



**UNIVERSITI PUTRA MALAYSIA**

***OPTIMISATION OF MEDIUM FORMULATION AND SCALING UP OF  
THREONINE AND TRYPTOPHAN PRODUCTION BY LACTIC ACID  
BACTERIA USING RESPONSE SURFACE METHODOLOGY***

**LIM YE HENG**

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By

**LIM YE HENG**

**Thesis Submitted to the School of Graduate Studies, Universiti  
Putra Malaysia, in Fulfilment of the Requirements for the Degree of  
Doctor of Philosophy**

**December 2017**

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Abstract of thesis presented to the Senate of Universiti Putra Malaysia in fulfilment of the requirement for the degree of Doctor of Philosophy

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**December 2017**

**Chair : Professor Foo Hooi Ling, PhD**  
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Increasing knowledge on the functions of amino acid (AA) in animal production has led to escalating demand of various amino acid. Threonine and tryptophan are among the most commonly employed feed AA due to their indispensable roles in enhancing the growth performance of livestock. Currently, AA production relies heavily on non-food-grade-microorganisms such as genetically modified *Corynebacterium glutamicum* and *Escherichia coli* which was a concern as the use of genetically modified *C. glutamicum* for production of amino acid was linked to over thousand cases of a deadly syndrome, eosinophilia myalgia syndrome (EMS). This has urged for search of safer alternatives by utilising food-grade-microorganisms. Recent studies reported that lactic acid bacteria (LAB) were capable to produce various AA owing to their well-established proteolytic system and presence of AA biosynthesis gene. Furthermore, they are reputed with the Generally Recognised as Safe (GRAS) status, making them an excellent candidate as food grade producer. However, there were limited studies regarding production of AA by using LAB. Hence, the objective of this study was to identify the threonine and tryptophan producing LAB and optimise the medium formulation via response surface methodology (RSM) approach, followed by scaling up their production by using constant impeller tip speed approach. It was hypothesised that threonine and tryptophan producing LAB could be identified and their production could be improved by optimisation of the medium formulation using RSM. Additionally, the production of threonine and tryptophan by the selected LAB could be scaled up constant impeller tip speed approach. In this study, 17 LAB isolates from Malaysian foods were identified phenotypically and genotypically. The isolates comprised of 3 species: *Pediococcus pentosaceus* (6 isolates), *Pediococcus acidilactici* (2 isolates), and *Lactobacillus plantarum* (9 isolates). Thereafter, the growth profile of the isolates were characterised and their proteolytic activity was determined qualitatively and quantitatively under 3 pH conditions by using skim milk agar hydrolysis assay, skim milk agar well diffusion assay and azocasein assay due

to the important role of proteolytic activity on amino acid production. All the LAB isolates exhibited versatile extracellular proteolytic system where proteolytic activity was detected over wide pH range. The highest extracellular proteolytic activity at pH 5 (15.76 U/mg) and pH 8 (19.42 U/mg) was detected in *L. plantarum* RG14. Meanwhile, *L. plantarum* RS5 and RI11 demonstrated the highest extracellular proteolytic activity of approximately 17 U/mg at pH 6.5.

The ability of the LAB isolates to produce AA was subsequently determined by cultivating in de Man Rogosa and Sharpe (MRS) medium. The production of amino acid was quantified by using high performance liquid chromatography (HPLC) analysis. The LAB isolates demonstrated the ability to produce numerous industrially important AA but the production was strain dependent. *P. pentosaceus* TL-3 demonstrated highest threonine productivity (12.88 mg/L/h) and identified as superior threonine producer in this study. Meanwhile, *P. acidilactici* TP-6 was selected as tryptophan producer with a productivity of 5.05 mg/L/h. The production of threonine and tryptophan by the selected LAB isolate was subsequently optimised by using RSM. The nutritional requirement for threonine and tryptophan production was first evaluated by using Plackett Burman Design (PBD) and subsequently optimised by using Central Composite Design (CCD). Molasses, meat extract,  $(\text{NH}_4)_2\text{SO}_4$  and  $\text{MnSO}_4$  were the most important components for threonine production by *P. pentosaceus* TL-3 with an optimum concentration of 30.79 g/L, 25.30 g/L, 8.59 g/L, and 0.098 g/L respectively. The net threonine produced recorded by *P. pentosaceus* TL-3 under optimised condition (125.98 mg/L) was improved by 2 fold whereas the cost of the optimised medium was reduced by 8.5% compared to MRS medium.

In comparison, the best combination of medium components for tryptophan production by *P. acidilactici* TP-6 were molasses, meat extract, urea and  $\text{FeSO}_4$ . The optimum concentration suggested by RSM were: molasses, 14.06 g/L; meat extract 23.68 g/L; urea, 5.56 g/L and  $\text{FeSO}_4$ , 0.024 g/L. Up to 68.05 mg/L of tryptophan was produced by *P. acidilactici* TP-6 under optimised condition, which was equivalent to 150% enhancement compared to the control. In contrast, the cost of the optimised medium was reduced by 11%. Furthermore, the production of threonine and tryptophan by the selected LAB isolate was successfully scaled up in 30 L stirred tank bioreactor based on constant impeller tip speed approach. Additionally, the net threonine and tryptophan produced in bioreactor cultivation was comparable to the predicted amount suggested by CCD. In conclusion, *P. pentosaceus* TL-3 and *P. acidilactici* TP-6 were identified as threonine and tryptophan producer respectively and the production of threonine and tryptophan by the selected producer strain was enhanced by 2 folds and 150% respectively through optimisation of their medium formulation using RSM approach. Additionally, the production of threonine and tryptophan was successfully scaled by based on constant impeller tip speed approach where the production in both small and large scale cultivation was comparable.

Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia  
sebagai memenuhi keperluan untuk ijazah Doktor Falsafah

**PENGOPTIMUMAN FORMULASI MEDIA DAN MEMPERTINGKATKAN  
SKALA PENGHASILAN TREONIN DAN TRIPTOFAN OLEH BAKTERIA ASID  
LAKTIK DENGAN MENGGUNAKAN KAEDAH RANGSANGAN PERMUKAAN**

Oleh

**LIM YE HENG**

**Disember 2017**

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Peningkatan pengetahuan terhadap fungsi asid amino dalam pengeluaran haiwan membawa kepada peningkatan permintaan pelbagai asid amino. Treonin dan triptofan merupakan asid amino yang paling biasa digunakan disebabkan peranan penting mereka dalam menambah baik prestasi pertumbuhan haiwan ternakan. Pada masa kini, penghasilan asid amino bergantung kepada mikroorganisma bukan gred makanan seperti *Corynebacterium glutamicum* serta *Escherichia coli* yang diubahsuai secara genetik dan ini merupakan satu kebimbangan oleh kerana penggunaan *C. glutamicum* yang diubahsuai secara genetik untuk penghasilan asid amino telah dikaitkan dengan lebih daripada ribuan kes sindrom maut iaitu sindrom eosinophila myalgia (EMS). Justeru, alternatif yang lebih selamat perlu dikaji. Baru-baru ini, terdapat laporan bahawa bakteria asid laktik (LAB) mampu menghasilkan pelbagai asid amino berikutan sistem proteolitik yang mantap serta kehadiran gen biosintesis asid amino. Tambahan pula, LAB dikenali dengan status umumnya yang diiktiraf sebagai selamat (GRAS), menjadikan mereka calon yang unggul sebagai penghasil asid amino gred makanan. Namun demikian, kajian berkaitan penghasilan asid amino dengan menggunakan LAB masih terhad. Oleh itu, objektif kajian ini adalah untuk mengenal pasti LAB yang boleh menghasilkan treonin dan triptofan serta mengoptimumkan rumusan media melalui kaedah rangsangan permukaan (RSM), disusuli dengan meningkatkan skala penghasilan dengan menggunakan kaedah kelajuan tip pendesak tetap. Hipotesis kajian ini adalah LAB yang mampu menghasilkan treonin dan triptofan dapat dikenal pasti dan penghasilannya dapat dipertingkatkan dengan pengoptimuman formulasi media dengan menggunakan RSM. Selain itu, penghasiln treonin dan triptofan oleh LAB terpilih boleh dipertingkatkan skala berdasarkan kaedah kelajuan tip pendesak tetap. Dalam kajian ini, 17 LAB yang pencilan daripada makanan Malaysia telah dikenal pasti secara fenotipik dan genotipik. LAB yang diisolasikan terdiri daripada 3 spesies: *Pediococcus pentosaceus* (6 strain), *Pediococcus acidilactici* (2 strain), dan

*Lactobacillus plantarum* (9 strain). Selepas itu, ciri profil pertumbuhan LAB dan aktiviti proteolitik mereka telah ditentukan secara kualitatif dan kuantitatif bawah 3 keadaan pH dengan menggunakan asai hidrolisis agar susu skim, kaedah penyebaran perigi agar susu skim serta kaedah azokasein. Semua LAB menunjukkan sistem proteolitik ekstrasel yang serba boleh di mana aktiviti proteolitik telah dikesan pada julat pH yang luas. Aktiviti proteolitik tertinggi ekstrasel pada pH 5 (15.76 U/mg) dan pH 8 (19.42 U/mg) dicatatkan oleh *L. plantarum* RG14. Sementara itu, *L. plantarum* RS5 dan RI11 menunjukkan aktiviti proteolitik ekstrasel tertinggi sekitar 17 U/mg pada pH 6.5.

Keupayaan LAB yang dipencilkan untuk menghasilkan asid amino kemudiannya ditentukan dengan pengkulturan dalam media de Man Rogosa dan Sharpe (MRS). Penghasilan asid amino diukur dengan menggunakan analisis kromatografi cecair berprestasi tinggi (HPLC). LAB yang dipencilkan menunjukkan keupayaan untuk menghasilkan pelbagai asid amino perindustrian penting namun penghasilan adalah bergantung pada strain. *P. pentosaceus* TL-3 menunjukkan produktiviti tertinggi treonin (12.88 mg/L/h) dan terpilih sebagai penghasil treonin unggul. Sementara itu, *P. acidilactici* TP-6 telah dipilih sebagai penghasil triptofan dengan produktiviti sebanyak 5.05 mg/L/h. Penghasilan treonin dan triptofan oleh LAB yang terpilih dioptimumkan seterusnya dengan menggunakan RSM. Pada mulanya, keperluan nutrien untuk penghasilan treonin dan triptofan dinilai dengan menggunakan reka bentuk *Plackett Burman* (PBD) dan kemudiannya dioptimumkan dengan menggunakan reka bentuk central composite (CCD). *Molasses*, ekstrak daging,  $(\text{NH}_4)_2\text{SO}_4$  dan  $\text{MnSO}_4$  adalah komponen yang paling penting untuk penghasilan treonin oleh *P. pentosaceus* TL-3 dengan kepekatan optimum sebanyak 30.79 g/L, 25.30 g/L, 8.59 g/L, dan 0.098 g/L masing-masing. Hasil treonin yang dicatatkan oleh *P. pentosaceus* TL-3 di bawah keadaan optimum (125.98 mg/L) telah meningkat sebanyak 2 kali ganda manakala kos media optimum telah dikurangkan sebanyak 8.5% berbanding dengan media MRS.

Di samping itu, kombinasi komponen media yang terbaik untuk penghasilan triptofan oleh *P. acidilactici* TP-6 adalah *molasses*, ekstrak daging, urea dan  $\text{FeSO}_4$ . Kepekatan optimum yang disyorkan oleh RSM adalah: ceng, 14.06 g/L; ekstrak daging, 23.68 g/L; urea, 5.56 g/L dan  $\text{FeSO}_4$ , 0.024 g/L. Sebanyak 68.05 mg/L triptofan telah dihasilkan oleh *P. acidilactici* TP-6 di bawah keadaan optimum, bersamaan dengan 150% peningkatan berbanding dengan kawalan. Sebaliknya, kos media optimum telah dikurangkan sebanyak 11%. Selain itu, penghasilan treonin dan triptofan oleh LAB terpilih telah berjaya dipertingkatkan skala dalam bioreaktor 30 L berdasarkan kaedah kelajuan tip pendesak tetap. Tambahan pula, hasil treonin dan triptofan yang dicatatkan di dalam pengkulturan bioreaktor adalah setanding dengan hasil yang diramalkan oleh RSM. Sebagai kesimpulan, *P. pentosaceus* TL-3 dan *P. acidilactici* TP-6 telah dikenal pasti sebagai penghasil treonin dan triptofan serta penghasilan treonin dan triptofan oleh penghasil terpilih telah dipertingkatkan sebanyak 2 kali ganda dan 150% masing-masing melalui pengoptimuman rumusan media mereka dengan menggunakan kaedah RSM. Tambahan pula, penghasilan treonin dan triptofan telah berjaya dipertingkatkan skala berdasarkan kaedah kelajuan

pendesak tetap di mana penghasilan di dalam skala kecil dan besar adalah setara.





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This thesis was submitted to the Senate of Universiti Putra Malaysia and has been accepted as fulfilment of the requirement for the degree of Doctor of Philosophy. The members of the Supervisory Committee were as follows:

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## LIST OF ABBREVIATIONS

% (v/v)	Percent volume per volume
% (w/v)	Percent weight per volume
(NH <sub>4</sub> ) <sub>2</sub> HC <sub>6</sub> H <sub>5</sub> O <sub>7</sub>	Ammonium citrate
(NH <sub>4</sub> ) <sub>2</sub> SO <sub>4</sub>	Ammonium sulphate
x g	Times gravity
° C	Degree in Celcius
µL	Microliter
2-(OH)-5-(NO <sub>2</sub> )C <sub>6</sub> H <sub>3</sub> COOH	2-hydroxy-5-nitrobenzoic acid
A230	Absorbance at wavelength 230 nm
A260	Absorbance at wavelength 260 nm
A280	Absorbance at wavelength 280 nm
AA	Amino acid
AAA	Auxiliary amino acid
Abs <sub>540nm</sub>	Absorbance at wavelength 540 nm
Ala	Alanine
ANOVA	Analysis of variance
Arg	Arginine
Asn	Asparagine
Asp	Aspartate
BLAST	Basic local alignment search tool
bp	Base pair
BSA	Bovine serum albumin
C.	<i>Corynebacterium</i>
C <sub>4</sub> H <sub>4</sub> KNaO <sub>6</sub> ·4H <sub>2</sub> O	Sodium tartrate tetrahydrate
C <sub>6</sub> H <sub>5</sub> OH	Phenol
CCD	Central composite design
CEP	Cell-envelope proteinase
CFS	Cell free supernatant
CFU/mL	Colony forming unit per millilitre
CuSO <sub>4</sub>	Copper (II) sulphate
Cy2	Cystine
DAD	Diode array detector
df	Degree of freedom
DNA	Deoxyribonucleic acid
DNS	Dinitrosalicylic acid
dNTPs	Deoxynucleotide triphosphates
Dpp	ATP-driven peptide transporter
DtpT	Di-/tripeptide transporter
E.	<i>Escherichia</i>
EBT	1,1'-ethylidene-bis-tryptophan
Em	Emission
EMP	Embden-Meyerhoff-Parnas
EMS	Eosinophils myalgia syndrome
Eq.	Equation
Ex	Excitation
FeSO <sub>4</sub>	Iron (II) sulphate

FLD	Fluorescence detector
FMOC	9-fluorenylmethyl chloroformate
g	Gram
GABA	Gamma-aminobutyric acid
Gln	Glutamine
Glu	Glutamate
Gly	Glycine
GRAS	Generally recognised as safe
h	Hour
HCl	Hydrochloric acid
His	Histidine
HIV	Human immunodeficiency virus
HPLC	High performance liquid chromatograph
Ile	Isoleucine
ISTD	Internal standard
K <sub>2</sub> HPO <sub>4</sub>	Dipotassium hydrogen phosphate
kb	Kilo base
KH <sub>2</sub> PO <sub>4</sub>	Potassium dihydrogen phosphate
k <sub>L</sub> a	Volumetric mass transfer coefficient
L	Litre
<i>L.</i>	<i>Lactobacillus</i>
LAB	Lactic acid bacteria
LC	Liquid chromatography
Leu	Leucine
LNAA	Large neutral amino acid
Lys	Lysine
M	Molar
Met	Methionine
MgCl <sub>2</sub>	Magnesium chloride
MgSO <sub>4</sub>	Magnesium sulphate
min	Minute
mL	Millilitre
mM	Millimolar
MnSO <sub>4</sub>	Manganese sulphate
MRS	de Man Rogasa and Sharpe
MSG	Monosodium glutamate
Na <sub>2</sub> SO <sub>3</sub>	Sodium sulfite
NaCl	Sodium chloride
NADP <sup>+</sup>	Nicotinamide adenine dinucleotide phosphate
NaH <sub>2</sub> PO <sub>4</sub>	Sodium dihydrogen phosphate
NaOAc	Sodium acetate
NaOH	Sodium hydroxide
NH <sub>4</sub> NO <sub>3</sub>	Ammonium nitrate
nm	Nanometre
NMF	Natural moisturising factors
OD <sub>600nm</sub>	Optical density at wavelength 600 nm
OPA	O-phthalaldehyde
Opp	Oligopeptides transporters
<i>P.</i>	<i>Pediococcus</i>
PBD	Plackett burman design
PCR	Polymerase chain reaction

Phe	Phenylalanine
PKP	Phosphoketolase pathway
PKU	Phenylketonuria
PMF	Protein motive force
pmol/ $\mu$ L	Picomole per microliter
$P_r$	Productivity
Pro	Proline
QPS	Qualified presumption of safety
$R^2$	Coefficient of determination
RC	Regenerated cellulose
rDNA	Ribosomal deoxyribonucleic acid
RNA	Ribonucleic acid
rpm	Rotation per minutes
RSM	Response surface methodology
SEM	Standard error mean
Ser	Serine
$S_f$	Final substrate concentration
$S_i$	Initial substrate concentration
$S_i-S_f$	Substrate consumed
SmF	Submerged fermentation
sp.	Species
SSF	Solid state fermentation
Std	Amino acid standard
Thr	Threonine
Trp	Tryptophan
Tyr	Tyrosine
UV	Ultraviolet
V	Volt
Val	Valine
$X_{max}$	Maximum cell concentration
$Y_{p/s}$	Product yield coefficient
$Y_{x/s}$	Growth yield coefficient
ZnSO <sub>4</sub>	Zinc sulphate

## CHAPTER 1

### INTRODUCTION

Amino acid is one of the most crucial nutrients to ensure survival of living organisms (Sundrum *et al.*, 2005). Increasing knowledge on the functions and properties of amino acid has led to diverse commercial applications and escalating global demand. Livestock industry is the largest consumer of amino acid and constitutes up to 56% of the total amino acid market, followed by food industry (32%) and other industries (12%) (Leuchtenberger *et al.*, 2005). Intensive development has favoured over feed amino acid due to their indispensable roles in enhancing the growth performance and well-being of livestock. Methionine and lysine were the first amino acid introduced to animal feed (Toride, 2004). However, threonine and tryptophan has gained increasing interest and applied widely in animal feed recently as they have positive impact on the growth performance of the animals (Xie *et al.*, 2014; Iwuji *et al.*, 2014; Duarte *et al.*, 2013; Świąch *et al.*, 2011).

Over the past decades, production of amino acid relied heavily on modified strains of *Corynebacterium glutamicum* and *Escherichia coli*. However, the use of genetically engineered microorganisms for production of tryptophan has caused over thousand cases of eosinophilia myalgia syndrome (EMS) due to production of a dimerisation product of tryptophan which is toxic. The tragedy eventually led to death of 27 victims while some of the victims experienced permanent disability. Hence, industries may still come to reluctant when using these microorganisms for production of amino acid, particularly in food industry. Moreover, use of pathogenic producer microorganisms such as *E. coli* are also raising concern (Venkitanarayanan *et al.*, 2016). This has motivated researchers to seek for safer alternatives by utilising food grade microorganisms such as lactic acid bacteria (LAB) for production of amino acid (Norfarina *et al.*, 2014; Zareian *et al.*, 2012).

LAB are one of the most important group of industrial microorganism. They are applied extensively for production of various cultured foods due to their Generally Recognised as Safe (GRAS) reputation (Blair & Regenstein, 2015; Mayo *et al.*, 2010) and ability to produce an array of metabolites that contribute greatly to improve the preservation properties and develop unique flavours and texture of the cultured foods (Margaret & Milind, 2006). Several studies revealed that LAB possessed a well-established proteolytic system which may contribute to production of amino acid (Savijoki *et al.*, 2006; Simova *et al.*, 2006; Hasan, 2003). Additionally, presence of active amino acid biosynthesis pathway and the relevant genes had been reported in LAB (Zareian *et al.*, 2012; Garault *et al.*, 2000).

Optimisation of fermentation parameters is a crucial step in bioprocess due to its impact on the product yield (Panda *et al.*, 2007) as well as influence on the economic feasibility of the fermentation process (Schmidt, 2005). Statistical

optimisation approaches are preferred over conventional optimisation approach due to its usefulness in optimising the process with minimal number of experiments. Response surface methodology (RSM) is one of the most commonly used statistical optimisation technique due to its high efficiency in determining the optimal conditions of a multivariate system and ability to explain the interactive effects of all the factors in a process with minimal experimental runs (Mander *et al.*, 2013; Elibol, 2004; Liu & Tzeng, 1998). RSM has been applied extensively for optimisation of various fermentation processes involving LAB (Deepak *et al.*, 2015; Tajabadi *et al.*, 2015; Anvari *et al.*, 2014; Hwang *et al.*, 2012; Kumar *et al.*, 2012; de Lima *et al.*, 2010; Li *et al.*, 2010; Norfarina, 2010). Another important aspect in bioprocess is the scalability of the process. The bioprocess must be able to be scaled up for industrial application. Constant impeller tip speed approach is one of the most commonly employed strategy for scaling up of bioprocesses involving microorganisms and enzymes (Keng *et al.*, 2008; Stephenie *et al.*, 2007; Hamdi *et al.*, 2000).

However, the major challenges to utilise LAB as a safer alternative for production of amino acid were lack of documentations regarding the ability of LAB to produce amino acid as well as the optimisation and scaling up of their production. Hence, the general objective of this study was to optimise the medium formulation for threonine and tryptophan production by using LAB isolated from Malaysian foods, followed by scaling up in pilot scale bioreactor. Meanwhile, the specific objectives of this study were as follows:

- 1) To identify and characterise the growth and proteolytic activity of LAB isolated from various Malaysian traditional foods and select the LAB isolate for threonine and tryptophan production.
- 2) To optimise the medium formulation for production of threonine and tryptophan by *Pediococcus pentosaceus* TL-3 and *Pediococcus acidilactici* TP-6 respectively via RSM approach.
- 3) To scale up the production of threonine and tryptophan by the *Pediococcus pentosaceus* TL-3 and *Pediococcus acidilactici* TP-6 by using constant impeller tip speed approach.

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