



**UNIVERSITI PUTRA MALAYSIA**

***ENHANCING RECOMBINANT T1 Lipase PRODUCTION IN  
Pichia guilliermondii***

**ABU MARY LADIDI**

**FBSB 2017 19**



**ENHANCING RECOMBINANT T1 *Lipase* PRODUCTION IN  
*Pichia guilliermondii***

**By**

**ABU MARY LADIDI**

**Thesis Submitted to the School of Graduate Studies, Universiti Putra Malaysia,  
in Fulfillments of the Requirements for the Degree of Master of Science**

**July 2017**

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## **DEDICATION**

I dedicate this research work to God Almighty the giver and sustainer of life and to my family who have being by my side.



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Abstract of thesis presented to the Senate of Universiti Putra Malaysia in fulfilment of the requirement for the Degree of Master of Science

**ENHANCING RECOMBINANT T1 *Lipase* PRODUCTION IN  
*Pichia guilliermondii***

By

**ABU MARY LADIDI**

**July 2017**

**Chairman : Professor Abu Bakar Salleh, PhD**  
**Faculty : Biotechnology and Biomolecular Sciences**

Lipases are hydrolytic enzymes, ranked as third most relevant industrial enzymes with 5 % share in global enzyme market. Thermostable T1 lipase from *Geobacillus zalihae* was previously expressed under the regulatory control of alcohol oxidase promoter 1 (AOXp 1) in the methylotrophic yeast *Pichia guilliermondii*, isolated from spoiled orange. Methanol was found to be a vital compound to induce the promoter activity in *Pichia pastoris*. In this study *P. guilliermondii* has shown the potential to express the recombinant lipase without methanol under the regulation of AOXp 1. This study sought to optimise medium conditions of thermostable lipase with and without methanol as inducer. The expression of T1 lipase without methanol was expected to reduce cost and toxicity effect of methanol.

Buffered and non-buffered media compositions were studied for T1 lipase production, the media were first, supplemented with methanol then without methanol. Buffer complex methanol medium was observed to be optimum for T1 lipase production with a 3-fold increase over non-buffered methanol medium. *One-factor-at-a-time*, conventional method of optimisation was used to identify significant data range for medium parameters. Using the observed data range, eight parameters which includes temperature, pH, inoculum size, biomass concentration, incubation time, shaking speed, culture volume and methanol concentration, were screened for lipase production in methanol medium using Plackett-Burman Design. Temperature, inoculum size, culture volume and incubation time, were observed to exert significant effect on lipase production. These parameters were optimised using Box-Behnken Design of Response Surface Methodology. Optimum levels of these parameters were predicted at temperature 34 °C, culture volume 190 mL, inoculum size 4 v/v and incubation time 24 h with an experimented lipase activity of 9.26 U/mL. Over 2-fold increase before optimization in methanol medium was observed and 6-fold increase over previous research work.

On the other hand, six parameters which include temperature, pH, inoculum size, incubation time, shaking speed and culture volume were screened for T1 lipase production in medium without methanol. Three parameters were observed to have significant effect on lipase production, then, these parameters were further optimised using Box-Behnken Design and their optimum levels were achieved at pH of 6, inoculum size 2 v/v and incubation time of 24 h as was predicted and the experimented lipase activity of 2.012 U/mL was observed. This result gave 4-fold increase over lipase production before optimisation in medium without methanol. Recombinant T1 lipase production was further scaled up to 3 L in the bioreactor for 128 h in methanol medium and lipase activity observed was 12 U/mL at 120 h incubation time. Meanwhile, scale up in medium without methanol yielded a lipase activity of 3.5 U/mL at 118 h incubation time. The result has proven that, the time taken for optimum lipase production without methanol was faster from the production with methanol in the bioreactor.

In conclusion, temperature, pH, inoculum size, incubation time and culture volume for recombinant T1 lipase production were optimised in shake-flask and applied to the bioreactor level. Lipase was also produced in medium without methanol both at the shake flask level and the bioreactor. Thus, medium parameters optimisation has proven to be very useful in enhancing thermostable lipase production in recombinant *P. guilliermondii*. Although T1 lipase has been expressed in medium without methanol but it is lower compared to medium with methanol.

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

**PENINGKATAN PENGELUARAN *Lipase* RENTAN HABA  
REKOMBINAN DALAM *Pichia guilliermondii* STRAIN SO**

Oleh

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Lipase adalah enzim hidrolitik, tersenarai sebagai enzim ketiga paling relevan dalam industri dengan berkongsi sebanyak 5% dalam pasaran enzim global. Lipase T1 rentan panas dari *Geobacillus zalihae* sebelum ini telah dihasilkan di bawah kawalan penganjur alkohol oksidase 1 (AOXp 1) di dalam *Pichia guilliermondii* yang merupakan yis metanotropik yang mana telah dipencilkan dari buah limau busuk. Metanol merupakan komponen penting untuk mengaruh aktiviti penganjur AOXp 1 dalam *Pichia pastoris*. Dalam kajian ini *P. guilliermondii* telah menunjukkan potensi untuk menghasilkan lipase rekombinan tanpa metanol di bawah aruhan AOXp 1. Kajian ini bertujuan untuk mengoptimumkan penghasilan lipase T1 rentan haba dengan kehadiran metanol dan tanpa methanol sebagai pengaruh. Penghasilan T1 lipase rekombinan tanpa aruhan metanol diharapkan akan mengurang kos dan kesan tosik daripada metanol.

Komposisi media yang berpenimbal dan tanpa penimbal dikaji untuk penghasilan T1 lipase. Pada permulaannya, media tersebut disertakan dengan metanol, kemudian tanpa metanol. Media berpenimbal kompleks dengan metanol didapati telah menghasilkan T1 lipase yang optimum dengan peningkatan sebanyak 3 kali ganda berbanding media yang tidak berpenimbal dan bermetanol. Satu-faktor-pada-satu-masa merupakan kaedah perngoptimuman konvensional telah digunakan untuk mengenalpasti julat data penting untuk parameter dalam media. Berdasarkan julat data yang diperolehi, lapan parameter seperti suhu, pH, saiz inokulasi, kepekatan biomas, masa pengeraman, kelajuan penggolak, isipadu kultur dan kepekatan metanol telah dicerap untuk pengeluaran lipase dalam medium metanol menggunakan Reka Bentuk Plackett-Burman. Suhu, saiz inokulasi, isipadu kultur dan masa pengeraman didapati memberikan kesan yang penting kepada penghasilan lipase. Parameter-parameter ini dioptimumkan menggunakan yang reka bentuk Box-Behnken daripada kaedah respon permukaan. Tahap optimum parameter ini telah diramalkan pada suhu 34 °C,

kandungan cecair 190 mL, saiz inokulum 4 v/v dan masa inkubasi 24 jam dengan hasil lipase 9.26 U/mL dengan eksperimen. Lebih 2 kali ganda peningkatan diperhatikan sebelum pengoptimuman dalam medium metanol dan 6 kali ganda peningkatan ke atas kerja penyelidikan yang sebelumnya.

Sebaliknya, enam parameter termasuk suhu, pH, saiz inokulasi, masa pengeraman, kelajuan penggolak dan isipadu kultur telah disaring untuk penghasilan lipase dalam media tanpa metanol. Tiga parameter telah diperhatikan mempunyai kesan ketara dalam pengeluaran lipase maka, parameter ini seterusnya telah dioptimumkan menggunakan Reka bentuk Box–Behnken dan tahap optimum mereka dicapai pada pH 6 saiz inokulum 2 v/v dan masa inkubasi 24 jam seperti yang dijangkakan dan hasil lipase eksperimen 2.01 U/mL direkodkan. Keputusan ini memberikan kenaikan 4 kali ganda ke atas pengeluaran lipase sebelum pengoptimuman dalam media tanpa metanol. Pengeluaran T1 lipase rekombinan bertambah pada skala besar sehingga 3 L bioreaktor pada 128 jam di dalam media metanol dan penghasilan lipase yang diperolehi adalah 12 U/mL pada 120 h masa inkubasi. Sementara itu, media tanpa methanol pada skala besar menghasilkan aktiviti lipase sebanyak 3.5 U/mL ketika masa inkubasi 118 jam. Hasilnya telah terbukti bahawa, masa yang diambil untuk pengeluaran optimum lipase tanpa methanol adalah lebih cepat daripada pengeluaran dengan methanol di dalam bioreaktor.

Kesimpulannya, parameter media untuk pengeluaran lipase T1 rekombinan dioptimumkan dalam kelalang pengoncang dan digunakan ke peringkat bioreaktor. Lipase juga dihasilkan dalam media tanpa metanol di kedua-dua peringkat kelalang pengoncang dan di dalam bioreaktor. Oleh itu, parameter media yang optimum telah terbukti menjadi sangat berguna dalam meningkatkan pengeluaran lipase rekombinan rentan haba *P. guilliermondii*. Walaupun enzim lipase T1 telah berjaya dihasilkan dalam media tanpa metanol namun penghasilannya adalah ia lebih rendah berbanding media dengan metanol.



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I certify that a Thesis Examination Committee has met on 28 July 2017 to conduct the final examination of Abu Mary Ladidi on his thesis entitled "Enhancing Recombinant T1 *Lipase* Production in *Pichia guilliermondii*" in accordance with the Universities 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 Master of Science.

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## LIST OF ABBREVIATIONS

%	Percentage
YPD	Yeast extract Peptone Dextrose
OD <sub>600</sub>	Optical Density at 600 nanometer
BMMY	Buffered complex - methanol medium yeast extract
BMY	Buffered complex - 1 medium yeast extract
BMGY	Buffered complex – glycerol medium yeast extract
YPT	Yeast extract Peptone Tryptic soy
YPTG	Yeast extract Peptone Tryptic soy Glycerol
YPTM	Yeast extract Peptone Tryptic soy Methanol
mL	Milliliter
Rpm	Rotation per minute
°C	Degree Celsius
h	hour
Df	Degree of freedom
CV	Coefficient of variation
R <sup>2</sup>	Regression coefficient
Std Dev	Standard deviation
SS	Sum of Square
w/v	Weight per volume
v/v	Volume per volume

mg	Milligram
PO <sub>2</sub>	partial pressure of oxygen
mM	Millimole
sec	Second
IT	Incubation time
AG	Agitation speed
CV	Culture volume
IS	Inoculum size
M	Molar
L	Liter
μg	Microgram
PBD	Plackett- Burman Design
BBD	Box-Behnken Design
CaCl <sub>2</sub>	Calcium Chloride

## CHAPTER 1

### INTRODUCTION

Yeasts provide an alternative platform for recombinant protein production (Gellissen *et al.*, 2005). Yeasts like the prokaryotic system are easy to modify genetically, with a simple fermentation profile. Furthermore, they secrete large amount of glycosylated proteins, a feature peculiar to the eukaryotic system (Gellissen *et al.*, 2005; Böer *et al.*, 2007).

Recently, a limited number of yeasts species able to metabolise methanol as their energy and carbon source were identified. They are being applied in the field of biotechnology for improved yield and cost-effective production of glycosylated recombinant proteins and secretion (Gellissen, 2000; Maleki *et al.*, 2010). These methanol metabolising yeasts are called the methylotrophic yeasts belonging to four genera. These are: *Pichia pastoris* (*Pichia*), *Hansenula polymorpha* (*Hansenula*), *Candida boidinii* (*Candida*) and *Pichia methanolica* (*Torulopsis*) (Hartner & Glieder, 2006).

*Pichia pastoris* is the model organism for the study of methanol utilization in the methylotrophic yeasts and is the most frequently used methylotrophic yeast for recombinant protein production (De Schutter *et al.*, 2009). Its ability to successfully express recombinant proteins is linked to the presence of a tightly regulated promoter system (alcohol oxidase 1) derivable from its methanol utilization pathway genes (Gellissen *et al.*, 2005; Hartner & Glieder, 2006; Böer, *et al.*, 2007; Ahmad *et al.*, 2014). The activation of AOX 1 promoter is fully dependent on methanol and its derivatives but is repressed by other carbon sources in *P. pastoris* (Inan & Meagher, 2001; Ahmad *et al.*, 2014). Methanol is hazardous and flammable in nature making it undesirable for the production of certain foods, pharmaceuticals and industrial products (Ahmad *et al.*, 2014).

A new yeast belonging to the genera *Pichia* of the methylotrophic yeast - *Pichia guilliermondii* (*Meyerozyma guilliermondii*), was isolated from spoiled orange recently in Malaysia (Oslan *et al.*, 2012). Thermostable lipase gene, strain T1, isolated from *Geobacillus zalihae* (Leow *et al.*, 2004), was cloned and expressed in *P. guilliermondii* using *P. pastoris* expression vector (pPICZαB-Invitrogen) (Appendix A). This led to the development of a recombinant host system in *P. guilliermondii* (Oslan *et al.*, 2015). *Pichia guilliermondii* is being developed as an alternative recombinant host to *P. pastoris*.

This study aim at enhancing thermostable T1 lipase production, in recombinant *P. guilliermondii*. The study also, seeks to address the cost of using methanol for gene expression and to optimise fermentation processes involve in protein production in *P. guilliermondii*.

### 1.1 Specific objectives

1. To study and evaluate buffered and non-buffered media compositions suitable for optimum growth of recombinant *P. guilliermondii* and T1 lipase expression. Also, to carryout univariate analysis of *one-factor-at-a-time* for generation of data units, for protein production in recombinant *P. guilliermondii* using the observed optimum medium.
2. To screen and optimise medium parameters in methanol supplemented medium and non-methanol supplemented medium using, Plackett-Burman Design (PBD) and Box-Behnken Design (BBD) of Response Surface Methodology (RSM).
3. To scale up thermostable lipase production at the bioreactor.

## REFERENCES

- Abusham, R. A., Rahman, R. N. Z. R., Salleh, A. B. and Basri, M. (2009). Optimization of physical factors affecting the production of thermo-stable organic solvent-tolerant protease from a newly isolated halo tolerant *Bacillus subtilis* strain Rand. *Microbial Cell Factories* 8(1): 1-9.
- Ahmad, M., Hirz, M., Pichler, H. and Schwab, H. (2014). Protein expression in *Pichia pastoris*: recent achievements and perspectives for heterologous protein production. *Applied Microbiology and Biotechnology* 98(12): 5301-5317.
- Ahmad, M., Hirz, M., Pichler, H. and Schwab, H. (2014). Protein expression in *Pichia pastoris*: recent achievements and perspectives for heterologous protein production. *Applied Microbiology and Biotechnology*, 98(12): 5301-5317.
- Akbari, V., Sadeghi, H. M. M., Jafarian-Dehkordi, A., Chou, C. P. and Abedi, D. (2015). Optimization of a single-chain antibody fragment overexpression in *Escherichia coli* using response surface methodology. *Research in pharmaceutical sciences*, 10(1): 75.
- Anasontzis, G. E., Salazar Penã, M., Spadiut, O., Brumer, H. and Olsson, L. (2014). Effects of temperature and glycerol and methanol-feeding profiles on the production of recombinant galactose oxidase in *Pichia pastoris*. *Biotechnology Progress*, 30(3): 728-735.
- Andualema, B. and Gessesse, A. (2012). Microbial lipases and their industrial applications: Review. *Biotechnology* 11(3): 100-118.
- Aravindan, R., Anbumathi, P. and Viruthagiri, T. (2007). Lipase applications in food industry.
- Ashengroph, M., Nahvi, I. and Amini, J. (2013). Application of Taguchi design and Response surface methodology for improving conversion of isoeugenol into vanillin by resting cells of *Psychrobacter* sp. CSW4. *Iranian Journal of Pharmaceutical Research*, 12(3): 411-421.
- Baharum, S. N., Salleh, A. B., Razak, C. N. A., Basri, M. and Rahman R.N.Z.R.A. (2003). Organic solvent tolerant lipase by *Pseudomonas* sp. strain S5: stability of enzyme in organic solvent and physical factors affecting its production. *Annals Microbiology*, 53, 75-83.
- Bakri, Y., Mekaeel, A. and Koreih, A. (2011). Influence of agitation speeds and aeration rates on the Xylanase activity of *Aspergillus niger* SS7. *Brazilian Archives of Biology and Technology*, 54(4): 659-664.b
- Bandaranayake, A. D. and Almo, S. C. (2014). Recent advances in mammalian protein production. *FEBS letters*, 588(2): 253-260.

- Bekatorou, A., Psarianos, C. and Koutinas, A. A. (2006). Production of food grade yeasts. *Food Technology and Biotechnology*, 44(3): 407-415.
- Bezerra, M. A., Santelli, R. E., Oliveira, E. P., Villar, L. S. and Escaleira, L. A. (2008). Response surface methodology (RSM) as a tool for optimization in analytical chemistry. *Talanta*, 76(5): 965-977.
- Bibhu, P. P., Mohd, A. and Saleem, J. (2007). Fermentation Process Optimization. *Research Journal of Microbiology*, 2: 201-208.
- Bill, R. M. (2014). Playing catch-up with Escherichia coli: using yeast to increase success rates in recombinant protein production experiments. *Frontiers in Microbiology*, 5.
- Branduardi, P., Valli, M., Brambilla, L., Sauer, M., Alberghina, L. and Porro, D. (2004). The yeast *Zygosaccharomyces bailii*: a new host for heterologous protein production, secretion and for metabolic engineering applications. *FEM Yeast Research*, 4(4-5): 493-504.
- Butler, M. and Meneses-Acosta, A. (2012). Recent advances in technology supporting biopharmaceutical production from mammalian cells. *Applied Microbiology and Biotechnology* 96(4): 885-894.
- Byrne, B. (2015). *Pichia pastoris* as an expression host for membrane protein structural biology. *Current Opinion in Structural Biology*, 32, 9-17.
- Cereghino, G. P. L., Cereghino, J. L., Ilgen, C and Cregg, J. M. (2002). Production of recombinant proteins in fermenter cultures of the yeast *Pichia pastoris*. *Current Opinion in Biotechnology*, 13(4): 329-332.
- Cereghino, J. L. and Cregg, J. M. (2000). Heterologous protein expression in the methylotrophic yeast *Pichia pastoris*. *FEM Microbiology Reviews*, 24(1): 45-66.
- Chauhan, M., and Garlapati, V. K. (2013). Production and characterization of a halo-, solvent-, thermo-tolerant alkaline lipase by *Staphylococcus arlettae* JPBW-1, isolated from rock salt mine. *Applied Biochemistry and Biotechnology*, 171(6): 1429-1443.
- Chisti, Y. (1999). Fermentation (industrial): basic considerations. *Encyclopedia of Food Microbiology*, 663-674.
- Ciafardini, G., Zullo, B. A., Cioccia, G., and Iride, A. (2006). Lipolytic activity of *Williopsis californica* and *Saccharomyces cerevisiae* in extra virgin olive oil. *International Journal of Food Microbiology*, 107(1): 27-32.
- Colla, L. M., Primaz, A. L., Benedetti, S., Loss, R. A., de Lima, M., Reinehr, C. O. and Costa, J. A. V. (2016). Surface response methodology for the optimization of lipase production under submerged fermentation by filamentous fungi. *Brazilian Journal of Microbiology* 47(2): 461-467.



- Collet, C., Adler, N., Schwitzguébel, J. P. and Péringer, P. (2004). Hydrogen production by *Clostridium thermolacticum* during continuous fermentation of lactose. *International Journal of Hydrogen Energy*, 29(14): 1479-1485.
- Couto, S. R. and Sanromán, M. A. (2006). Application of solid-state fermentation to food industry-a review. *Journal of Food Engineering*, 76(3): 291-302.
- Cox, M.M.J. (2004). Commercial production in insect cells: One company's perspective. *Bioprocess International*:1-5.
- Dalton, A. C. and Barton, W. A. (2014). Over-expression of secreted proteins from mammalian cell lines. *Protein Science* 23(5): 517-525.
- De los Reyes-Gavilán, C. G., Fernández, M., Hudson, J. A. and Korpela, R. (2015). Role of microorganisms present in dairy fermented products in health and disease. *BioMed Research International*, 2015.
- De Schutter, K., Lin, Y.C., Tiels, P., Van Hecke, A., Glinka, S., Weber-Lehmann, J., Rouzé, P., Van de Peer, Y. and Callewaert, N. (2009). Genome sequence of the recombinant protein production host *Pichia pastoris*. *Nature Biotechnology*, 27(6): pp.561-566.
- Deepak, V., Kalishwaralal, K., Ramkumarpandian, S., Babu, S. V., Senthilkumar, S. R. and Sangiliyandi, G. (2008). Optimization of media composition for Nattokinase production by *Bacillus subtilis* using response surface methodology. *Bioresource Technology*, 99(17): 8170-8174.
- Demirci, A., Izmirlioglu, G. and Ercan, D. (2014). Fermentation and enzyme technologies in food processing. *Food Processing: Principles and Applications, Second Edition*, 107-136.
- Dinarvand, M., Rezaee, M., Masomian, M., Jazayeri, S. D., Zareian, M., Abbasi, S. and Ariff, A. B. (2013). Effect of C/N ratio and media optimization through response surface methodology on simultaneous productions of intra-and extracellular inulinase and invertase from *Aspergillus niger* ATCC 20611. *BioMed Research International*, 2013.
- Djekrif-Dakhmouche, S., Gheribi-Aoulmi, Z., Meraihi, Z. and Bennamoun, L. (2006). Application of a statistical design to the optimization of culture medium for  $\alpha$ -amylase production by *Aspergillus niger* ATCC 16404 grown on orange waste powder. *Journal of Food Engineering*, 73(2): 190-197.
- Drake, D. R. and Brogden, K. A. (2002). Continuous-culture chemostat systems and flowcells as methods to investigate microbial interactions.
- El-Gendy, N., Madian, H. R. and Amr, S. S. (2012). Design and Optimization of a Process for Sugarcane Molasses Fermentation by *Saccharomyces cerevisiae* Using Response Surface Methodology. *International Journal of Microbiology*, (2013): 815631-815631.

- Fakruddin, M., Mohammad Mazumdar, R., Bin Mannan, K. S., Chowdhury, A. and Hossain, M. N. (2012). Critical factors affecting the success of cloning, expression, and mass production of enzymes by recombinant *E. coli*. *ISRN Biotechnology*, 2013.
- Fang, Z., Xu, L., Pan, D., Jiao, L., Liu, Z., and Yan, Y. (2014). Enhanced production of *Thermomyces lanuginosus* lipase in *Pichia pastoris* via genetic and fermentation strategies. *Journal of Industrial Microbiology and Biotechnology*, 41(10): 1541-1551.
- Farias, C.M., de Souza, O.C., Sousa, M.A., Cruz, R., Magalhães, O.M.C., de Medeiros, É.V., Moreira, K.A. and de Souza-Motta, C.M. (2015). High-level lipase production by *Aspergillus candidus* URM 5611 under solid state fermentation (SSF) using waste from *Siagrus coronata* (Martius) Becari. *African Journal of Biotechnology*, 14(9): 820-828.
- Ferreira, S.C., Bruns, R.E., Ferreira, H.S., Matos, G.D., David, J.M., Brandao, G.C., da Silva, E.P., Portugal, L.A., Dos Reis, P.S., Souza, A.S. and Dos Santos, W.N.L. (2007). Box-Behnken design: an alternative for the optimization of analytical methods. *Analytica Chimica Acta*, 597(2): pp.179-186.
- Frenzel, A., Hust, M. and Schirrmann, T. (2013). Expression of recombinant antibodies. *Frontiers in Immunology*, 4.
- Gangadharan, D., Sivaramakrishnan, S., Nampoothiri, K. M., Sukumaran, R. K. and Pandey, A. (2008). Response surface methodology for the optimization of alpha amylase production by *Bacillus amyloliquefaciens*. *Bioresource Technology*, 99(11): 4597-4602.
- Gellissen, G. (Ed.). (2006). Production of recombinant proteins: Novel microbial and eukaryotic expression systems. *John Wiley and Sons*. Page: 429
- Gellissen, G. and Veenhuis, M. (2001). The methylotrophic yeast *Hansenula polymorpha*: its use in fundamental research and as a cell factory. *Yeast* (Chichester, England) 18(3): i-iii.
- Gellissen, G., Kunze, G., Gaillardin, C., Cregg, J. M., Berardi, E., Veenhuis, M. and Van Der Klei, I. (2005). New yeast expression platforms based on methylotrophic *Hansenula polymorpha* and *Pichia pastoris* and on dimorphic *Arxula adenivorans* and *Yarrowia lipolytica*—a comparison. *FEM Yeast Research* 5(11): 1079-1096.
- Gerngross, T. U. (2004). Advances in the production of human therapeutic proteins in yeasts and filamentous fungi. *Nature Biotechnology* 22(11): 1409-1414.
- Gomes, A., Byregowda, S., Veeregowda, B. and Balamurugan, V. (2016). An overview of heterologous expression host systems for the production of recombinant proteins. *Adv. Animal Veterinary Science*, 4(7): 346-356.

- Graf, A., Dragosits, M., Gasser, B. and Mattanovich, D. (2009). Yeast systems biotechnology for the production of heterologous proteins. *FEM Yeast Research* 9(3): 335-348.
- Guengerich, L., Kang, H. A., Behle, B., Gellissen, G. and Suckow, M. (2004). A platform for heterologous gene expression based on the methylotrophic yeast *Hansenula polymorpha*. In *Genetics and Biotechnology* (pp. 273-287).
- Gunasekaran, V. and Das, D. (2005). Lipase fermentation: progress and prospects. *Indian Journal of Biotechnology* 4(4): 437-445.
- Gupta, R., Beg, Q. and Lorenz, P. (2002). Bacterial alkaline proteases: molecular approaches and industrial applications. *Applied Microbiology and Biotechnology*, 59(1), 15-32.
- Gurung, N., Ray, S., Bose, S. and Rai, V. (2013). A broader view: microbial enzymes and their relevance in industries, medicine, and beyond. *BioMed Research International*, 2013.
- Haddar, A., Fakhfakh-Zouari, N., Hmidet, N., Frikha, F., Nasri, M. and Kamoun, A. S. (2010). Low-cost fermentation medium for alkaline protease production by *Bacillus mojavensis* A21 using hulled grain of wheat and sardinella peptone. *Journal of Bioscience and Bioengineering*, 110(3): 288-294.
- Hartner, F. S. and Glieder, A. (2006). Regulation of methanol utilisation pathway genes in yeasts. *Microbial Cell Factories* vol. 5, article 39.
- Harzevili, F. D. and Chen, H. (2015). *Microbial biotechnology: Progress and trends*. Boca Raton: CRC Press, Taylor and Francis Group, CRC Press is an imprint of the Taylor and Francis Group, 412pp.
- Heijnen, J. J. and Van Dijken, J. P. (1992). In search of a thermodynamic description of biomass yields for the chemotrophic growth of microorganisms. *Biotechnology and Bioengineering*, 39(8): 833-858.
- Hensing, M. C. M., Rouwenhorst, R. J., Heijnen, J. J., Van Dijken, J. P. and Pronk, J. T. (1995). Physiological and technological aspects of large-scale heterologous-protein production with yeasts. *Antonie Van Leeuwenhoek* 67(3): 261-279.
- Hölker, U., Höfer, M. and Lenz, J. (2004). Biotechnological advantages of laboratory-scale solid-state fermentation with fungi. *Applied Microbiology and Biotechnology*, 64(2): 175-186.
- Holmes, W. J., Darby, R. A., Wilks, M. D., Smith, R. and Bill, R. M. (2009). Developing a scalable model of recombinant protein yield from *Pichia pastoris*: the influence of culture conditions, biomass and induction regime. *Microbial Cell Factories* 8(1): 1-14.

- Hsu, A. F., Jones, K., Foglia, T. A. and Marmer, W. N. (2002). Immobilized lipase-catalysed production of alkyl esters of restaurant grease as biodiesel. *Biotechnology and Applied Biochemistry* 36(3): 181-186.
- Inan, M. and Meagher, M. M. (2001). Non-repressing carbon sources for alcohol oxidase (AOX1) promoter of *Pichia pastoris*. *Journal of Bioscience and Bioengineering*, 92(6): 585-589.
- Irfan, M., Nadeem, M. and Syed, Q. (2014). One-factor-at-a-time (OFAT) optimization of xylanase production from *Trichoderma viride*-IR05 in solid-state fermentation. *Journal of Radiation Research and Applied Sciences*, 7(3): 317-326.
- Jafari, A. R., Sarrafzadeh, M. H., Alemzadeh, I. and Vosoughi, M. (2007). Effect of stirrer speed and aeration rate on the production of glucose oxidase by *Aspergillus niger*. *Journal of Biological Sciences*, 7(2), 270-275.
- Jia, J., Yang, X., Wu, Z., Zhang, Q., Lin, Z., Guo, H., Lin, C.S.K., Wang, J. and Wang, Y. (2015). Optimization of fermentation medium for extracellular lipase production from *Aspergillus niger* using response surface methodology. *BioMed Research International*, 2015.
- Kakde, R. B. and Chavan, A. M. (2011). Extracellular lipase enzyme production by seed-borne fungi under the influence of physical factors. *International Journal of Biology*, 3(1): 94.
- Kang, H. A, Gellissen G: *Hansenula polymorpha*. In Production of recombinant proteins: novel microbial and eukaryotic expression systems. Edited by: Gellissen G. Weinheim: Wiley-VCH; 2005:111-142.
- Kanmani, P., Karthik, S., Aravind, J. and Kumaresan, K. (2012). The use of response surface methodology as a statistical tool for media optimization in lipase production from the dairy effluent isolate *Fusarium solani*. *ISRN Biotechnology*, 2013.
- Kaszycki, P., Tyszka, M., Malec, P., and Kołoczek, H. (2001). Formaldehyde and methanol biodegradation with the methylotrophic yeast *Hansenula polymorpha*. An application to real wastewater treatment. *Biodegradation*, 12(3): 169-177.
- Kaushik, R., Saran, S., Isar, J. and Saxena, R. K. (2006). Statistical optimization of medium components and growth conditions by response surface methodology to enhance lipase production by *Aspergillus carneus*. *Journal of Molecular Catalysis B: Enzymatic*, 40(3): 121-126.
- Khoo, M. L. and Ibrahim, C. O. (2003). Development of alkaline lipase for the formulation of detergent. In Proceedings of *National Biology Conference* (pp. 14-16).

- Kim, H., Yoo, S. J. and Kang, H. A. (2014). Yeast synthetic biology for the production of recombinant therapeutic proteins. *FEM Yeast Res*, 15, 1-16.
- Kost, T. A., Condreay, J. P. and Jarvis, D. L. (2005). Baculovirus as versatile vectors for protein expression in insect and mammalian cells. *Nature Biotechnology*, 23(5): 567.
- Krishna, C. (2005). Solid-state fermentation systems—an overview. *Critical Reviews in Biotechnology*, 25(1-2): 1-30.
- Kumar, M., Joshi, A., Kashyap, R. and Khanna, S. (2011). Production of xylanase by *Promicromonospora* sp MARS with rice straw under non sterile conditions. *Process Biochemistry*, 46(8): 1614-1618.
- Kumari, A., Mahapatra, P. and Banerjee, R. (2009). Statistical optimization of culture conditions by response surface methodology for synthesis of lipase with *Enterobacter aerogenes*. *Brazilian Archives of Biology and Technology*, 52(6): 1349-1356.
- Kunze, G., Kang, H. A. and Gellissen, G. (2009). *Hansenula polymorpha* (*Pichia angusta*): Biology and Applications. *Springer* Netherlands, pp. 47-64.
- Kurtovic, I., Marshall, S. N., Zhao, X. and Simpson, B. K. (2009). Lipases from mammals and fishes. *Reviews in Fisheries Science* 17(1): 18-40.
- Lackner, A., Genta, K., Koppensteiner, H., Herbacek, I., Holzmann, K., Spiegl-Kreinecker, S. and Grusch, M. (2008). A bicistronic baculovirus vector for transient and stable protein expression in mammalian cells. *Analytical Biochemistry* 380(1): 146-148.
- Lan, D., Qu, M., Yang, B. and Wang, Y. (2016). Enhancing production of lipase MAS1 from marine *Streptomyces* sp. strain in *Pichia pastoris* by chaperones co-expression. *Electronic Journal of Biotechnology*, 22, 62-67.
- Leow, T. C., Rahman, R. N. Z. R. A., Basri, M. and Salleh, A. B. (2004). High level expression of thermostable lipase from *Geobacillus species* strain T1. *Bioscience, Biotechnology and Biochemistry* 68(1): 96-103.
- Li, F., Vijayasankaran, N., Shen, A., Kiss, R. and Amanullah, A. (2010). Cell culture processes for monoclonal antibody production. In *MAbs* (Vol. 2, No. 5, pp. 466-479). Taylor & Francis.
- Li, J., Tang, C., Shi, H. and Wu, M. (2011). Cloning and optimized expression of a neutral endoglucanase gene (nce15A) from *Volvariella volvacea* WX32 in *Pichia pastoris*. *Journal of Bioscience and Bioengineering*, 111(5): 537-540.
- Li, T., Chen, X. B., Chen, J. C., Wu, Q. and Chen, G. Q. (2014). Open and continuous fermentation: Products, conditions and bioprocess economy. *Biotechnology Journal*, 9(12): 1503-1511.

- Longobardi, G. P. (1994). Fed-batch versus batch fermentation. *Bioprocess and Biosystems Engineering*, 10(5): 185-194.
- Macauley-Patrick, S. and Finn, B. (2008). Modes of fermenter operation. Practical fermentation technology. John Wiley and Sons, Ltd., West Sussex, England, 69-95.
- Macauley-Patrick, S., Fazenda, M. L., McNeil, B. and Harvey, L. M. (2005). Heterologous protein production using the *Pichia pastoris* expression system. *Yeast*, 22(4): 249-270.
- Machida, M., Yamada, O. and Gomi, K. (2008). Genomics of *Aspergillus oryzae*: learning from the history of Koji mold and exploration of its future. *DNA Research* 15(4): 173-183.
- Maleki, A., Roohvand, F., Tajerzadeh, H., Khanahmad, H., Nobari, M. B., Beirut, A. and Najafabadi, A. R. (2010). High expression of methylotrophic yeast-derived recombinant human erythropoietin in a pH-controlled batch system. *Avicenna Journal of Medical Biotechnology*, 2(4), 197.
- Manderson, D., Dempster, R. and Chisti, Y. (2006). Production of an active recombinant Aspin antigen in *Escherichia coli* for identifying animals resistant to nematode infection. *Enzyme and Microbial technology*, 38(5): 591-598.
- Mattanovich, D., Branduardi, P., Dato, L., Gasser, B., Sauer, M. and Porro, D. (2012). Recombinant protein production in yeasts. Recombinant gene expression, 329-358.
- Mohajeri, A., Abdolalizadeh, J., Pilehvar-Soltanahmadi, Y., Kiafar, F. and Zarghami, N. (2016). Expression and Secretion of Endostar Protein by *Escherichia Coli*: Optimization of Culture Conditions Using the Response Surface Methodology. *Molecular Biotechnology*, 58(10): 634-647.
- Mourabet, M., El Rhilassi, A., El-Boujaady, H., Bennani-Ziatni, M. and Taitai, A. (2014). Use of response surface methodology for optimization of fluoride adsorption in an aqueous solution by Brushite. *Arabian Journal of Chemistry*.
- Muralidhar, R. V., Chirumamila, R. R., Marchant, R. and Nigam, P. (2001). A response surface approach for the comparison of lipase production by *Candida cylindracea* using two different carbon sources. *Biochemical Engineering Journal*, 9(1): 17-23.
- Nallathambi, V., Jayaraman, A., Angayarkanni, P. and Govindasamy, N. (2016). Application of statistical designs for the optimization of medium constituents for the production of pterin deaminase from *Aspergillus terreus*. *International Journal of Recent Scientific Research*, 7(4): 10678-10685.
- Nasser, M. W., Pooja, V., Abdin, M. Z. and Jain, S. K. (2003). Evaluation of yeast as an expression system. *Indian Journal of Biotechnology* 2(4): 477-493.

- Negruță, O., Csutak, O., Stoica, I., Rusu, E., and Vassu, T. (2010). Methylotrophic yeasts: diversity and methanol metabolism. *Romanian Biotechnological Letters*, 15(4).
- Nevoigt, E. (2008). Progress in metabolic engineering of *Saccharomyces cerevisiae*. *Microbiology and Molecular Biology Reviews*, 72(3): 379-412.
- O'Reilly, D. R., Miller, L. K. and Luckow, V. A. (1992). *Baculovirus Expression Vector: A Laboratory Manual* WH Freeman and Company. New York. 347 pp.
- Oslan, S. N., Salleh, A. B., Rahman, R. A., Zaliha, R. N., Leow, T. C., Sukamat, H. and Basri, M. (2015). A newly isolated yeast as an expression host for recombinant lipase. *Cellular and Molecular Biology Letters* 20(2): 279-293.
- Oslan, S. N., Salleh, A. B., Rahman, R. A., Zaliha, R. N., Leow, A. T. C. and Basri, M. (2014). *Pichia pastoris* as a host to overexpress the thermostable T1 lipase from *Geobacillus zalihae*. *GSTF Journal of Biosciences*, 3(1): 7-17.
- Oslan, S. N., Salleh, A. B., Rahman, R. N. Z. R. A., Basri, M. and Chor, A. L. T. (2012). Locally isolated yeasts from Malaysia: Identification, phylogenetic study and characterization. *Acta Biochimica Polonica*, 59(2): 225-229.
- Pandey, A. (2003). Solid-state fermentation. *Biochemical Engineering Journal*, 13(2): 81-84.
- Paques, F. W., Pio, T. F., Carvalho, P. D. O. and Macedo, G. A. (2008). Characterization of the lipase from *Carica papaya* residues. *Braz. Journal Food Technol*, 11(1), 20-27.
- Porro, D. and Mattanovich, D. (2004). Recombinant protein production in yeasts. *Recombinant Gene Expression: Reviews and Protocols*, 241-258.
- Porro, D., Sauer, M., Branduardi, P. and Mattanovich, D. (2005). Recombinant protein production in yeasts. *Molecular Biotechnology* 31(3): 245-259.
- Prielhofer, R., Cartwright, S. P., Graf, A. B., Valli, M., Bill, R. M., Mattanovich, D. and Gasser, B. (2015). *Pichia pastoris* regulates its gene-specific response to different carbon sources at the transcriptional, rather than the translational, level. *BMC Genomics*, 16(1): 1.
- Qiu, P., Cui, M., Kang, K., Park, B., Son, Y., Khim, E., Jang, M. and Khim, J. (2014). Application of Box-Behnken design with response surface methodology for modeling and optimizing ultrasonic oxidation of arsenite with H<sub>2</sub>O<sub>2</sub>. *Central European Journal of Chemistry*, 12(2): pp.164-172.
- Rai, M. and Padh, H. (2001). Expression systems for production of heterologous proteins. *Current Science-Bangalore* 80(9): 1121-1128.

- Rajakumara, E., Acharya, P., Ahmad, S., Sankaranaryanan, R. and Rao, N. M. (2008). Structural basis for the remarkable stability of *Bacillus subtilis* lipase (Lip A) at low pH. *Biochimica et Biophysica Acta (BBA)-Proteins and Proteomics*, 1784(2), 302-311.
- Rajendran, A., Palanisamy, A. and Thangavelu, V. (2008). Evaluation of medium components by Plackett-Burman statistical design for lipase production by *Candida rugosa* and kinetic modeling. *Chinese Journal of Biotechnology*, 24(3): 436-444.
- Ramezani-Rad, M., Hollenberg, C.P., Lauber, J., Wedler, H., Griess, E., Wagner, C., Albermann, K., Hani, J., Piontek, M., Dahlems, U. and Gellissen, G. (2003). The *Hansenula polymorpha* (strain CBS4732) genome sequencing and analysis. *FEM Yeast Research*, 4(2): 207-215.
- Rao, K. J., Kim, C. H. and Rhee, S. K. (2000). Statistical optimization of medium for the production of recombinant hirudin from *Saccharomyces cerevisiae* using response surface methodology. *Process Biochemistry*, 35(7): 639-647.
- Reddy, L. V. A., Wee, Y. J., Yun, J. S. and Ryu, H. W. (2008). Optimization of alkaline protease production by batch culture of *Bacillus* sp. RKY3 through Plackett-Burman and response surface methodological approaches. *Bioresource Technology*, 99(7): 2242-2249.
- Rhee, S. J., Lee, J. E. and Lee, C. H. (2011). Importance of lactic acid bacteria in Asian fermented foods. *Microbial Cell Factories*, 10(1), S5.
- Rodrigues, L., Teixeira, J., Oliveira, R. and Van Der Mei, H. C. (2006). Response surface optimization of the medium components for the production of biosurfactants by probiotic bacteria. *Process Biochemistry* 41(1): 1-10.
- Rosano, G. L. and Ceccarelli, E. A. (2014). Recombinant protein expression in *Escherichia coli*: advances and challenges. *Frontiers in Microbiology*, 5.
- Salihu, A. and Alam, M. Z. (2015). Solvent tolerant lipases: a review. *Process Biochemistry* 50(1): 86-96.
- Salihu, A., Alam, M. Z., AbdulKarim, M. I. and Salleh, H. M. (2011). Optimization of lipase production by *Candida cylindracea* in palm oil mill effluent based medium using statistical experimental design. *Journal of Molecular Catalysis B: Enzymatic*, 69(1): 66-73.
- Salis, A., Solinas, V. and Monduzzi, M. (2003). Wax esters synthesis from heavy fraction of sheep milk fat and cetyl alcohol by immobilised lipases. *Journal of Molecular Catalysis B: Enzymatic* 21(4): 167-174.
- Satyanarayana, T. and Kunze, G. (Eds.). (2009). Yeast biotechnology: diversity and applications (Vol. 78). *Dordrecht: Springer*.



- Saxena, K., Dutta, A., Klein-Seetharaman, J. and Schwalbe, H. (2012). Isotope labeling in insect cells. *Protein NMR Techniques*, 37-54.
- Schaefer, S., Piontek, M., Ahn, S.J., Papendieck, A., Janowicz, Z.A., Timmermans, I. and Gellissen, G. (2002). *Hansenula polymorpha*–Biology and Applications. 175-210.
- Schmidt, F. R. (2005). Optimization and scale up of industrial fermentation processes. *Applied Microbiology and Biotechnology*, 68(4), 425-435.
- Seth, S., Chakravorty, D., Dubey, V. K. and Patra, S. (2014). An insight into plant lipase research–challenges encountered. *Protein Expression and Purification* 95: 13-21.
- Sethi, B. K., Nanda, P. K. and Sahoo, S. (2016). Characterization of biotechnologically relevant extracellular lipase produced by *Aspergillus terreus* NCFT 4269.10. *Brazilian Journal of Microbiology*, 47(1), 143-149.
- Sethi, B. K., Rout, J. R., Das, R., Nanda, P. K. and Sahoo, S. L. (2013). Lipase production by *Aspergillus terreus* using mustard seed oil cake as a carbon source. *Annals of Microbiology*, 63(1), 241-252.
- Sharma, S., and Kanwar, S. S. (2014). Organic solvent tolerant lipases and applications. *The Scientific World Journal*, 2014.
- Sharma, R., Soni, S. K., Vohra, R. M., Gupta, L. K. and Gupta, J. K. (2002). Purification and characterisation of a thermostable alkaline lipase from a new thermophilic *Bacillus* sp. RSJ-1. *Process Biochemistry*, 37(10): 1075-1084.
- Shimada, Y., Ogawa, J., Watanabe, Y., Nagao, T., Kawashima, A., Kobayashi, T. and Shimizu, S. (2003). Regiospecific analysis by ethanolysis of oil with immobilized *Candida antarctica* lipase. *Lipids* 38(12): 1281-1286.
- Sibirny, A. A. and Boretsky, Y. R. (2009). *Pichia guilliermondii*. In *Yeast biotechnology: diversity and applications* (pp. 113-134). Springer Netherlands.
- Silva, C. J. S. M. and Roberto, I. C. (2001). Improvement of xylitol production by *Candida guilliermondii* FTI 20037 previously adapted to rice straw hemicellulosic hydrolysate. *Letters in Applied Microbiology*, 32(4): 248-252.
- Singh, V., Haque, S., Niwas, R., Srivastava, A., Pasupuleti, M, and Tripathi, C. K. M. (2016). Strategies for fermentation medium optimization: an in-depth review. *Frontiers in Microbiology*, 7.
- Singhania, R. R., Sukumaran, R. K., Patel, A. K., Larroche, C. and Pandey, A. (2010). Advancement and comparative profiles in the production technologies using solid-state and submerged fermentation for microbial cellulases. *Enzyme and Microbial Technology*, 46(7): 541-549.

- Sørensen, H. P. (2010). Towards universal systems for recombinant gene expression. *Microbial Cell Factories* 9(1): 1-4.
- Stanbury, P. F., Whitaker, A. and Hall, S. J. (2013). Principles of fermentation technology. *Elsevier*.
- Steensels, J., Snoek, T., Meersman, E., Nicolino, M. P., Voordeckers, K. and Verstrepen, K. J. (2014). Improving industrial yeast strains: exploiting natural and artificial diversity. *FEM Microbiology Reviews*, 38(5): 947-995.
- Stöckmann, C., Scheidle, M., Dittrich, B., Merkelbach, A., Hehmann, G., Melmer, G., Klee, D., Büchs, J., Kang, H.A. and Gellissen, G. (2009). Process development in *Hansenula polymorpha* and *Arxula adenivorans*, a re-assessment. *Microbial Cell Factories*, 8(1): p.22.
- Takagi, S., Tsutsumi, N., Terui, Y. and Kong, X. (2012). U.S. Patent No. 8,236,528. Washington, DC: U.S. Patent and Trademark Office.
- Thakur, V., Tewari, R. and Sharma, R. (2014). Evaluation of production parameters for maximum lipase production by *Pseudomonas stutzeri* MTCC 5618 and scale-up in bioreactor. *Chinese Journal of Biology*, 2014.
- Treanor, J.J., Wilkinson, B.E., Masseur, F., Hu-Primmer, J., Battaglia, R., O'Brien, D., Wolff, M., Rabinovich, G., Blackwelder, W. and Katz, J.M. (2001). Safety and immunogenicity of a recombinant hemagglutinin vaccine for H5 influenza in humans. *Vaccine*, 19(13): 1732-1737.
- Vakhlu, J. and Kour, A. (2006). Yeast lipases: enzyme purification, biochemical properties and gene cloning. *Electronic Journal of Biotechnology*, 9(1): 69-85.
- Van Dijk, R., Faber, K. N., Kiel, J. A., Veenhuis, M. and van der Klei, I. (2000). The methylotrophic yeast *Hansenula polymorpha*: a versatile cell factory. *Enzyme and Microbial Technology*, 26(9): 793-800.
- Van Ooyen, A. J., Dekker, P., Huang, M., Olsthoorn, M. M., Jacobs, D. I., Colussi, P. A. and Taron, C. H. (2006). Heterologous protein production in the yeast *Kluyveromyces lactis*. *FEM Yeast Research*, 6(3): 381-392.
- Verbelen, P. J., De Schutter, D. P., Delvaux, F., Verstrepen, K. J. and Delvaux, F. R. (2006). Immobilized yeast cell systems for continuous fermentation applications. *Biotechnology Letters*, 28(19): 1515-1525.
- Vici, A.C., da Cruz, A.F., Facchini, F.D., de Carvalho, C.C., Pereira, M.G., Fonseca-Maldonado, R., Ward, R.J., Pessela, B.C., Fernandez-Lorente, G., Torres, F.A. and Jorge, J.A. (2015). Beauveria bassiana Lipase A expressed in *Komagataella (Pichia) pastoris* with potential for biodiesel catalysis. *Frontiers in Microbiology*, 6, p.1083.

Wurm, F. M. (2004). Production of recombinant protein therapeutics in cultivated mammalian cells. *Nature Biotechnology* 22(11): 1393-1398.

Yin, T., Miao, L.L., Guan, F.F., Wang, G.L., Peng, Q., Li, B.X., Guan, G.H. and Li, Y. (2010). Optimized medium improves expression and secretion of extremely thermostable bacterial xylanase, XynB, in *Kluyveromyces lactis*. *Journal of Microbiology and Biotechnology*, 20(11), pp.1471-1480.

Yurimoto, H., Oku, M. and Sakai, Y. (2011). Yeast methylotrophy: metabolism, gene regulation and peroxisome homeostasis. *International Journal of Microbiology*.

Yurimoto, H., Oku, M. and Sakai, Y. (2011). Yeast methylotrophy: metabolism, gene regulation and peroxisome homeostasis. *International Journal of Microbiology*, 2011.

