

Cellulase Production by *Aspergillus niger* ATCC 16888 on Copra Waste from Coconut Milk Process In Layered Packed-bed Bioreactor



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Solid-state fermentation (SSF) has been used to produce value-added products from agricultural and food processing wastes, but the majority of SSF production has been conducted on a small scale due to the limitations of heat and mass transfer. This research aimed to reduce the temperature changes in packed-bed bioreactor during the cellulase fermentation of *Aspergillus niger* ATCC 16888 on copra waste from coconut milk process. The three configurations of bioreactor used in this study consisted of one- (control), two- and three-layered packed-bed bioreactors. Total volumes of packed-bed bioreactor and bed material were 127 and 57 L, respectively. The sterilized air 30 °C was forced from the bottom of bioreactor at 0.20 vvm. The highest cellulase production (5.0 ± 0.4 FPU g^{-1}_{DS}) was obtained from the bottom zone of the three-layered bioreactor after fermenting for 1 d. However, the greatest growth was found at the top one, in which a large amount of aerial mycelium was detected.

Keywords:

cellulase, solid-state fermentation, packed-bed bioreactor, *Aspergillus niger*

Introduction

Nowadays, solid-state fermentation (SSF) is used to produce value-added bio-products from agro-industrial and food wastes, which contain carbohydrates and other nutrients for microbial metabolisms¹. There are many researchers studying SSF for enzyme production; for instance, protease², peroxidase³, α -amylase⁴, etc. Cellulase is an enzyme that can be produced by SSF⁵. Although both bacteria and fungi can produce cellulase, the filamentous fungi *Aspergillus niger* and *Trichoderma reesei* have been mostly used for commercial cellulase production⁶. Recently, many microorganisms have been studied for cellulase production from different agricultural materials, such as soy residues⁷, babassu cake⁸, sugar cane bagasse⁹ and Palmyra palm¹⁰. Cerda *et al.*¹¹ indicated that the microbial diversity benefited the economics of cellulase fermentation using wastes as substrates. However, filamentous fungi are most suitable for SSF because they can grow and carry out their metabolisms in low moisture conditions¹².

Coconut, an economic crop of Thailand, has been used to produce many food products; for in-

stance, coconut juice and coconut milk. In the process of coconut milk, copra waste was obtained up to 30 %¹³. To add value of this waste, a solid-state fermentation (SSF) process is attractive due to low production cost. Since copra waste contains high content of cellulose ~ 72 %¹⁴, which is a good substrate for cellulose production by fungi¹⁵, this research focused on using copra waste as a solid substrate for a fungal cellulase production by SSF, because cellulolytic enzymes have been currently useful in many industries such as textile, paper, detergent, and energy industries^{16,17}. However, most researchers into SSF have conducted experiments on the laboratory scale, such as flasks^{2,18–20}, glass columns²¹, plate bioreactor²², bag bioreactor²³, etc. due to scale-up problems, especially the accumulation of heat generated by microbial respiration and other metabolisms²⁴. One of the bioreactor types which is easy and cheap to operate is the packed-bed bioreactor. However, the main problems in its application are the difficulties in parameter control, including temperature and moisture distribution within the bioreactor^{25,26} which is affected by porosity, bed height, and fluid velocity²⁷. Recently, Biz *et al.*²⁸ indicated that the porosity of the substrate bed was an important factor for a temperature control of the packed-bed column and the problem found in the pilot-scale packed-bed bioreactor was heat ac-

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cumulation due to a compaction of the bed during the SSF²⁹. Therefore, this work aimed to increase the heat removal from packed-bed bioreactor during the SSF of cellulase by *A. niger* by dividing the packed-bed into two and three layers, and determining its effect on the fungal growth and the cellulase production.

Materials and methods

Bioreactor construction

The 127-L packed-bed bioreactor (45 cm diameter and 80 cm height) was used to ferment the 57-L substrate in this research. The schematics of the packed-bed bioreactors used in this research are shown in Fig. 1. The aluminium screen baskets used as the substrate container were connected to the cylinder stainless steel frame. The diameter of the baskets for the control (CB), two-layered (2LB) and three-layered (3LB) bioreactors were 45 cm, but their heights were different – 36 cm (1 basket), 18 cm (2 baskets) and 12 cm (3 baskets), respectively. After filling the substrate in the baskets, they were assembled to the frame. The positions of the baskets were adjustable along the height of the frame by screws. The 45.5 cm diameter aluminium cylinder (1.5 mm thickness) with holes and stoppers was made to cover the side and act as a wall of the bioreactor, which was able to collect the fermented sample and measure the temperature during SSF. An aluminium lid punctured for air was then used to cover the bioreactor. The bioreactor was connected to a rotary air pump (LP-20, Resun, China) at the bottom of the bioreactors for forcing air. The air ve-

locity was controlled by gas rotameter (Model K-1011, Nitto Denko Corporation, Japan).

Fungal strain and inoculum preparation

The strain, *Aspergillus niger* ATCC 16888, was purchased from the American Type Culture Collection (ATCC) and treated according to the procedure provided by the ATCC. Its spore suspension was prepared by cultivating *A. niger* on PDA in 86-mm-diameter petri dish at 30 °C for 7 d, and collecting the spores using sterilized water. The suspension was filtrated using Whatman™ No.1 filter paper (GE Healthcare UK limited, China) under aseptic conditions. The concentration of spores in the suspension was adjusted to 10⁶ spores mL⁻¹ by adding sterilized water.

Solid substrate preparation

Copra waste obtained from a local market was dried at 80 °C for 6 h. It was packed in a closed container to prevent oxidations from air and light, and then kept at lower than 5 °C. An amount of 5 kg of substrate containing 60 % moisture was used for one batch of SSF. It was prepared by adding 2.8 kg distilled water into 2.0 kg dry copra waste, and autoclaved at 121 °C for 30 min, and cooled to room temperature in an aseptic area. The sterilized substrate was inoculated and mixed with 200 mL spore suspension (~0.2 kg, total spore number = 2 – 10⁸) to achieve the 60 % moisture content and 10⁵ spores g⁻¹ dry substrate using a 50-L horizontal rotary drum that was built to mix at 10 rpm for 10 min (~15–20 L/batch). The moisture content of substrate was measured using the moisture analyzer before the experiment.

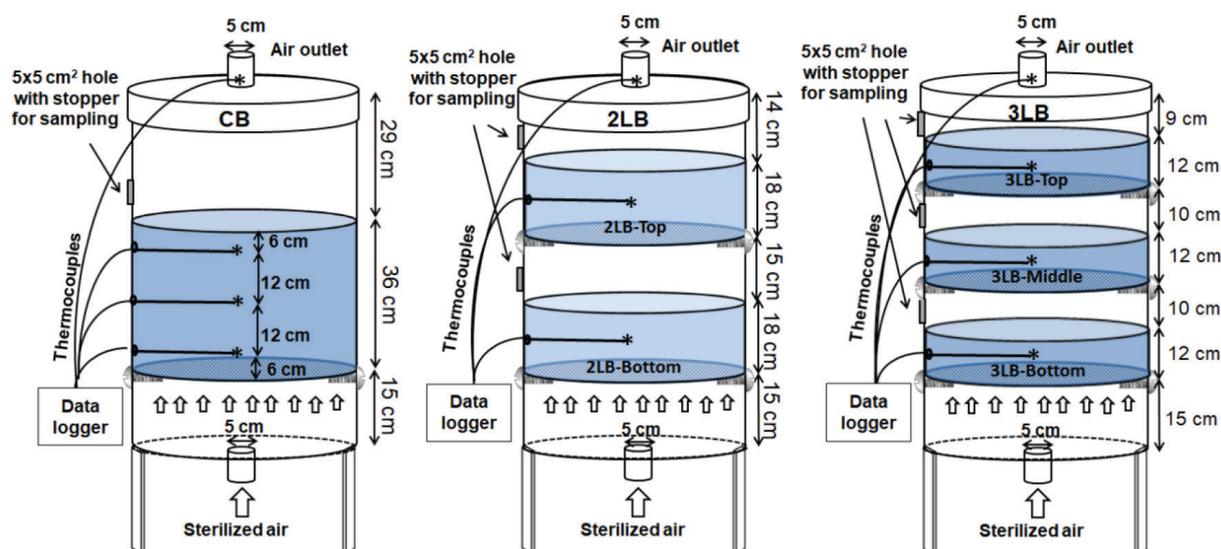


Fig. 1 – Schematics of packed-bed bioreactors used in this study (a) Control packed-bed bioreactor, CB; (b) Two-layered packed-bed bioreactor, 2LB; (c) Three-layered packed-bed bioreactor, 3LB; The * symbols represent the positions for temperature measurement

Solid-state fermentation

Five kg of prepared substrate mixed with spore suspension (total volume = 57 L) was filled in the studied packed-bed bioreactors consisting of the CB, 2LB, and 3LB. Each bed in the 2LB contained 2.5 kg or 28.5 L of the prepared substrate. The same total weight and volume of bed was applied in the 3LB, where each bed was filled with 1.7 kg or 19 L of the substrate. During 6 days of fermentation, the humidified air (30 °C) sterilized by the PTFE membrane (Millex®-FG, Millipore corporation, USA) was aerated from the bottom of bioreactor at a rate of 0.2 vvm or 11.4 L min⁻¹. The fermented sample was collected randomly (5 samples per bed) by puncturing the bed along the height of the bed using 1.0 cm diameter glass tube at intervals of 1 d to analyze moisture content by oven drying, cellulase activity by FPase assay, and fungal growth by glucosamine measurement. Each treatment was performed in triplicate. Temperatures of outlet air and beds in each layer were monitored during SSF using the thermocouples Type T (1.5 mm diameter) connected to the data logger (Model 34970A, Keysight Technologies, USA).

Enzyme extraction

Cellulase was extracted from the fermented sample according to the modified method of Ang *et al.*³⁰ About 4 g fermented sample was soaked in 100 mL sodium acetate buffer (pH 6.5, 0.05 M) at room temperature (~25 °C), and stirred for 30 min with a magnetic stirrer. After filtrating through Whatman™ No.1 filter paper, the filtrate was used for FPase assay.

Analysis

Glucosamine determination for fungal growth

About 5 g of fermented sample was dried at 80 °C for 6 h, then the amount of 10 mg of dried fermented sample was hydrolyzed with 60 % sulfuric acid solution, and the glucosamine content analyzed using a colorimetric method³¹ by measuring the absorbance of reaction solution at 650 nm with the UV-Visible spectrophotometer (Model Genesys 10S, Thermo Fisher Scientific, USA). The standard solution was prepared using HCl-glucosamine solution with different concentrations. The growth of *A. niger* was reported as a glucosamine content in dry fermented copra waste (mg g⁻¹_{DS}). The analyses were performed in triplicate. The values of glucosamine content were plotted against the fermentation time, and their values in the growth phase were used to calculate the specific growth rate using exponential equation as in Eq. 1.

$$X_t = X_0 e^{\mu t} \quad (1),$$

where X_t is the glucosamine content (mg g⁻¹ sample) at time t , X_0 is the glucosamine content (mg g⁻¹ sample) at the initial time ($t = 0$), and μ is the specific growth rate (d⁻¹) in the growth phase.

FPase assay

Cellulase activity (FPA) was analyzed according to Ghose³². The amount of 0.5 mL crude enzyme and 1.0 mL (0.05 M) citrate buffer, pH 5.0, were added to test tubes containing 5 mg of Whatman No.1 filter paper (6 x 1 cm²). The test tube was incubated in a water bath at 50 °C for 1 h. Then, 3 mL of 3, 5- dinitrosalicylic acid was added into the tube and boiled for 15 min to stop the reaction. After cooling to ambient temperature, its volume was filled to 10 mL by adding distilled water and the absorbance of the filtered solution measured at 540 nm wavelength against a blank of sugar-free sample using the UV-Visible spectrophotometer (Model Genesys 10S, Thermo Fisher Scientific, USA). Standard curve was conducted using a standard glucose solution with known concentration. One unit of enzyme activity (U) was the amount of enzyme that released one micromole of glucose in one minute. The FPase activity was reported as the unit of enzyme in one gram of dry substrate (FPU g⁻¹_{DS}).

Results and discussion

Temperature distribution in packed-bed bioreactor

During the SSF, the temperatures of each bed in the studied bioreactors were measured as shown in Fig. 2. The bed temperature in the CB increased from 30.0 ± 0.5 °C at the beginning to the maximum of 52.0 ± 3.0 °C at day 1 and slowly decreased to 38.2 ± 1.6 °C at day 6. The increasing temperature also occurred in the study of Pitol *et al.*²⁴, who found that the bed temperature increased up to 47 °C from 32 °C during the SSF of *A. niger* in 30 kg wheat bran substrate using the pilot-scale packed-bed bioreactor (50 cm height). Casciatori and Frassetto³³ indicated that increasing aeration rate helped reduce the metabolic heat during the SSF, but the decreasing moisture content of the bed became a problem. The temperature at the center of the CB was higher than the lower and upper positions throughout the SSF. To decrease the bed temperature, the bed of the CB was divided into two and three beds as in the 2LB and 3LB. The temperatures in both layers of the 2LB (2LB-Bottom and 2LB-Top) slightly decreased from that in the CB. The highest temperature in the 2LB-Bottom and 2LB-Top were

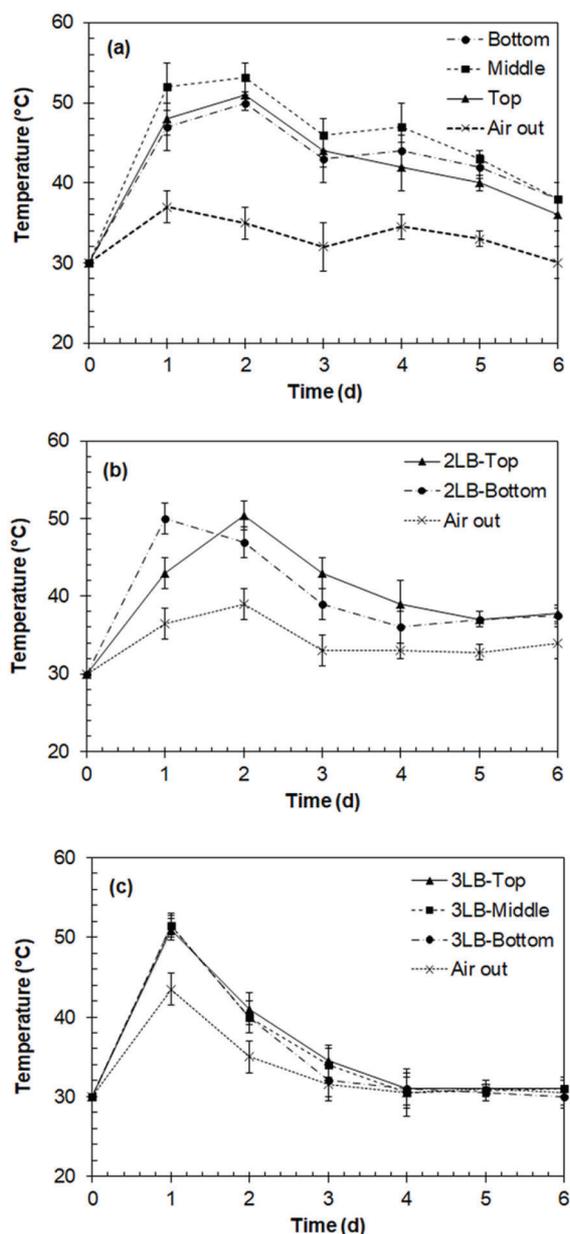


Fig. 2 – Temperatures of beds and outlet air during SSF of *A. niger* (a) Control packed-bed bioreactor, CB; (b) Two-layered packed-bed bioreactor, 2LB; (c) Three-layered packed-bed bioreactor, 3LB

48.7 ± 0.8 °C at day 1, and 50.4 ± 1.9 °C at day 2, and reduced to 37.5 ± 0.8 °C and 37.8 ± 1.1 °C at day 6, respectively. For the 3LB, it was found that all temperatures of the three beds were quite similar. The temperature of each bed in the 3LB measured at day 1 changed from 30.0 ± 0.5 °C ($t = 0$ d) to 51.5 ± 1.5 , 51.5 ± 1.2 and 51.3 ± 1.0 °C for the 3LB-Bottom, -Middle and -Top, respectively. From all studied packed-bed bioreactors, the maximum temperatures were around 50 °C at day 1, except that of the 2LB-Top, which increased up to day 2. The greatest reduction rate of temperature (highest slope of decreasing temperature in Fig. 2(c)) was detected in the 3LB. It was found that more heat in

the 3LB was removed by flowing air, since the highest temperature of its outlet air (43.5 ± 2.1 °C) was higher than that of the CB (37.0 ± 1.5 °C) and the 2LB (39.1 ± 1.9 °C at day 2). This indicated that dividing the packed bed had increased heat removal in the packed-bed bioreactor due to a reduction in bed compaction.

Growth of *A. niger*

The fungal growth of *A. niger*, shown in Fig. 3, is reported as glucosamine content in fermented substrate. The highest glucosamine content of the sample from the culture in the CB was found on day 2 as 76.9 ± 11.5 mg g⁻¹_{DS}, which means its specific growth rate in 0–1 d was 7.7 d⁻¹. The growth phase changed to stationary after day 2. For SSF in the 2LB, the fungal growth separately measured in each layer showed that the maximum glucosamine contents in the fermented samples were detected on day 2 as 73.9 ± 8.6 and 84.2 ± 5.9 mg g⁻¹_{DS}; however, their growth phases from 0–1 d showed the specific growth rates as 7.1 and 8.3 d⁻¹ for the 2LB-Bottom and -Top, respectively. The characteristics of growth curves obtained from the lower (2LB-Bottom) and upper (2LB-Top) beds of the two-layered bioreactor were similar to that of the CB. The highest glucosamine content in the fermented samples from the control was not significantly different from both layers of the 2LB. The different fungal growth in different layers of the bioreactor was clearly present in the 3LB, as shown in Fig. 3(c). The maximum glucosamine contents in the cultures of each bed of the 3LB were 33.3 ± 3.7 mg g⁻¹_{DS} at day 1, 62.4 ± 6.8 mg g⁻¹_{DS} at day 2 and 67.4 ± 5.4 mg g⁻¹_{DS} at day 1 for the 3LB-Bottom, 3LB-Middle and 3LB-Top, respectively. The highest specific growth rate found in the 3LB was 8.8 d⁻¹ analyzed from the 3LB-Top. Even if the maximum glucosamine contents in the fermented samples from the 3LB-Bottom and -Middle were different, their specific growth rates were similar as 8.1 d⁻¹.

Obviously, a higher amount of aerial mycelium, observed by white color of *A. niger* mycelium, was found in the higher bed position of bioreactor including the top of the CB, the 2LB-Top and the 3LB-Top. Therefore, the large amount of glucosamine in the samples cultured on the higher position of the bioreactor might be due to the growth of aerial mycelium of *A. niger*. As a role of aerial mycelium related to oxygen supply^{34–36}, it could be implied that low partial pressure of oxygen occurring at the top of bioreactor enhanced aerial mycelium generation in order to increase oxygen supply.

The results of the bed temperatures in Fig. 2 show the direct correlation between bed temperature and fungal growth. When the temperature reached around 50 °C, the growth stopped and then

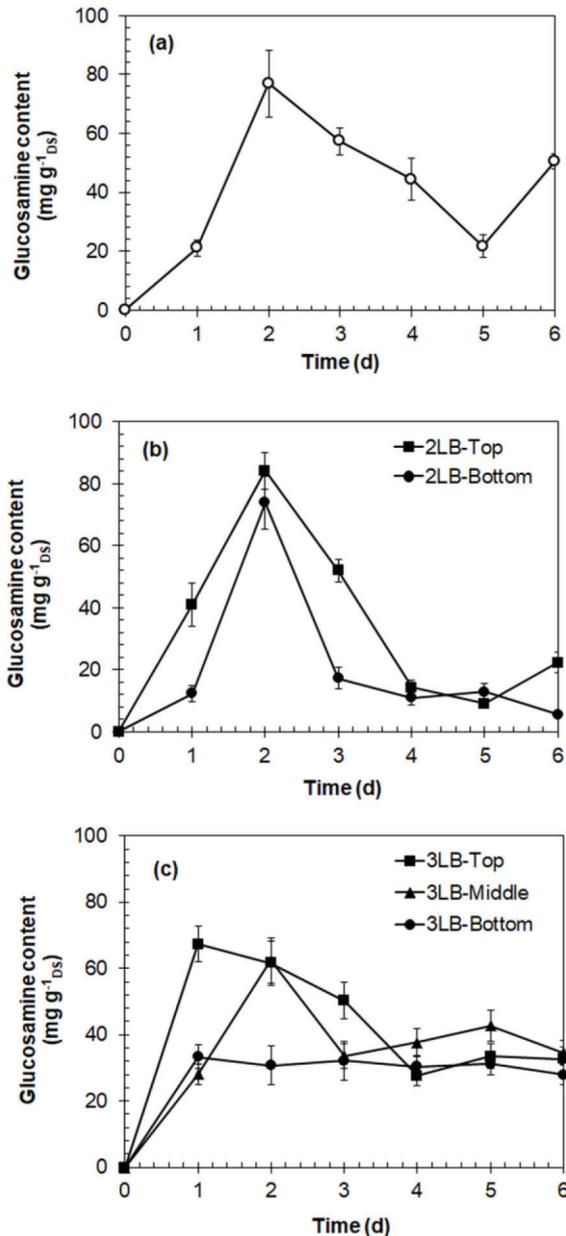


Fig. 3 – Glucosamine contents in fermented substrate. The glucosamine contents represent the fungal growth (a) Control packed-bed bioreactor, CB; (b) Two-layered packed-bed bioreactor, 2LB; (c) Three-layered packed-bed bioreactor, 3LB.

decreased (death phase), because the fungal hyphal branching decreased at the temperature higher than 45 °C³⁷. After the growth had stopped, the temperature of all beds decreased due to a decrease in heat generation from fungal respiration³⁸. The decrease in glucosamine content indicated self-degradation of fungi, where the fungal cell was destroyed by many fungal self-produced enzymes, including cellulase³⁹. The glucosamine obtained from a degradation of chitin was used as carbon and nitrogen sources for the next generation of *A. niger*. Moreover, the new spores of *A. niger* were detected in

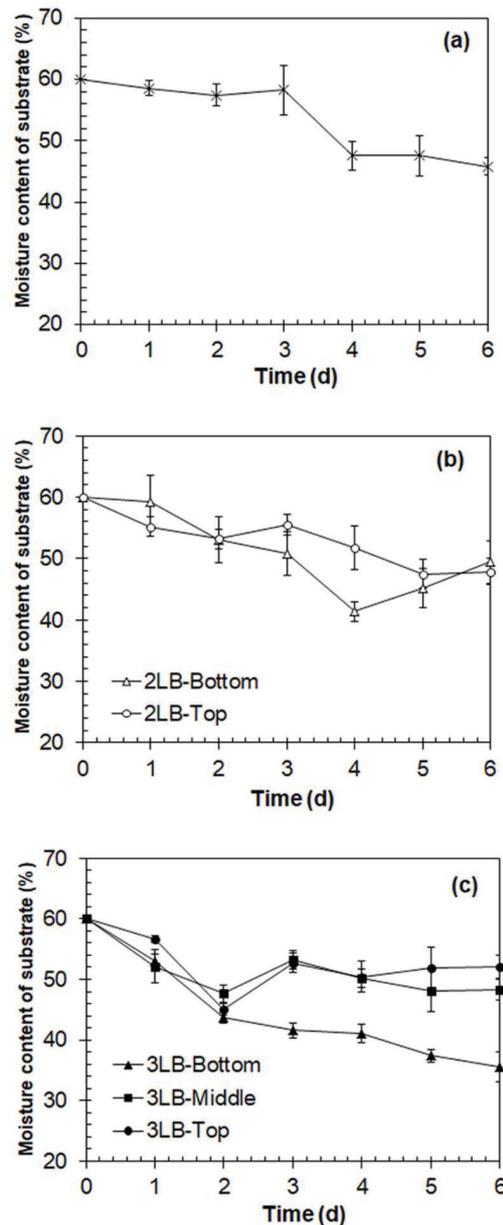


Fig. 4 – Moisture contents of fermented substrate (a) Control packed-bed bioreactor, CB; (b) Two-layered packed-bed bioreactor, 2LB; (c) Three-layered packed-bed bioreactor, 3LB

the substrate on days 4–5 causing increased content of glucosamine on day 6.

In addition, the fungal growth without the degradation of glucosamine was only found in the SSF at the 3LB-Bottom due to low moisture content of substrate, which was unsuitable for its growth and metabolisms. Selvakumar *et al.*⁴⁰ also reported that the optimum initial moisture content for the growth of *A. niger* on tea waste was about 60 %, and found that the growth decreased when initial moisture content was lower than 45 %. Moreover, a reduction in moisture content of the substrate affected the

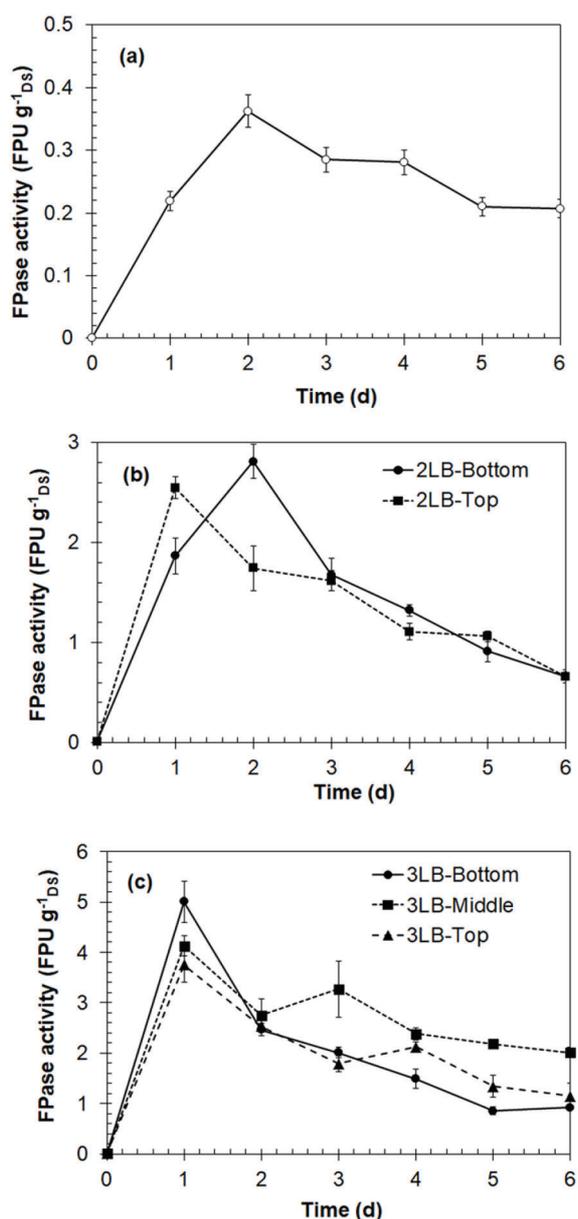


Fig. 5 – Cellulase production of *A. niger* in the SSF (a) Control packed-bed bioreactor, CB; (b) Two-layered packed-bed bioreactor, 2LB; (c) Three-layered packed-bed bioreactor, 3LB

oxygen and nutrient transfers during the SSF⁴¹. As shown in Fig. 4, the lower moisture content of substrate was found at the layer near the bottom because the aeration was forced at the bottom of bioreactor leading to moisture transfer from the bottom bed to the top bed of bioreactor.

Cellulase (FPase) activity

The cellulase production of *A. niger* shown in Fig. 5 suggests that the maximum FPase activity obtained from the CB was only 0.36 ± 0.03 FPU g⁻¹_{DS} at day 2, while that in the 2LB-Bottom and -Top were higher, i.e., 2.81 ± 0.22 and 2.55 ± 0.18 FPU g⁻¹_{DS}, respectively. Even if the FPase activity mea-

sured from the bed of the 2LB-Bottom was significantly higher than that of the 2LB-Top, the time to reach the maximum value was delayed 1 d. According to the temperature results presented in Fig. 2, the FPase activity corresponded to the change in bed temperature. When the bed temperature increased to 50 °C, which was the maximum value, the FPase activity in the fermented samples decreased. This phenomenon also occurred in the 3LB. The highest FPase activity in the 3LB was detected in the 3LB-Bottom as 5.01 ± 0.41 FPU g⁻¹_{DS} followed by that in the 3LB-Middle and the 3LB-Top as 4.13 ± 0.21 and 3.75 ± 0.34 FPU g⁻¹_{DS}, respectively. The results obtained from the 3LB clearly showed that the highest cellulase production was obtained in the bottom layer, higher than that in the middle and top layers. Since the aeration was forced from the bottom of bioreactor, the air in the bottom zone contained a higher oxygen concentration, which reduced along the height of bioreactor due to fungal respiration utilizing oxygen. This showed the importance of oxygen for cellulase production. Even if the highest biomass production was detected at the top of the bioreactor, the highest cellulase production was found at the bottom. This might be because the aerial mycelium did not support the cellulase production. These results are in agreement with the work of Rahardjo *et al.*³⁶, indicating that the secretion of the enzyme increased in the cultivation without aerial mycelium. Dividing substrate bed into many layers reduced the compactness of bed caused by the mass of upper bed, leading to an increase in the possibility of fungi contacting oxygen in forced air.

Conclusion

Copra waste from coconut milk processing was a good substrate for cellulase fermentation by *A. niger*. The most important factors were the controls of bed temperature, water migration, and oxygen concentration. Using a multi-layer packed-bed bioreactor helped increase the heat removal, and consequently, the cellulase production increased. The quality of air forced through the bed along the packed bed should be adjusted in order to obtain higher production yield of cellulase from all layers of a multi-packed-bed bioreactor. The thickness of the bed should be considered to relate with the temperature and moisture uniformity, as well as oxygen partial pressure.

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