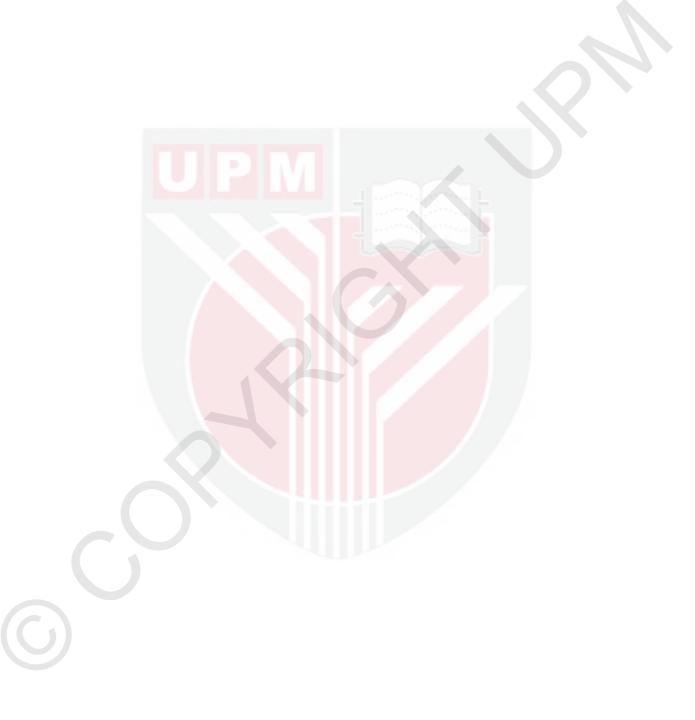
AZIN AMIRI

Thesis Submitted to the School of Graduate Studies, Universiti Putra Malaysia, in Fulfilment of the Requirement for the Degree of Master of Science

March 2014

Dedicated to

my beloved parents



Abstract of thesis presented to the Senate of Universiti Putra Malaysia in fulfillment of the requirement for the degree of Master of Science

IMPROVEMENT OF CYCLODEXTRIN GLYCOSYLTRANSFERASE BIOSYNTHESIS BY RECOMBINANT *Lactococcus lactis* NZ:NSP:CGT

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March 2014

Chairman:Assoc. Prof. Rosfarizan Mohamad, PhDFaculty:Biotechnology and Biomolecular Sciences

Cyclodextrin glycosyltransferase (CGTase) is a distinctive enzyme that has the capability of producing cyclodextrin (CD) from starch. The CD as the product of CGTase has numerous applications in various industries such as foods, cosmetics and toiletries, textiles and agrochemistry. Therefore, CGTase is considered as an industrially important enzyme and its production improvement is very crucial. So, essential efforts to increase its activity are desirable. CGTase production has never been investigated in Generally Regarded as Safe (GRAS) organism, Lactococcus lactis despite its advantages. The CGTase biosynthesis by recombinant Lactococcus lactis NZ:NSP:CGT using different carbon sources ((corn starch), potato (dextrin from starch), tapioca starch and several soluble potato starches) and nitrogen sources (yeast extract, meat extract, peptone from meat, peptone from soymeal and peptone from casein) was carried out in batch cultivation using 250 mL shake-flask. Statistical optimization was performed using artificial neural network technique in order to optimize the culture condition (temperature) and medium compositions (carbon and nitrogen sources concentrations) to achieve maximum CGTase production. The experimental data from the aforementioned fermentation experiments were analyzed in order to obtain the kinetic parameter values and establish the basis of a kinetic model. The optimum parameters obtained were used to run batch fermentation in a 2L stirred tank bioreactor. The best carbon source leading to maximum CGTase biosynthesis was determined as Nacalai Tesque GR soluble potato starch. The maximum CGTase activity and productivity obtained by this carbon source were 7.99 U/mL and 1 U/mL.h, respectively. Yeast extract (Merck) was selected as the best nitrogen source due to its highest CGTase activity (9.88 U/mL) and productivity (0.99 U/mL.h) obtained. In screening stage of CGTase fermentation, carbon source concentration, nitrogen source concentration and temperature were recognized as three significant fermentation parameters. The optimum values for these parameters were determined through statistical optimization as 20°C for temperature and 3.82 and 5.67% (w/v) of soluble starch and yeast extract concentrations, respectively. The maximum CGTase activity obtained using the optimum values was 22.09 U/mL, which was closed to the predicted value (24.17 U/mL). The models used in this study were based on unstructured model equations including logistic and Luedeking-Piret, which were suitable to explain the growth, substrate consumption and CGTase production by L. lactis NZ:NSP:CGT in batch cultivation. According to the results, CGTase

production is a growth-associated process. Production of CGTase in 2L stirred tank bioreactor (15.36 U/mL) was lower than shake-flask, which shows the essential optimization studies in bioreactor scale.



Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai memenuhi keperluan ijazah Master Sains

PENAMBAHBAIKAN BIOSINTESIS SIKLODEKSTRIN GLIKOSILTRANSFERASE OLEH *Lactococcus lactis* NZ:NSP:CGT REKOMBINAN

Oleh

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Mac 2014

Pengerusi:Assoc. Prof. Rosfarizan Mohamad, PhDFakulti:Bioteknologi dan Sains Biomolekul

Siklodekstrin glikosiltranferase (CGTase) ialah enzim yang jelas berbeza dan berkeupayaan menghasilkan siklodekstrin (CD) daripada kanji. CD sebagai produk CGTase mempunyai banyak kegunaan dalam pelbagai industri seperti industri makanan, kecantikan dan kelengkapan kebersihan diri, pakaian dan agrokimia. Oleh itu, CGTase adalah dianggap sebagai enzim yang penting dalam industri dan penghasilannya adalah amat penting. Justeru itu, penghasilan CGTase daripada organisma yang boleh dianggap selamat (GRAS) seperti Lactococcus lactis NZ:NSP:CGT menggunakan sumber karbon berbeza; (kanji jagung, ubi kentang (dekstrin daripada kanji), kanji ubi kavu dan beberapa jenis kanji boleh larut daripada ubi kayu) dan sumber nitrogen (ekstrak yis, ekstrak daging, pepton daging, pepton minyak kacang soya dan pepton kasein) telah dilakukan dalam pengkulturan sesekelompok menggunakan 250 mL kelalang kon. Pengoptimuman secara statistik telah dibuat menggunakan teknik rangkaian neural tiruan untuk mengoptimumkan keadaan kultur dan komposisi media untuk memperoleh penghasilan CGTase yang paling maksima. Data eksperimen daripada eksperimen fermentasi yang disebutkan telah dianalisis untuk memperoleh nilai parameter kinetik dan membuat satu model kinetik asas. Parameter optimum yang diperoleh telah digunakan untuk melakukan fermentasi kelompok di dalam bioreaktor tangki berpengaduk 2L. Sumber karbon terbaik yang dapat menghasilkan CGTase secara maksima adalah kanji boleh larut daripada ubi kayu Nacalai Tesque GR. Aktiviti dan penghasilan maksima CGTase yang diperoleh menerusi sumber karbon ini adalah 7.99 U/mL bagi aktiviti dan 1.00 U/mL.h untuk penghasilannya. Ekstrak yis (Merck) dipilih sebagai sumber nitrogen terbaik disebabkan aktiviti (9.88 U/mL) dan produktiviti (0.99 U/mL.h) CGTase yang tinggi diperoleh daripadanya. Dalam peringkat saringan fermentasi CGTase, kepekatan sumber karbon, kepekatan sumber nitrogen dan suhu telah dikenalpasti sebagai parameter fermentasi yang penting. Nilai optimum untuk parameter ini telah ditentukan menerusi pengoptimuman secara statistik di mana suhunya adalah 20°C, 3.82% (w/v) kanji boleh larut dan 5.67% (w/v) kepekatan ekstrak yis. Aktiviti CGTase paling maksima yang diperoleh dengan menggunakan nilai optimum adalah 22.09 U/mL, di mana berdekatan dengan nilai jangkaan (24.17 U/mL). Model yang digunakan dalam kajian ini adalah berdasarkan persamaan model tidak berstruktur termasuk Logistik dan Luedeking-Piret yang sesuai untuk menerangkan proses pertumbuhan, penggunaan sumber dan penghasilan CGTase oleh L. lactis NZ:NSP:CGT dalam pengkulturan sesekelompok. Berdasarkan kepada keputusan, penghasilan CGTase adalah proses berkaitan dengan pertumbuhan. Penghasilan CGTase di dalam bioreaktor tangki berpengaduk 2L (15.36 U/mL) adalah lebih rendah berbanding di dalam kelalang kon dimana menunjukkan pengoptimuman adalah penting untuk dikaji dalam skala bioreaktor.



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I certify that a Thesis Examination Committee has met on (**6 of March**) to conduct the final examination of Azin Amiri on her thesis entitled "**Improvement of Cyclodextrin Glycosyltransferase Biosynthesis by Recombinant** *Lactococcus lactis* **NZ:NSP:CGT**" in accordance with the University and University Colleges Act 1971 and the Constitution of the Universiti Putra Malaysia [P.U. (A) 106] 15 March 1998. The Committee recommends that the student be awarded the M.Sc degree.

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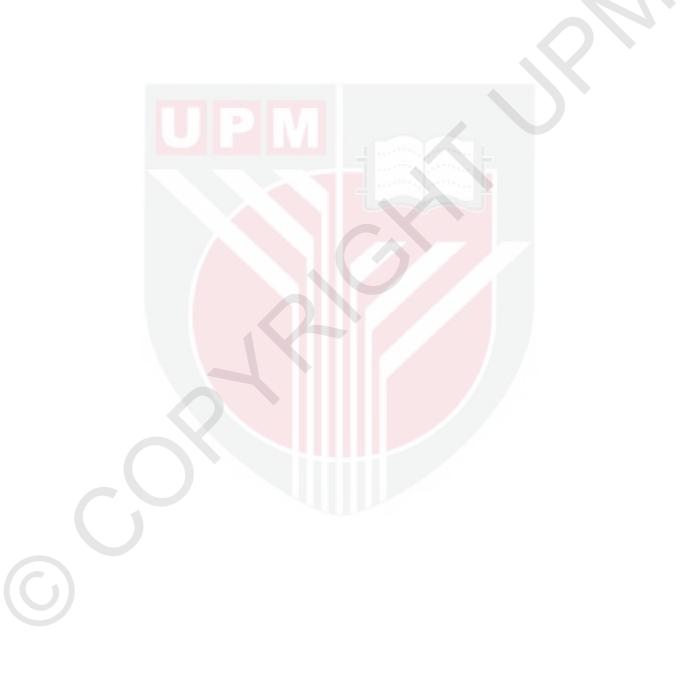
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LIST OF ABBREVIATIONS

AAD	Absolute Average Deviation
CCD	Central Composite Design
CD	Cyclodextrin
CGTase	Cyclodextrin Glycosyltransferase
DCW	Dry Cell Weight
DOT	Dissolved Oxygen Tension
E. coli	Escherichia coli
EC	Enzyme Classification
Eq	Equation
F. S.	Friendemann Schmidt
GRAS	Generally Recognized as Safe
IBP	Incremental Back Propagation
IUBMB	International Union of Biochemistry and Molecular Biology
L. lactis	Lactococcus lactis
LAB	Lactic Acid Bacteria
max	Maximum
OD	Optical Density
PBD	Placket-Burman Design
Pr	Productivity
RMSE	Minimum Root Square Error
RSM	Response Surface Methodology
μ	Specific Growth Rate

CHAPTER 1

INTRODUCTION

Cyclodextrin glycosyltransferase (EC 2.4.1.19; CGTase), is a "carbohydrateconverting", "bacterial glycosyltransferase" (Subramaniam *et al.*, 2012). CGTase is an enzyme that catalyzes the conversion of starch and related substances to cyclodextrins through cyclization reaction (Ibrahim *et al.*, 2005). It catalyzes other transferase reactions including disproportion and coupling beside cyclization. CGTase displays minor hydrolysis activity as well (Rahman *et al.* 2004).

Cyclodextrin (CD) is a cyclic malto-oligosaccharide molecule, which is formed of 6 to 60 glucose monomers (Vassileva *at al.*, 2003). CDs most commonly synthesized are α -, β - and γ -CDs, which consist of 6, 7 and 8 glucose units (Mahat *et al.*, 2004). CD holds a hydrophobic central cavity and a hydrophilic outer surface (Vassileva *et al.*, 2003).

According to Dodziuk (2006a, b) and Uekama *et al.* (2006), various molecules can enter the CDs' cavity and almost all the applications of CDs include their inclusion complex formation capability leading to their wide uses in different industries. For instance, CDs are applied in foods, cosmetics and toiletries, textiles and agrochemistry. In food industry, CDs are employed for stabilization by powdering (flavor or spices, fish oil, coffee, green tea), taste modification, anti-oxidation and improvement of bioavailability. CDs applications are also expanded to pharmaceuticals. They are employed to study different properties of drugs such as release control, site-specific drug delivery, absorption enhancement and so on. They can also be utilized in gene therapies. CDs also assist in improving the solubility and stability, reducing volatility and masking odors and tastes which result in increased popularity with their extensive use in various industries (Sian *et al.*, 2005). The vast applications of CD, increases the attentions focused on CGTase.

Microorganisms synthesize CGTase in order to catalyze the conversion of starch present in their environment to cyclodextrin for the purpose of growth and survival (Wang *et al.*, 2005). Although, enhancement of CGTase biosynthesis is of great interest (due to improvement of CD production), there is no report on CGTase production by *L. lactis* species. Therefore, no studies on optimization of culture conditions and medium compositions for the purpose of maximum CGTase production by *L. lactis* have been conducted. Recently, Subramaniam *et al.* (2012) has constructed a recombinant *L. lactis* strain capable of producing CGTase. The CGTase gene originally from *Bacillus* sp. G1 (Illias *et al.*, 2003) was cloned in *Escherichia coli* (Ong *et al.*, 2008). According to Subramaniam *et al.* (2012), the CGTase quality might be reduced in *Bacillus* and *E. coli* due to presence of some impurities such as proteases. Therefore, CGTase production studies in *L. lactis* are desirable.

The CGTase activity obtained by recombinant *L. lactis* NZ:NSP:CGT strain was very low and needed improvement through fermentation techniques. There are various methods available for fermentation optimization. One of the useful techniques is mathematical optimization with different tools available. Artificial

neural network (ANN) is one of the softwares employed for optimization of various product formation processes through fermentation process. There is no literature available in regard to CGTase biosynthesis using ANN. In this study, ANN was employed for the purpose of optimization of CGTase biosynthesis by recombinant *L. lactis* NZ:NP:CGT.

In most fermentation processes, mathematical models are required to control, optimize, simulate and scale up of the process in lots of unit operations (Rosfarizan, 2000). Information on the kinetics and modeling of CGTase production is very scarce. Therefore, a set of experiments is necessary in order to develop better understanding of CGTase fermentation process. The results could be used for estimation of kinetic parameters, which are prerequisite for mathematical model development. The model helps in better understanding of the whole process as well as control of CGTase biosynthesis by *L. lactis* NZ:NSP:CGT. This study is one of the first attempts in terms of studying the CGTase fermentation kinetics and modeling.

Generally, the focus of this study was on development of a process aimed at establishing high performance CGTase fermentation using a newly constructed recombinant *L. lactis* NZ:NSP:CGT strain. The objectives of this research were;

- 1. To evaluate the influences of different types of carbon and nitrogen sources on CGTase biosynthesis by *L. lactis* NZ:NSP:CGT in shake-flask culture.
- 2. To improve the CGTase biosynthesis by *L. lactis* NZ:NSP:CGT through medium compositions and environmental conditions optimization by statistical experimental design techniques.
- 3. To perform the kinetics and modeling of CGTase fermentation by *L. lactis* NZ:NSP:CGT in shake-flask scale followed by evaluation of CGTase biosynthesis in a 2L stirred tank bioreactor system.

REFERENCES

- Ahmed, E. and El-Refai, H. (2010). Cyclodextrin glucosyltransferase production by *Bacillus megaterium* NCR: evaluation and optimization of culture conditions using factorial design. *Indian journal of microbiology*. 50: 303–308.
- Albers, E. and Müller, B. (1992). Complexation of steroid hormones with cyclodextrin derivatives: substituent effects of the guest molecule on solubility and stability in aqueous solution. *Journal of pharmaceutical sciences*. 81(8): 756–761.
- Ammor, M., Flórez, A. B. and Mayo, B. (2007). Antibiotic resistance in nonenterococcal lactic acid bacteria and bifidobacteria. *Food microbiology*. 24(6): 6–15.
- Ariff, A. B. (1993). The Influence of Mode Operation on the Production of Glucoamylase by *Aspergillus owamori*. PhD Thesis, Univeriti Putra Malaysia.
- Ayadi-Zouari, D., Kammoun, R., Jemli, S., Chouayekh, H. and Bejar, S. (2012). Secretion of cyclodextrin glucanotransferase in *E. coli* using *Bacillus* subtilis lipase signal peptide and optimization of culture medium. Indian journal of experimental biology. 50(1): 72–9.
- Bahey-El-Din, M. and Gahan, C. (2010). *Lactococcus lactis*: from the dairy industry to antigen and therapeutic protein delivery. *Discovery medicine*. 9: 455–461.
- Baş, D. and Boyacı, İ. H. (2007). Modeling and optimization II: Comparison of estimation capabilities of response surface methodology with artificial neural networks in a biochemical reaction. *Journal of Food Engineering*. 78(3): 846–854.
- Benjamin, K., Emmanuel, A., David, A. and Benjamin, Y. (2008). Genetic Algorithms Using for a Batch Fermentation Process Identification. *Journal* of Applied Sciences. 8(12): 2272–2278.
- Birol, G., Undey, C. and Cinar, A. (2002). A modular simulation package for fedbatch fermentation: penicillin production. *Computers & chemical engineering*. 26: 1553–1565.
- Blanco, K., Lima, C. de and Monti, R. (2012). *Bacillus lehensis*—an alkali-tolerant bacterium isolated from cassava starch wastewater: optimization of parameters for cyclodextrin glycosyltransferase production. *Annals of Microbiology*. 62: 329–337.
- Bolotin, A., Wincker, P., Mauger, S., Jaillon, O., Malarme, K., Weissenbach, J., Ehrlich, S. D. and Sorokin, A. (2001). The complete genome sequence of the lactic acid bacterium *Lactococcus lactis* ssp. *lactis* IL1403. *Genome research*. 11(5): 731–53.

- Bouguettoucha, A., Balannec, B. and Amrane, A. (2011). Unstructured Models for Lactic Acid Fermentation- A Review. *Food Technology and Biotechnology*. 49(1): 3–12.
- Burhan, N., Sapundzhiev, T. and Beschkov, V. (2005). Mathematical modelling of cyclodextrin-glucanotransferase production by batch cultivation. *Biochemical Engineering Journal*. 24(1): 73–77.
- Cheigh, C. I., Choi, H. J., Park, H., Kim, S. B., Kook, M. C., Kim, T. S., Hwang, J. K. and Pyun, Y. R. (2002). Influence of growth conditions on the production of a nisin-like bacteriocin by *Lactococcus lactis* subsp. *lactis* A164 isolated from kimchi. *Journal of Biotechnology*. 95(3): 225–35.
- Choonia, H. S. and Lele, S. S. (2013). Kinetic modeling and implementation of superior process strategies for β -galactosidase production during submerged fermentation in a stirred tank bioreactor. *Biochemical Engineering Journal*. 77: 49–57.
- Cotter, P. and Hill, C. (2003). Surviving the acid test: responses of gram-positive bacteria to low pH. *Microbiology and Molecular Biology Reviews*. 67(3): 429–453.
- Dalié, D., Deschamps, A. and Richard-Forget, F. (2010). Lactic acid bacteria– Potential for control of mould growth and mycotoxins: A review. *Food Control*. 21: 370–380.
- Deegan, L., Cotter, P., Hill, C. and Ross, P. (2006). Bacteriocins: biological tools for bio-preservation and shelf-life extension. *International Dairy Journal*. 16(9): 1058-1071.
- Desai, K. M., Akolkar, S. K., Badhe, Y. P., Tambe, S. S. and Lele, S. S. (2006). Optimization of fermentation media for exopolysaccharide production from *Lactobacillus plantarum* using artificial intelligence-based techniques. *Process Biochemistry*. 41(8): 1842–1848.
- Dodziuk, H. (2006a). Applications Other Than in the Pharmaceutical Industry. In H. Dodziuk (Ed.), *Cyclodextrins and Their Complexes* (pp. 450-473). KGaA, Weinheim: WILEY-VCH Verlag GmbH &Co.
- Dodziuk, H. (2006b). Molecules with Holes-Cyclodextrins. In H. Dodziuk (Ed.), *Cyclodextrins and Their Complexes* (pp. 1-30). KGaA, Weinheim.
- Del Valle, E. M. M. (2004). Cyclodextrins and their uses: a review. *Process Biochemistry*. 39(9): 1033–1046.
- Falch, E. A. (1991). Production and application. *Biotechnology Advance*. 9: 643–658.
- Farliahati, M., Mohamed, M., Rosfarizan, M., Puspaningsih, N. and Ariff, A. (2009). Kinetics of xylanase fermentation by recombinant *Escherichia coli* DH5α in shake flask culture. *American Journal of Biochemistry and Bioechnology*. 5(3): 110–118.

- Feng, T., Zhuang, H. and Ran, Y. (2011). The Application of Cyclodextrin Glycosyltransferase in Biological Science. *Journal of Bioequivalence & Bioavailability*. 03(09): 202–206.
- Freitas, T. L. De, Monti, R. and Contiero, J. (2004). Production of CGTase by a *Bacillus alkalophilic* CGII strain isolated from wastewater of a manioc flour industry. *Brazilian Journal of Microbiology*. 35: 255–260.
- Frelet-barrand, A., Boutigny, S., Kunji, E. R. S. and Rolland, N. (2010). Heterologous Expression of Membrane Proteins. In I. Mus-Veteau (Ed.), (Vol. 601, pp. 67–85). Totowa, NJ: Humana Press.
- Gastón, J. A. R., Szerman, N., Costa, H., Krymkiewicz, N. and Ferrarotti, S. A. (2009). Cyclodextrin glycosyltransferase from *Bacillus circulans* DF 9R: Activity and kinetic studies. *Enzyme and Microbial Technology*. 45(1): 36– 41.
- Gawande, B. and Patkar, A. (2001). Purification and properties of a novel raw starch degrading-cyclodextrin glycosyltransferase from *Klebsiella pneumoniae* AS-22. *Enzyme and Microbial Technology*. 28(9-10): 735–743.
- Gawande, B., Singh, R. and Chauhan, A. (1998). Optimization of cyclomaltodextrin glucanotransferase production from *Bacillus firmus*. *Enzyme and Microbial Technology*. 22(4): 288-291.
- Gawande, B., Sonawane, A., Vogdand, V. and Patkar, A. (2003). Optimization of Cyclodextrin Glycosyltransferase Production from *Klebsiella pneumoniae* AS- 22 in Batch, Fed- Batch, and Continuous Cultures. *Biotechnolog Progress*. 19(6): 1679–1702.
- Ghosh, S. and Chakraborty, R. (2012). Study on fermentation conditions of palm juice vinegar by response surface methodology and development of a kinetic model. *Brazilian Journal of Chemical Engineering*. 29(2).
- Grothe, E., Moo-Young, M. and Chisti, Y. (1999). Fermentation optimization for the production of poly (NL -hydroxybutyric acid) microbial thermoplastic. *Enzyme and Microbial Technology*. 25: 132–141.
- Guo, W., Zhang, Y., Lu, J., Jiang, L., Teng, L. and Wang, Y. (2010). Optimization of fermentation medium for nisin production from *Lactococcus lactis* subsp. *lactis* using response surface methodology (RSM) combined with artificial neural network-genetic algorithm (ANN-GA). *African Journal of Biotechnology*. 9(38): 6264–6272.
- Hatti-Kaul, R. (2002). Enzyme production. In *Encyclopedia of Life Support Systems* (Vol. V). UNESCO.
- Higuti, I. H., Anunciação, P. and José, A. (2004). Studies on Alkalophilic CGTase-Producing Bacteria and Effect of Starch on Cyclodextrin-Glycosyltransferase Activity. *Brazilian Archives of Biology and Techonology*. 47: 135–138.

- Hong, P., Illias, R. and Mau, K. (2012). Domain replacement to elucidate the role of B domain in CGTase thermostability and activity. *Process Biochemistry*. 47(12): 2123–2130.
- Hoopen, Hjg., Gulik, Wm., Schlatmann, J., Moreno, P., Vinke, J., Heijnen, J. and Verpoorte, R. (1994). Ajmalicine production by cell cultures of Catharanthus roseus: from shake flask to bioreactor. Plant Cell, Tissue and Organ Culture. 38: 85–91.
- Ibrahim, H. M., Yusoff, W. M. W., Hamid, A. A., Illias, R. M., Hassan, O. and Omar, O. (2005). Optimization of medium for the production of βcyclodextrin glucanotransferase using Central Composite Design (CCD). *Process Biochemistry*. 40(2): 753–758.
- Illias, R., Tien, S., Rahman, R., Rashid, N., Yusoff, W., Hamid, A., Hassan, O. and Kamaruddin, K. (2003). Application of factorial design to study the effects of temperature, initial pH and agitation on the production of cyclodextrin glucanotransferase from alkalophilic *Bacillus* sp. G1. *Science Asia*. 29: 135–140.
- Kaneko, T., Kato, T., Nakamura, N. and Horikoshi, K. (1987). Spectrophotometric determination of cyclization activity of β-cyclodextrin-forming cyclomaltodextrin glucanotransferase. *Journal of Japan Society of Starch Sciences.* 34: 45–48.
- Keller, A. and Gerhardt, P. (2004). Continuous lactic acid fermentation of whey to produce a ruminant feed supplement high in crude protein. *Biotechnology* and Bioengineering. 17(7): 997–1018.
- Khoramnia, A., Lai, O. M., Ebrahimpour, A., Tanduba, C. J., Voon, T. S. and Mukhlis, S. (2010). Thermostable lipase from a newly isolated *Staphylococcus* xylosus strain; process optimization and characterization using RSM and ANN. *Electronic Journal of Biotechnology*. 13(5).
- Khurana, S., Kapoor, M., Gupta, S. and Kuhad, R. C. (2007). Statistical optimization of alkaline xylanase production from Streptomyces violaceoruber under submerged fermentation using response surface methodology. *Indian journal of microbiology*. 47(2): 144–52.
- Khuri, A. and Mukhopadhyay, S. (2010). Response surface methodology. *Wiley Interdisciplinary Reviews: Computational Statistics*. 2(2): 128–149.
- Kim, P., Hassan, O., Ahmad, A., Muhammad, N. and Illias, R. (2007). Excretory over-expression of Bacillus sp . G1 cyclodextrin glucanotransferase (CGTase) in *Escherichia coli*: Optimization of the cultivation conditions by response surface methodology. *Enzyme and Microbial Technology*. 40: 1256–1263.
- Kirk, O., Borchert, T. V. and Fuglsang, C. C. (2002). Industrial enzyme applications. *Current Opinion in Biotechnology*. 13(4): 345–351.

- Kojić, M., Lozo, J., Begović, J. and Topisirović, L. (2007). Characterization of *lactococci* isolated from homemade kefir. *Archives of Biological Science*. 59(1): 13–22.
- Kono, T. and Asai, T. (2004). Kinetics of fermentation processes. *Biotechnology and Bioengineering*. 11(3): 293–321.
- Kuo, C. C., Lin, C. A., Chen, J. Y., Lin, M. T. and Duan, K. J. (2009). Production of cyclodextrin glucanotransferase from an alkalophilic *Bacillus* sp. by pH stat fed-batch fermentation. *Biotechnology Letters*. 31:1723–1727.
- Li, S., Yang, X., Yang, S., Zhu, M. andWang, X. (2012). Technology Prospecting on Enzymes : Application, Marketing and Engineering Abstract : Enzymes are protein molecules functioning as specialized catalysts for chemical reactions. They have contributed greatly to the traditional and modern chemical industr. *Computational and Structural Biotechnology Journal*. 2(3).
- Ling, L. (2004). *Pilot-Scale Production of Lactobacillus rhamnosus ATCC 7469*, PhD Thesis, Universiti Putra Malaysia.
- Liu, J., Weng, L., Zhang, Q., Xu, H. and Ji, L. (2003). Short communication A mathematical model for gluconic acid fermentation by *Aspergillus niger*. *Biochemical Engineering Journal*. 14: 137–141.
- Lo, P., Tan, C., Hassan, O., Ahmad, A., Mahadi, N. and Illias, R. (2009). Improvement of excretory overexpression for *Bacillus* sp. G1 cyclodextrin glucanotransferase (CGTase) in recombinant *Escherichia coli* through medium optimization. *Biotechnology*. 8(2): 184–193.
- Loftsson, T. and Duchêne, D. (2007). Cyclodextrins and their pharmaceutical applications. *International journal of pharmaceutics*. 329(1-2): 1–11.
- Low, K. O., Mahadi, N. M., Rahim, R. A., Rabu, A., Abu Bakar, F. D., Murad, A. M. A. and Illias, R. M. (2011). An effective extracellular protein secretion by an ABC transporter system in *Escherichia coli*: statistical modeling and optimization of cyclodextrin glucanotransferase secretory production. *Journal of industrial microbiology & biotechnology*. 38(9): 1587–97.
- Luedeking, R. and Piret, E. (2004). A kinetic study of the lactic acid fermentation. Batch process at controlled pH. *Journal of Biochemical and Molecular Toxicology*. 1(4): 392–412.
- Mahat, M., Illias, R. and Rahman, R. (2004). Production of cyclodextrin glucanotransferase (CGTase) from alkalophilic *Bacillus* sp. TS1-1: media optimization using experimental design. *Enzyme and Microbial Technology*. 35: 467–473.
- Makarova, K. and Slesarev, A., *et al.* (2006). Comparative genomics of the lactic acid bacteria. *Proceedings of the Natural Academic Science of United States of America.* 103(42): 15611-15616.

- Mall, P., Mohanty, B. K. and Patankar, D. B. (2010). Physiochemical Parameters Optimization for Enhanced Nisin Production by *Lactococcus lactis* (MTCC 440). *Brazilian Archives of Biology and Technology*. 53(1): 203– 209.
- Manabe, K., Kageyama, Y., Morimoto, T., Shimizu, E., Takahashi, H., Kanaya, S., Ozaki, K. and Ogasawara, N. (2013). Improved production of secreted heterologous enzyme in *Bacillus subtilis* strain MGB874 via modification of glutamate metabolism and growth conditions. *Microbial cell factories*. 12(1): 18.
- Mander, P., Choi, Y. H., Seong, J. H., Na, B. H., Cho, S. S., Lee, H. J. and Yoo, J. C. (2013). Statistical optimization of a multivariate fermentation process for enhancing antibiotic activity of Streptomyces sp. CS392. Archives of pharmacal research, (2013). doi:10.1007/s12272-013-0140-4
- Mata-alvarez, J. and Mitchell, D. A. (n.d.). Mathematical modeling in biotechnology. In *Encyclopedia of Life Support Systems* (Vol. II).
- Matte, C. R., Nunes, M. R., Benvenutti, E. V., Schöffer, J. D. N., Ayub, M. A. Z. and Hertz, P. F. (2012). Characterization of cyclodextrin glycosyltransferase immobilized on silica microspheres via aminopropyltrimethoxysilane as a "spacer arm." *Journal of Molecular Catalysis B: Enzymatic.* 78: 51–56.
- Menocci, V., Goulart, A., Adalberto, P., Tavano, O., Marques, D., Contiero, J. and Monti, R. (2008). Cyclodextrin glycosyltransferase production by new bacillus sp. strains isolated from brazilian soil. Brazilian Journal of Microbiology. 39: 682–688.
- Mierau, I., Olieman, K., Mond, J. and Smid, E. J. (2005). Optimization of the *Lactococcus lactis* nisin-controlled gene expression system NICE for industrial applications. *Microbial cell factories*, 4(16).
- Miro, A. M., Krasowska, A., Murzyn, A. and Marcin, Ł. (2012). Production of the Bacillus licheniformis SubC protease using *Lactococcus lactis* NICE expression system. *SpringerPlus*. 1(54): 1–10.
- Monod, J. (1942). Researches sur les croissances des cultures bacteriennes. 2nd edition. Hermann and Cie, Paris.
- More, S. S., Niraja, R., Evelyn, C., Byadgi, A. M., Shwetha, V. and Mangaraj, S. Das. (2012). Isolation, Purifi cation and Biochemical Characterization of CGTase from *Bacillus halodurans*. Croatian Journal of Food Technology, Biotechnology and Nutrition. 7(1-2): 90–97.
- Muthulakshmi, C., Gomathi, D., Kumar, D. G., Ravikumar, G., Kalaiselvi, M. and Uma, C. (2011). Production, Purification and Characterization of Protease by Aspergillus flavus under Solid State Fermentation. Jordan Journal of Biological Sciences. 4(3): 137–148.

- Nagata, Y. and Chu, K. H. (2003). Optimization of a fermentation medium using neural networks and genetic algorithms. *Biotechnology letters*. 25(21): 1837–42.
- Nakamura, L. K. (1981). *Lactobacillus amylovorus*, a new starch-hydrolyzing species from cattle waste-corn fermentations. *International Journal of Systematic Bacteriology*. 31(1): 56-63.
- Nelofer, R., Ramanan, R. N., Rahman, R. N. Z. R. A., Basri, M. and Ariff, A. B. (2012). Comparison of the estimation capabilities of response surface methodology and artificial neural network for the optimization of recombinant lipase production by *E. coli* BL21. *Journal of industrial microbiology & biotechnology*. 39(2): 243–54.
- Noreen, N., Hooi, W. Y., Baradaran, A., Rosfarizan, M., Sieo, C. C., Rosli, M. I., Yusoff, K. and Raha, A. R. (2011). *Lactococcus lactis* M4, a potential host for the expression of heterologous proteins. *Microbial cell factories*. 10(1): 28.
- Olempska-Beer, Z. S., Merker, R. I., Ditto, M. D. and DiNovi, M. J. (2006). Foodprocessing enzymes from recombinant microorganisms--a review. *Regulatory toxicology and pharmacology : RTP.* 45(2): 144–158.
- Oliveira, A. P., Nielsen, J. and Förster, J. (2005). Modeling *Lactococcus lactis* using a genome-scale flux model. *BMC microbiology*. 5: 39.
- Ong, R. M., Goh, K. M., Mahadi, N. M., Hassan, O., Rahman, R. N. Z. R. A. and Illias, R. M. (2008). Cloning, extracellular expression and characterization of a predominant beta-CGTase from *Bacillus* sp. G1 in *E. coli*. *Journal of industrial microbiology & biotechnology*. 35(12): 1705–14.
- Orla-Jensen, S. (1919). The lactic acid bacteria.
- Papagianni, M. (2012). Metabolic engineering of lactic acid bacteria for the production of industrially important compounds. *Computational and Structural Biotechnology Journal*.
- Pearl, R. and Reed, L. (1920). On the rate of growth of the population of the United States since 1790 and its mathematical representation. *Proceeding of the National Academy of Sciences of the United States of America*. 6(6): 275– 288.
- Pinto, F., Flôres, S., Ayub, M. and Hertz, P. (2007). Production of cyclodextrin glycosyltransferase by alkaliphilic Bacillus circulans in submerged and solid-state cultivation. *Bioprocess and biosystems Enginnering*. 30: 377–382.
- Prakasham, R. S., Rao, R. S., Rao, C. S. and Sarma, P. N. (2005). Cyclodextrin glycosyl transferases from *Bacillus circulance* and *Bacillus* sp. *Indian Journal of Biotechnology*. 4: 347–352.
- Rahman, R., Illias, R. and Nawawi, M. (2004). of growth medium for the production of cyclodextrin glucanotransferase from *Bacillus*

stearothermophilus HR1 using response surface methodology. Process Biochemistry. 39(12): 2053-2060.

- Ramli, N., Abd-Aziz, S., Hassan, M., Alitheen, N. and Kamaruddin, K. (2010). Potential cyclodextrin glycosyltransferase producer from locally isolated bacteria. *African Journal of Biotechnology*. 9(43): 7317–7321.
- Rasheed, A., Kumar, A., & Sravanthi, V. (2008). Cyclodextrins as Drug Carrier Molecule: A Review. *Scientia Pharmaceutica* .76(4): 567–598.
- Ravinder, K., Prabhakar, T. and Bhavanidevi, R. (2012). Optimization of process parameters for the production of cyclodextrin glycosyltransferase by newly isolated *bacillus* sp. tpr71h by conventional. *International journal of advanced biotechnology and research*. 3(2): 578–584.
- Roels, J. (2007). Mathematical models and the design of biochemical reactors. Journal of Chemical Technology and Biotechnology. 32(1): 59–72.
- Rosfarizan, M. (2000). Kinetics, Modelling and Scaling-Up of Kojic Acid Fermentation by Aspergillus flavus 44-1 Using Different Carbon Sources, PhD Thesis, Universiti Putra Malaysia.
- Rosso, A. and Ferrarotti, S. (2002). Optimisation of batch culture conditions for cyclodextrin glucanotransferase production from *Bacillus circulans* DF 9R. *Microbial Cell Factories.* 9: 1-9.
- Salvetti, E., Torriani, S., and Felis, G. (2012). The Genus Lactobacillus: A Taxonomic Update. Probiotics and Antimicrobial Proteins. 4(4): 217-226.
- Samaržija, D., Antunac, N. and Havranek, J. L. (2001). Taxonomy, physiology and growth of *Lactococcus lactis*: a review. *Mljekarstvo*. 51(1): 35–48.
- Sanchez, S. and Demain, A. L. (2011). Enzymes and Bioconversions of Industrial, Pharmaceutical, and Biotechnological Abstract : Organic Process Research & Development. 15(1): 224–230.
- Sarrouh, B., Santos, T. M., Miyoshi, A., Dias, R. and Azevedo, V. (2012). Up-To-Date Insight on Industrial Enzymes Applications and Global Market. J *Bioprocess Biotechniq*. doi:10.4172/2155-9821.S4-002.
- Schleifer, K. and Ludwig, W. (1995). Phylogenetic relationships of lactic acid bacteria. *The genera of lactic acid bacteria*. 2: 7–18.
- Scott, C. (2004). Microbial Fermentation The Oldest Form of Biotechnology. *Bioprocess International*. 2: 8–20.
- Sharma, S., Garg, A., & Singh, G. (2010). Optimization of fermentation conditions for bacteriocin production by *Lactococcus lactis* CCSULAC1 on modified MRS medium. *Intertnational Journal of Dairy Science*. 5(1): 1–9.
- Shene, C., Mir, N., Andrews, B. and Asenjo, J. (2000). Effect of the growth conditions on the synthesis of a recombinant beta-1,4-endoglucanase in

continuous and fed-batch culture. *Enzyme and Microbial Technology*. 27(3-5): 248–253.

- Sian, H. K., Said, M., Hassan, O., Kamaruddin, K., Ismail, a. F., Rahman, R. A., Mahmood N. A. N. and Illias, R. M. (2005). Purification and characterization of cyclodextrin glucanotransferase from alkalophilic *Bacillus* sp. G1. *Process Biochemistry*. 40(3-4): 1101–1111.
- Silva, S. de, Petterson, B., Muro, M. de and Priest, F. (1998). A DNA Probe for the Detection and Identification of *Bacillus sporothermodurans* Using the 16S-23S rDNA Spacer Region and Phylogenetic Analysis of Some Field Isolates of *Bacillus* which form Highly Heat Resistant Spores. *Systematic* and Applied Microbiology. 21(3): 398–407.
- Sinclair, C. and Cantero, D. (1990). Fermentation modeling. In E. B. McNeil & L. M. Harvey (Eds.), *Fermentation: a practical approach* (pp. 65–112). Oxford: Oxford University Press.
- Sivakumar, N. and Shakilabanu, S. (2013). Original Research Article Production of cyclodextrin glycosyl transferase by Bacillus megaterium. *International Journal of Current Microbiology and Applied Sciences*. 2(7): 44–55.
- Song, H., Jang, S. H., Park, J. M. and Lee, S. Y. (2008). Modeling of batch fermentation kinetics for succinic acid production by *Mannheimia* succiniciproducens. Biochemical Engineering Journal. 40(1): 107–115.
- Souza, R. D., Pandeya, D. R. and Hong, S. (2012). Review: *Lactococcus Lactis* : An efficient Gram positive cell factory for the production and secretion of recombinant protein. *Biomedical Research*. 23(1): 1–7.
- Stanbury, P. (1988). Fermentation technology. In E. B. McNeil & L. M. Harrey (Eds.), *Extraction*. Wiley InterScience.
- Subramaniam, M., Baradaran, A., Illias, RM., Rosfarizan, M., Khatijah, Y. and Raha, A. (2012). Effect of Signal Peptides on the Secretion of β-Cyclodextrin Glucanotransferase in *Lactococcus lactis* NZ9000. *Journal of Molecular Microbiology and Biotechnology*. 22: 361–372.
- Surampalli, R. and Tyagi, R. (2004). *Advances in Water and Wastewater Treatment*. Reston, Virginia: American Society of Civil Engineering.
- Szente, L., Szejtli, J. and Szeman, J. (1993). Fatty acid-cyclodextrin complexes : properties and applications. *Journal of Inclusion Phenomena and Molecular Recognition in Chemistry*. 16: 339–354.
- Tan, J., Ramanan, R., Ling, T., Shuhaimi, M. and Ariff, A. (2011). Comparison of predictive papabilities of Response Surface Methodology and Artificial Neural Network for optimization of periplasmic interferon-α2b production by recombinant *Escherichia coli*. *Minerva Biotecnologica*. 23: 1–2.
- Thatai, A., Kumar, M. and K.J. Mukherjee. (1999). A single step purification process from *Bacillus* sp. isolated from soil. *Preparative Biochemistry and Biotechnology*. 29: 35 47.

- Teuber, M. and Geis, A. (2006). The genus *Lactococcus*. In *The Prokaryotes* (3rd. Ed., Vol. 4, pp. 205–228).
- Todar, K. (2006). Todar's online textbook of bacteriology (p. 1421). Madison.
- Tsao, C. C. (2008). Comparison between response surface method- ology and radial basis function network for core-center drill in drilling composite materials. *International Journal of Advanced Manufacturing Technology*. 37:1061–1068.
- Uekama, K., Hirayama, F., & Arima, H. (2006). Pharmaceuticals Applications of Cyclodextrins and Their Derivatives. In H. Dodziuk (Ed.), Cyclodextrins and Their Complexes (pp. 381-422). KGaA, Weinheim: WILEY-VCH Verlag GmbH &Co.
- Vassileva, A., Burhan, N. and Beschkov, V., Spasova, D., Radoevska, S., Ivanova, V. and Tonkova, A. (2003). Cyclodextrin glucanotransferase production by free and agar gel immobilized cells of *Bacillus circulans* ATCC 21783. *Process Chemistry*. 38(11): 1585-1591.
- Verhulst, P.F. (1838). Notice sur la loi que la population suit dans son accroissement. *Corr. Mathematica et Physica*. 10: 113–121.
- Villatoro-Hernández, J., Kuipers, O. P., Saucedo-cárdenas, O. and Montes-de-ocaluna, R. (2012). Recombinant Gene Expression. In A. Lorence (Ed.), (3rd Ed., Vol. 824, pp. 155–165). New York, NY: Springer New York.
- Wachenheim, D. E., Patterson, J. A. and Ladisch, M. R. (2003). Analysis of the logistic function model: derivation and applications specific to batch cultured microorganisms. *Bioresource Technology*. 86: 157–164.
- Walker, J. M. and Ditor, S. E. E. (2005). *Microbial Processes and Products*. (B. Jose-Luis, Ed.). Totowa, New Jeresey: Humana Press Inc.
- Wang, F., Du, G., Li, Y. and Chen, J. (2005). Optimization of Cultivation Conditions for the Production of γ-Cyclodextrin Glucanotransferase by *Bacillus macorous. Food Biotechnology*. 18(2): 251–264.
- Wouters, J. A., Kamphuis, H. H., Kuipers, O. P. and Vos, W. M. De. (2000). Changes in Glycolytic Activity of *Lactococcus lactis* Induced by Low Temperature. *Applied and Environmental Microbiology*. 66(9): 3686–3691.
- Xu, L.-J., Liu, Y.-S., Zhou, L.-G. and Wu, J.-Y. (2011). Modeling of *Fusarium redolens* Dzf2 mycelial growth kinetics and optimal fed-batch fermentation for beauvericin production. *Journal of industrial microbiology* & *biotechnology*. 38(9): 1187–92.
- Yap, P., Ariff, A., Woo, K. and Hii, S. (2010). Production of cyclodextrin glycosyltransferase (CGTase) by *Bacillus lehensis* S8 using sago starch as carbon source. *Journal of Biological Sci*ences. 10(7): 676–681.
- Youssefi, S. H., Emam-Djomeh, Z. and Mousavi, S. M. (2009.) Comparison of artificial neural network (ANN) and response surface methodology (RSM)

in the prediction of quality parameters of spray- dried pomegranate juice. *Dry Technology*. 27:910–917.

- Zain, W., Illias, R. and Salleh, M. (2007). Production of cyclodextrin glucanotransferase from alkalophilic *Bacillus* sp. TS1-1: Optimization of carbon and nitrogen concentration in the feed medium using central composite design. *Biochemical Engineering Journal*. 33(1): 26–33.
- Zhang, G., Mills, D. A. and Block, D. E. (2009). Development of chemically defined media supporting high-cell-density. *Applied and Environmental Microbiology*. 75(4): 1080–1087.
- Zhen-Gui, Z., Di, L. and Zhi-Bi, H. (1998). Comparison of cell growth and alkaloid production of *Catharanthus roseus* cells cultured in shake-flask and bioreactor. *Acta Botanica Sincia*. 40(1): 51–55.
- Zhu, L.-W., Wang, C.-C., Liu, R.-S., Li, H.-M., Wan, D.-J. and Tang, Y.-J. (2012). Actinobacillus succinogenes ATCC 55618 fermentation medium optimization for the production of succinic acid by response surface methodology. Journal of biomedicine & biotechnology, 2012, 626137. doi:10.1155/2012/626137.