#### REFERENCES

- Ana-Rita, F. D. and Ian, W. D. (1996) Pyrolysis of sugar cane bagasse in a wire mesh reactor. <u>Industrial & Engineering Chemistry Research</u>, 35(4), 1263– 1268.
- Andren, R.K., Mandels, M., and Modeiros, J. E. (1976) Production of sugars from waste cellulose by by enzymatic hydrolysis: primary evaluation of substrates. <u>Process Biochemistry</u>, 11(4), 2-11.
- Chen, M., Xiaa, L., and Xueb, P. (2007) Enzymatic hydrolysis of corncob and ethanol production from cellulosic hydrolysate. <u>International</u> <u>Biodeterioration and Biodegradation</u>, 59, 85–89.
- Chunmei, P., Shufang, Z., Yaoting, F., and Hongwei, H. (2010) Bioconversion of corncob to hydrogen using anaerobic mixed Microflora. <u>Internation of Hydrogen Energy</u>, 35, 2663–2669.
- Dasari, R., Eric, K., Berson, R., and Mielenz, J. R. (2007) The Effect of Particle Size on Hydrolysis Reaction Rates and Rheological Properties in Cellulosic Slurries. <u>Applied biochemistry and biotechnology</u>, 137, 289-299.
- Dimian, A.C. and Bildea, C.S. (2008) <u>Chemical Process Design: Computer-Aided</u> <u>Case Studies</u>. Wiley-VCH.
- Galbe, M. and Zacchi, G. (2002). A review of the production of ethanol from softwood. Application Microbiology and Biotechnology, 59(6), 618-628.
- Galbe, M. and Zacchi, G. (2007) Pretreatment of lignocellulosic materials for efficient bioethanol production. <u>Advanced Biochemical</u> <u>Engineering/Biotechnology</u>, 108, 41–65.
- Kumar, P., Barrett, D.M., Delwiche, M.J., and Stroeve, P. (2009) Methods for pretreatment of lignocellulosic biomass for efficient hydrolysis and biofuel production. <u>Industrial & Engineering Chemistry Research</u>, 48(8), 3713– 3729.
- Lee, D., Yu, A.H.C., Wong, K.K.Y., and Saddler, J.R. (1994) Evaluation of the enzymatic susceptibility of cellulosic substrates using specific hydrolysis rates and enzyme adsorption. <u>Application Biochemical and Biotechnology</u>, 45(45), 407–415.

- Li, L., FrÖhlich, J., Pfeiffer, P., and KÖnig, H. (2003) Termite gut symbiotic archaezoa are becoming living metabolic fossils. <u>Eukaryot Cell</u>, 2, 1091–1098.
- Lin, L., Yan, R., Liu, Y., and Jiang, W. (2010) In-depth investigation of enzymatic hydrolysis of biomass wastes based on three major components: Cellulose, hemicelluloses and lignin. <u>Bioresource Technology</u>, 101, 8217-8223.
- Mousdale, D.M. (2008) <u>Biofuels : Biotechnology, Chemistry, and Sustainable</u> <u>Development</u>. New York: Taylor and Francis Group.
- Okano, K., Kitagaw, M., Sasaki, Y., and Watanabe, T. (2005) Conversion of Japanese red cedar (*Cryptomeria japonica*) into a feed for ruminants by white-rot basidiomycetes. <u>Animal Feed Science and Technology</u>, 120, 235– 243.
- Rivas, B., Torre, P., Domnguez, J.M., Perego, P., Converti, A., and Paraj, J.C. (2003) Carbon material and bioenergetic balances of xylitol production from corncobs by *Debaryomyces hansenii*. <u>Biotechnology Progress</u>, 19(3), 706–713.
- Saha, B. C. (2003) Hemicellulose bioconversion. <u>Industrial Microbiology and</u> <u>Biotechnology</u>, 30, 279-291.
- Sorahi, A., Keikhosro, K., Morteza, Khanahmadi., and Mohammad, J.T. (2009) Ethanol production by *Mucor indicus* and *Rhizopus oryzae* from rice straw by separate hydrolysis and fermentation. <u>Biomass and Bioenergy</u>, 33(5), 828-833.
- Sun, Y. and Cheng, J. (2002) Hydrolysis of lignocellulosic materials for ethanol production: a review. <u>Bioresource Technology</u>, 83, 1-11.
- Swatloski, R.P., Roger, R.D., and Holbrey, J.D. (2002) Dissolution and processing of cellulose using ionic liquids. <u>US Patent 6,824,599</u>, September 27.
- Taherzadeh, M.J. and Karimi, K. (2007a) Acid-based hydrolysis process for ethanol from ligrocellulosic material. <u>Bioresources Technology</u>, 2(3), 472-499.
- Taherzadeh, M.J. and Karimi, K. (2007b) Enzyme-based hydrolysis processes for ethanol from lignocellulosic Material. <u>Bioresources Technology</u>, 2(4), 707-738.

- Taechapoempol. K., Sreethawong, T., Rangsunvigit, P., Namprohm, W., Thamprajamchit, B., Rengpipat, S., and Chavadej, S. (2011) Cellulase producing bacteria from Thai higher termites, *Microcerotermes* sp.: enzymatic activities and ionic liquid tolerance. <u>Applied Biochemistry and Biotechnology</u>, 164(2), 204-19.
- Torres, B.R., Aliakbariana, B., Torrea, P., Peregoa, P., Doninguezb J.M., Zilli, M., and Converti, A. (2009) Vanillin bioproduction from alkaline hydrolyzate of corn cob by *Escherichia coli* JM109/pBB1. <u>Enzyme\_and\_Microbial</u> <u>Technology</u>, 44, 154–158.
- Yeh, A., Huang Y., and Chen, S.H. (2010) Effect of particle size on the rate of enzymatic hydrolysis of cellulose. <u>Carbohydrate Polymers</u>, 79(1), 192-199.
- Yoon, H.H. (1998) Pretreatment of lignocellulosic biomass by autohydrolysis and aqueous ammonia percolation. <u>Korean Journal of Chemical Engineering</u>, 15(6), 631-636.
- Werner, C., "Cellulosic Ethanol State-of-the-Art Conversion Processes" <u>Environmental and Energy Study Institute</u>. Acessed on 8 Jan. 2006. <a href="http://www.ef.org/documents/ce\_conversion\_factsheet\_ef\_eesi\_final\_1-08-07.pdf">http://www.ef.org/documents/ce\_conversion\_factsheet\_ef\_eesi\_final\_1-08-07.pdf</a>>.
- Worasamutprakarn, C. (2010) <u>Conversion of Cellulose to Glucose by Microbes</u> <u>Isolated from Higher Termites</u>. M.S. Thesis, The Petroleum and Petrochemical College, Chulalongkorn University.

Ł

## **APPENDICES**

## Appendix A Standard Calibration Curve

1. Glucose Calibration Curve

• • •

• :





| Table A1 Gluc | ose calibration curve |
|---------------|-----------------------|
|---------------|-----------------------|

ŧ

| Glucose concentration (g/L) | area(glucose) |
|-----------------------------|---------------|
| 0                           | 0             |
| 1.25                        | 798652        |
| 2.5                         | 1738632       |
| 5                           | 3808316       |
| 10                          | 7439522       |

## Appendix B Media for Microorganisms

| 1. 65 modified DSMZ broth medium 2 |      |   |  |
|------------------------------------|------|---|--|
| Approximate Formula* Per Liter     |      |   |  |
| Carboxymethyl Cellulose (CMC)      | 5.0  | g |  |
| Yeast extract                      | 4.0  | g |  |
| Malt extract                       | 10.0 | g |  |
|                                    |      |   |  |

Dissolve and adjust pH to 7.2

Autoclave at 121°C and pressure at 15 pounds/square inch for 15 minutes

# 2. 65 modified DSMZ agar medium 2

| Approximate Formula* Per Liter |                     |      |   |
|--------------------------------|---------------------|------|---|
| Carboxyme                      | hyl cellulose (CMC) | 5.0  | g |
| Yeast extrac                   | it .                | 4.0  | g |
| Malt extract                   |                     | 10.0 | g |
| Agar                           |                     | 12.0 | g |
|                                |                     |      |   |

Dissolve and adjust pH to 7.2

4

ł

Autoclave at 121°C and pressure at 15 pounds/square inch for 15 minutes

# Appendix C Reagent Preparations

4

| 1. 0.85%(w/v) NaCl in 1000 mL                        |       |    |
|--|-------|----|
| Sodium chloride (NaCl)                               | 8.5   | g  |
| Distilled water                                      | 1000  | mL |
| 2. Hydrochloric acid 1 N in 100 mL                   |       |    |
| Hydrochloric acid )HCl conc.)                        | 8.29  | mL |
| Distilled water                                      | 91.71 | mL |
|  | •     |    |
| 3. Sodium hydroxide 0.5 N in 1000 mL                 |       |    |
| Sodium hydroxide (NaOH)                              | 5.0   | g  |
| Distilled water                                      | 1000  | mL |
|  |       |    |
| 4. Sulfuric acid 0.72 N in 1000 mL                   |       |    |
| Sulfuric acid (H <sub>2</sub> SO <sub>4</sub> conc.) | .72   | mL |
| Distilled water                                      | .28   | mL |
|  |       |    |

#### Appendix D Bacteria Concentration

Bacteria concentration was determined using total nitrogen test kit.

1. The bacteria concentration from enzymatic hydrolysis

During enzymatic hydrolysis, bacteria growth was monitored by withdrawing samples from the hydrolysis reactor periodically. Solid that obtained from centrifuging of the sample, contained of corncob and bacteria. Method that can calculate weight of bacteria and corncob is shown in equation D1.

wt. Solid = wt. Corncob + wt. Bacteria (D1)

Then, a concentration of bacteria was determined by the total nitrogen test kit.

#### 1.1 The amount of nitrogen in bacteria

ł

The amount of nitrogen in each strain was determined in triplicates by using the total nitrogen test kit. Figure F1 shows procedure for determination



Figure D1 Diagram for determination the amount of nitrogen in bacteria.

## Procedure

Nitrogen total persulfate digestion method is conducted in order to check amount of nitrogen which directly related to amount of bacteria during hydrolysis.



Figure D2 Procedure for analyzing amount of nitrogen.



Figure D2 Procedure for analyzing amount of nitrogen (continued).

# Appendix E Experiment Data of Enzymatic Hydrolysis

**Table E1** Glucose produced from the hydrolysis of 60 mesh particle size corncoband strain A 002 bacteria at 37 °C.

| Hr | Area   | Glucose<br>concentration<br>(wt%) | Glucose<br>concentration<br>(g/L) |
|----|--------|-----------------------------------|-----------------------------------|
| 0  | 229544 | 0.03086                           | 0.30862                           |
| 1  | 116684 | 0.01569                           | 0.15688                           |
| 2  | 52858  | 0.00711                           | 0.07107                           |
| 3  | 22666  | 0.00305                           | 0.03047                           |
| 4  | 128946 | 0.01734                           | 0.17337                           |
| 5  | 247563 | 0.03328                           | 0.33285                           |
| 6  | 547499 | 0.07361                           | 0.73612                           |
| 7  | 641769 | 0.08629                           | 0.86286                           |
| 8  | 675620 | 0.09084                           | 0.90837                           |
| 9  | 804021 | 0.10810                           | 1.08101                           |
| 10 | 762416 | 0.10251                           | 1.02507                           |
| 11 | 732903 | 0.09854                           | 0.98539                           |
| 12 | 519497 | 0.06985                           | 0.69847                           |
| 15 | 160586 | 0.02159                           | 0.21591                           |
| 18 | 52666  | 0.00708                           | 0.07081                           |
| 24 | 50023  | 0.00673                           | 0.06726                           |

4

**Table E2** Glucose produced from the hydrolysis of 40 mesh particle size corncoband strain A 002 bacteria at 37 °C.

| Hr | Area   | Glucose<br>concentration<br>(wt%) | Glucose<br>concentration<br>(g/L) |
|----|--------|-----------------------------------|-----------------------------------|
| 0  | 229544 | 0.03086                           | 0.30862                           |
| 1  | 36916  | 0.00496                           | 0.04963                           |
| 2  | 103369 | 0.01390                           | 0.13898                           |
| 3  | 70140  | 0.00943                           | 0.09430                           |
| 4  | 23722  | 0.00319                           | 0.03189                           |
| 5  | 233477 | 0.03139                           | 0.31391                           |
| 6  | 185312 | 0.02492                           | 0.24915                           |
| 7  | 312634 | 0.04203                           | 0.42034                           |
| 8  | 451986 | 0.06077                           | 0.60770                           |
| 9  | 725897 | 0.09760                           | 0.97597                           |
| 10 | 683000 | 0.09183                           | 0.91830                           |
| 11 | 690582 | 0.09285                           | 0.92849                           |
| 12 | 575000 | 0.07731                           | 0.77309                           |
| 15 | 554778 | 0.07459                           | 0.74590                           |
| 18 | 456580 | 0.06139                           | 0.61387                           |
| 24 | 284908 | 0.03831                           | 0.38306                           |

--

| Hr  | Area   | Glucose<br>concentration<br>(wt%) | Glucose<br>concentration<br>(g/L) |
|-----|--------|-----------------------------------|-----------------------------------|
| 0   | 229544 | 0.03086                           | 0.30862                           |
| 1   | 249085 | 0.03349                           | 0.33490                           |
| . 2 | 252909 | 0.03400                           | 0.34004                           |
| 3   | 260915 | 0.03508                           | 0.35080                           |
| . 4 | 263551 | 0.03543                           | 0.35435                           |
| 5   | 318393 | 0.04281                           | 0.42808                           |
| 6   | 468493 | 0.06299                           | 0.62989                           |
| .7  | 523020 | 0.07032                           | 0.70320                           |
| . 8 | 650002 | 0.08739                           | 0.87393                           |
| . 9 | 703599 | 0.09460                           | 0.94599                           |
| 10  | 577300 | 0.07762                           | 0.77618                           |
| -11 | 456188 | 0.06133                           | 0.61335                           |
| 12  | 478158 | 0.06429                           | 0.64289                           |
| 15  | 401121 | 0.05393                           | 0.53931                           |
| 18  | 254973 | 0.03428                           | 0.34281                           |
| 24  | 178520 | 0.02400                           | 0.24002                           |

**Table E3** Glucose produced from the hydrolysis of 60 mesh particle size corncoband strain A 002 bacteria at 30 °C.

| Hr | Area   | Glucose<br>concentration<br>(wt%) | Glucose<br>concentration<br>(g/L) |
|----|--------|-----------------------------------|-----------------------------------|
| 0  | 229544 | 0.03086                           | 0.30862                           |
| 1  | 172888 | 0.02324                           | 0.23245                           |
| 2  | 268048 | 0.03604                           | 0.36039                           |
| 3  | 298586 | 0.04015                           | 0.40145                           |
| 4  | 339069 | 0.04559                           | 0.45588                           |
| 5  | 353190 | 0.04749                           | 0.47487                           |
| 6  | 424803 | 0.05711                           | 0.57115                           |
| 7  | 505844 | 0.06801                           | 0.68011                           |
| 8  | 527788 | 0.07096                           | 0.70961                           |
| 9  | 662401 | 0.08906                           | 0.89060                           |
| 10 | 625263 | 0.08407                           | 0.84067                           |
| 11 | 595295 | 0.08004                           | 0.80038                           |
| 12 | 375203 | 0.05045                           | 0.50446                           |
| 15 | 375203 | 0.05045                           | 0.50446                           |
| 18 | 344951 | 0.04638                           | 0.46379                           |
| 24 | 298990 | 0.04020                           | 0.40199                           |

**Table E4** Glucose produced from the hydrolysis of 40 mesh particle size corncoband strain A 002 bacteria at 30 °C.

| Hr | Area   | Glucose<br>concentration<br>(wt%) | Glucose<br>concentration<br>(g/L) |
|----|--------|-----------------------------------|-----------------------------------|
| 0  | 229544 | 0.03086                           | 0.30862                           |
| 1  | 306312 | 0.04118                           | 0.41184                           |
| 2  | 218858 | 0.02943                           | 0.29426                           |
| 3  | 137435 | 0.01848                           | 0.18478                           |
| 4  | 272330 | 0.03661                           | 0.36615                           |
| 5  | 267331 | 0.03594                           | 0.35943                           |
| 6  | 290858 | 0.03911                           | 0.39106                           |
| 7  | 354695 | 0.04769                           | 0.47689                           |
| 8  | 369550 | 0.04969                           | 0.49686                           |
| 9  | 391898 | 0.05269                           | 0.52691                           |
| 10 | 231627 | 0.03114                           | 0.31142                           |
| 11 | 245620 | 0.03302                           | 0.33024                           |
| 12 | 258539 | 0.03476                           | 0.34761                           |
| 15 | 267514 | 0.03597                           | 0.35967                           |
| 18 | 196442 | 0.02641                           | 0.26412                           |
| 24 | 56888  | 0.00765                           | 0.07649                           |

ł

 
 Table E5
 Glucose produced from the hydrolysis of 60 mesh particle size corncob
and strain M 015 bacteria at 37 °C.

| Hr | Area   | Glucose<br>concentration<br>(wt%) | Glucose<br>concentration<br>(g/L) |
|----|--------|-----------------------------------|-----------------------------------|
| 0  | 229544 | 0.03086                           | 0.30862                           |
| 1  | 255125 | 0.03430                           | 0.34302                           |
| 2  | 176015 | 0.02367                           | 0.23665                           |
| 3  | 172123 | 0.02314                           | 0.23142                           |
| 4  | 211878 | 0.02849                           | 0.28487                           |
| 5  | 322648 | 0.04338                           | 0.43380                           |
| 6  | 328453 | 0.04416                           | 0.44161                           |
| 7  | 328017 | 0.04410                           | 0.44102                           |
| 8  | 359922 | 0.04839                           | 0.48392                           |
| 9  | 365885 | 0.04919                           | 0.49193                           |
| 10 | 331961 | 0.04463                           | 0.44632                           |
| 11 | 325514 | 0.04377                           | 0.43766                           |
| 12 | 300236 | 0.04037                           | 0.40367                           |
| 15 | 226688 | 0.03048                           | 0.30478                           |
| 18 | 163065 | 0.02192                           | 0.21924                           |
| 24 | 135891 | 0.01827                           | 0.18271                           |

Ł

**Table E6** Glucose produced from the hydrolysis of 40 mesh particle size corncoband strain M 015 bacteria at 37 °C.

. . . . . .

| Hr | Area   | Glucose<br>concentration<br>(wt%) | Glucose<br>concentration<br>(g/L) |
|----|--------|-----------------------------------|-----------------------------------|
| 0  | 229544 | 0.03086                           | 0.30862                           |
| 1  | 58330  | 0.00784                           | 0.07842                           |
| 2  | 59856  | 0.00805                           | 0.08048                           |
| 3  | 60012  | 0.00807                           | 0.08069                           |
| 4  | 61792  | 0.00831                           | 0.08308                           |
| 5  | 88924  | 0.01196                           | 0.11956                           |
| 6  | 95865  | 0.01289                           | 0.12889                           |
| 7  | 70140  | 0.00943                           | 0.09430                           |
| 8  | 84075  | 0.01130                           | 0.11304                           |
| 9  | 225050 | 0.03026                           | 0.30258                           |
| 10 | 185312 | 0.02492                           | 0.24915                           |
| 11 | 103369 | 0.01390                           | 0.13898                           |
| 12 | 67168  | 0.00903                           | 0.09031                           |
| 13 | 23336  | 0.00314                           | 0.03138                           |
| 14 | 50680  | 0.00681                           | 0.06814                           |
| 15 | 25756  | 0.00346                           | 0.03463                           |
| 16 | 29520  | 0.00397                           | 0.03969                           |
| 18 | 18965  | 0.00255                           | 0.02550                           |
| 24 | 13334  | 0.00179                           | 0.01793                           |

Table E7Glucose produced from the hydrolysis of 60 mesh particle size corncoband strain M 015 bacteria at 30 °C.

| Hr          | Area   | Glucose<br>concentration<br>(wt%) | Glucose<br>concentration<br>(g/L) |
|-------------|--------|-----------------------------------|-----------------------------------|
| 0           | 229544 | 0.03086                           | 0.30862                           |
| $1^{\circ}$ | 11000  | 0.00148                           | 0.01479                           |
| 2 -         | 13560  | 0.00182                           | 0.01823                           |
| 3           | 14520  | 0.00195                           | 0.01952                           |
| 4 =         | 15349  | 0.00206                           | 0.02064                           |
| 5           | 15902  | 0.00214                           | 0.02138                           |
| 6           | 17390  | 0.00234                           | 0.02338                           |
| 7           | 70207  | 0.00944                           | 0.09439                           |
| 8           | 64243  | 0.00864                           | 0.08638                           |
| 9           | 180628 | 0.02429                           | 0.24286                           |
| 10          | 145200 | 0.01952                           | 0.19522                           |
| 11          | 108328 | 0.01456                           | 0.14565                           |
| 12          | 104772 | 0.01409                           | 0.14087                           |
| 13          | 120690 | 0.01623                           | 0.16227                           |
| 14          | 120036 | 0.01614                           | 0.16139                           |
| 15          | 99958  | 0.01344                           | 0.13439                           |
| 16          | 78596  | 0.01057                           | 0.10567                           |
| 18          | 56000  | 0.00753                           | 0.07529                           |
| 24          | 52223  | 0.00702                           | 0.07021                           |

**Table E8** Glucose produced from the hydrolysis of 40 mesh particle size corncoband strain M 015 bacteria at 30 °C.

| Hr | Total bacterial<br>(g/L) | Solid(g) | Nitrogen<br>bacteria (g/L) | Bacterial<br>(g/L) |
|----|--------------------------|----------|----------------------------|--------------------|
| 0  | 10.3                     | 10.244   | 0.056                      | 0.496774           |
| 2  | 14.2                     | 14.124   | 0.076                      | 0.674194           |
| 3  | 23.1                     | 23       | 0.1                        | 0.887097           |
| 5  | 14.4                     | 14.278   | 0.122                      | 1.082258           |
| 7  | 14                       | 13.84    | 0.16                       | 1.419355           |
| 8  | 9.9                      | 9.708    | 0.192                      | 1.703226           |
| 9  | 10                       | 9.8      | 0.2                        | 1.774194           |
| 12 | 16.6                     | 16.34    | 0.26                       | 2.306452           |
| 18 | 24.4                     | 24.1     | 0.3                        | 2.66129            |
| 24 | 14.9                     | 14.582   | 0.318                      | 2.820968           |

**Table E9** Bacteria evolution from the enzymatic hydrolysis of 60 mesh size corncobwith strain A 002 bacteria at 37 °C.

**Table E10** Bacteria evolution from the enzymatic hydrolysis of 40 mesh sizecorncob with strain A 002 bacteria at 37 °C.

| Hr | Total bacterial<br>(g/L) | Solid(g) | Nitrogen<br>bacteria (g/L) | Bacterial<br>(g/L) |
|----|--------------------------|----------|----------------------------|--------------------|
| 0  | 6.2                      | 6.142    | 0.058                      | 0.514516           |
| 3  | 10.3                     | 10.23    | 0.07                       | 0.620968           |
| 5  | 11.5                     | 11.41    | 0.09                       | 0.798387           |
| 6  | 11.7                     | 11.592   | 0.108                      | 0.958065           |
| 9  | 7.2                      | 7.004    | 0.196                      | 1.73871            |
| 12 | 8                        | 7.798    | 0.202                      | 1.791935           |
| 18 | 7.7                      | 7.482    | 0.218                      | 1.933871           |
| 24 | 15.6                     | 15.404   | 0.196                      | 1.73871            |

Ł

| Hr | Total<br>bacterial (g/L) | Solid(g) | Nitrogen<br>bacteria (g/L) | Bacterial<br>(g/L) |
|----|--------------------------|----------|----------------------------|--------------------|
| 0  | 4.9                      | 4.825    | 0.075                      | 0.665323           |
| 2  | 10.3                     | 10.14    | 0.16                       | 1.419355           |
| 5  | 6                        | 5.824    | 0.176                      | 1.56129            |
| 7  | 8.8                      | 8.6      | 0.2                        | 1.774194           |
| 9  | 7.6                      | 7.36     | 0.24                       | 2.129032           |
| 12 | 9                        | 8.75     | 0.25                       | 2.217742           |
| 18 | 9.5                      | 9.252    | 0.248                      | 2.2                |
| 24 | 9.4                      | 9.16     | 0.24                       | 2.129032           |

**Table E11** Bacteria evolution from the enzymatic hydrolysis of 60 mesh sizecorncob with strain A 002 bacteria at 30 °C.

**Table E12** Bacteria evolution from the enzymatic hydrolysis of 40 mesh sizecorncob with strain A 002 bacteria at 30 °C.

| Hr | Total<br>bacterial<br>(g/L) | Solid(g) | Nitrogen<br>bacteria (g/L) | Bacterial<br>(g/L) |
|----|-----------------------------|----------|----------------------------|--------------------|
| 0  | 8.1                         | 7.998    | 0.102                      | 0.904839           |
| 3  | 10.4                        | 10.242   | 0.158                      | 1.401613           |
| 7  | 2.6                         | 2.466    | 0.134                      | 1.18871            |
| 9  | 5.7                         | 5.52     | 0.18                       | 1.596774           |
| 12 | 4                           | 3.832    | 0.168                      | 1.490323           |
| 14 | 5.7                         | 5.492    | 0.208                      | 1.845161           |
| 16 | 7.4                         | 7.186    | 0.214                      | 1.898387           |
| 18 | 4                           | 3.782    | 0.218                      | 1.933871           |
| 24 | 5.9                         | 5.7      | 0.2                        | 1.774194           |

-1

**Table E13** Bacteria evolution from the enzymatic hydrolysis of 60 mesh sizecorncob with strain M 015 bacteria at 37 °C.

| Hr | Total<br>bacterial (g/L) | Solid(g) | Nitrogen<br>bacteria (g/L) | Bacterial<br>(g/L) |
|----|--------------------------|----------|----------------------------|--------------------|
| 0  | 1.2                      | 1.131    | 0.069                      | 0.432997           |
| 3  | 4.4                      | 4.290    | 0.110                      | 0.694737           |
| 7  | .1                       | 0.820    | 0.180                      | 1.136842           |
| 10 | 2.7                      | 2.540    | 0.160                      | 1.010526           |
| 12 | 2.9                      | 2.706    | 0.194                      | 1.225263           |
| 16 | 3.1                      | 2.856    | 0.244                      | 1.541053           |
| 18 | 2.9                      | 2.656    | 0.244                      | 1.541053           |
| 24 | 8.3                      | 8.054    | 0.246                      | 1.553684           |
|    |                          |          |                            |                    |

**Table E14**Bacteria evolution from the enzymatic hydrolysis of 40 mesh sizecorncob with strain M 015 bacteria at 37 °C.

| Hr | Total<br>bacterial (g/L) | Solid(g) | Nitrogen<br>bacteria (g/L) | Bacterial<br>(g/L) |
|----|--------------------------|----------|----------------------------|--------------------|
| 0  | 2                        | 1.963    | 0.1                        | 0.631579           |
| 3  | 1.1                      | 0.962    | 0.138                      | 0.871579           |
| 4  | 4.6                      | 4.45     | 0.15                       | 0.947368           |
| 6  | 12                       | 11.742   | 0.258                      | 1.629474           |
| 7  | 11.6                     | 11.296   | 0.304                      | 1.92               |
| 9  | 30                       | 29.688   | 0.312                      | 1.970526           |
| 12 | 23.2                     | 22.88    | 0.32                       | 2.021053           |
| 18 | 11                       | 10.694   | 0.306                      | 1.932632           |
| 24 | 7.6                      | 7.286    | 0.314                      | 1.983158           |

Table E15Bacteria evolution from the enzymatic hydrolysis of 60 mesh sizecorncob with strain M 015 bacteria at 30 °C.

| Hr | Total<br>bacterial (g/L) | Solid(g) | Nitrogen<br>bacteria(g/L) | Bacterial<br>(g/L) |
|----|--------------------------|----------|---------------------------|--------------------|
| 0  | 13                       | 12.948   | 0.052                     | 0.328421           |
| 1  | 16.3                     | 16.14    | 0.16                      | 1.010526           |
| 3  | 53.7                     | 53.46    | 0.24                      | 1.515789           |
| 4  | 27.3                     | 27.054   | 0.246                     | 1.553684           |
| 9  | 13.7                     | 13.4     | 0.3                       | 1.894737           |
| 12 | 51.5                     | 51.25    | .0.25                     | 1.578947           |
| 18 | 28.2                     | 27.92    | 0.28                      | 1.768421           |
| 24 | 22.7                     | 22.428   | 0.272                     | 1.717895           |
|    |                          |          | 97 H                      |                    |

Table E16Bacteria evolution from the enzymatic hydrolysis of 40 mesh sizecorncob with strain M 015 bacteria at 30 °C.

| Hr | Total<br>bacterial (g/L) | Solid(g) | Nitrogen<br>bacteria(g/L) | Bacterial<br>(g/L) |
|----|--------------------------|----------|---------------------------|--------------------|
| 0  | 3.6                      | 3.582    | 0.018                     | 0.113684           |
| 1  | 34.9                     | 34.784   | 0.116                     | 0.732632           |
| 3  | 7.5                      | 7.196    | 0.304                     | 1.92               |
| 4  | 2.7                      | 2.388    | 0.312                     | 1.970526           |
| 9  | 27                       | 26.58    | 0.42                      | 2.652632           |
| 10 | 37.8                     | 37.358   | 0.442                     | 2.791579           |
| 18 | 45.2                     | 44.748   | 0.452                     | 2.854737           |
| 24 | 30.3                     | 29.856   | 0.444                     | 2.804211           |

-2

### **CURRICULUM VITAE**

| Name:          | Ms. Wannaporn Eourarekullart |
|----------------|------------------------------|
| Date of Birth: | December 29, 1986            |

Nationality: Thai

Ł

#### **University Education:**

2005-2008 Bachelor Degree of Engineering, Department of Chemical Engineering, Faculty of Engineering, Mahidol University, Nakhonpathom, Thailand Work Experience:

2008Position:Process Engineer (Student Internship)Company:Thai Synthetic Rubber Co.,Ltd

#### **Proceedings:**

 Eourarekullart, W., Rangsunvigit, P., Sreethawong, T., Chavadej, S., and Rengpipat, S., (2011, April 26) Conversion of Corncob to Sugars by Microbial Hydrolysis. <u>Proceedings of The 2<sup>nd</sup> Research Symposium on Petroleum,</u> <u>Petrochemicals, and Advanced Materials and The 17<sup>th</sup> PPC Symposium on Petroleum, Petrochemicals, and Polymers, Bangkok, Thailand.</u>

