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Short Communication

Bioethanol Levels of Dragon Fruit (*Hylocereus polyrhizus*) Peel with the Addition of Blend Crude Cellulase Enzyme from *Trichoderma reesei* and *Aspergillus niger*

Trianik Widyaningrum^{1,2*}, Masreza Parahadi³

- 1) Department of Biology Education, Faculty of Teacher Training and Education, Ahmad Dahlan University, Kampus 4, Jl. Prof. Dr. Soepomo, Janturan, Yogyakarta 55164, Indonesia.
- 2) Department of Biology, Faculty of Mathematics and Natural Sciences, Brawijaya University, Gedung Biologi, Jl. Veteran, Malang 65145, Indonesia.
- 3) Department of Postgraduate Program in Biotechnology, Postgraduate School, Universitas Gadjah Mada, Jl. Teknik Utara, Berek Yogyakarta 55281, Indonesia.

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ABSTRACT

The petroleum fuel crisis shows that Indonesia's fossil energy reserves are limited. It is necessary to develop an environmentally, friendly and sustainable alternative energy, one of which is bioethanol. This study aims to determine the bioethanol levels of dragon fruit (*Hylocereus polyrhizus*) peel with the treatment of cellulase enzymes from *Trichoderma reesei* and *Aspergillus niger*. This research was an experimental study that uses steps such as making dragon fruit peel substrate and filtrate, cellulose degradation with enzymes from *Trichoderma reesei* and *Aspergillus niger* and inoculating with yeast (*Saccharomyces cerevisiae*) with a fermentation time of 96 hours and then measured reducing sugar levels with the method of DNS, distillation and the measurement of bioethanol levels using alcohol meters. The results have shown that using enzymes from *Trichoderma reesei* and *Aspergillus niger* can increase the reduction of 49.68% sugar levels with the treatment of *T. reesei*: *A. niger* (3: 1) and produce the highest bioethanol level, which is 2.46% in the treatment of *T. reesei*: *A. niger* (2: 1).

Keywords: *Aspergillus niger*, dragon fruit peel, cellulase, *Trichoderma reesei*

The increasing demand for ethanol for various industrial purposes such as alternative sources of energy, industrial solvents, cleansing agents and preservatives has necessitated increased production of this alcohol (Ali et al., 2011). In the current time, the importance of alternative energy sources has become even more necessary not only due to the continuous depletion of limited fossil fuel stock but also for a safe and better environment. With the inevitable depletion of the world's energy supply, there has been an increasing worldwide interest in alternative sources of energy (Lynd et al., 2017; Chandel et al. 2007; Wyman, 1999; Herrera 2004; Herrera, 2006; Lin, 2006; Vertes & Inui, 2006; Schubert, 2006; Dien, 2003).

Currently, biomass-derived ethanol is produced at an industrial scale from sucrose and starch; however, this poses concerns about the potential competition with food and feed supplies (Hahn-Hägerdal et al, 2006; Field, Campbell, J. E., & Lobell, 2008). Hence, other alternatives such as the production on fallow fields of crops and grasses to produce biofuels have recently attracted attention. In particular, the lignocellulosic materials such as agricultural wastes are considered to be the main potential sources of biomass for second generation bioethanol production (Hu., et al., 2008; Hahn-Hägerdal et al., 2006; Sakai et al, 2007; Merino & Cherry 2007; Goh et al, 2010).

Bioethanol is one of the renewable alternative fuels that have the potential to be developed in Indonesia. Bioethanol is produced from biomass fermentation processes aided by microorganisms.

*Corresponding author

Tel.: +62 816682123

Email: trianikwidyaningrum@pbio.uad.ac.id

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The requirement to make bioethanol is a biological material that has sugar content (glucose, starch, and fibre) (Hambali et al, 2007) among other is dragon fruit peel. Dragon fruit including cactus or Cactaceae family, red dragon fruit peel contains sugar component around 8.4% and also other complex carbohydrates like cellulose around 68.3% (Jamilah et al., 2011).

Table 1. Sugar levels of Dragon fruit peel after treatment of *T.reesei* and *A.niger*

Cellulase ratio between <i>T. reesei</i> and <i>A. niger</i>	Sugar level (mg/mL)
Control	49.41 ± 3.94 ^a
1:0	94.31 ± 3.56 ^{de}
0:1	91.25 ± 2.29 ^d
1:1	73.60 ± 1.77 ^b
2:1	96.95 ± 4.98 ^{de}
1:2	84.44 ± 4.69 ^c
3:1	98.20 ± 2.83 ^e
1:3	76.66 ± 3.56 ^b

The same letter within each row do not differ significantly ($p > 0.05$) according to the Duncan test.

The cellulose degradation process can be done chemically or biologically using cellulolytic organisms originate from bacteria or fungi, degradation of cellulose into simpler sugars in the form of both cellobiose and glucose with the help of a catalyst. Hydrolysis can be carried out chemically (acid) or enzymatically, enzymatic hydrolysis using cellulase enzymes. Cellulase enzymes can be produced from cellulolytic microbes both mold and bacteria, while molds commonly used from *Trichoderma* and *Aspergillus*. *Trichoderma reesei* has been widely used for the production of commercial cellulase (Vandana and

Anahit, 2014). Cellulase is a multi-component enzyme comprising of endoglucanase, which attacks cellulose in the amorphous zone and releases oligomers such as cellobiohydrolase, that liberate cellobiose from reducing and non-reducing ends also β -glucosidase, which hydrolyze cellobiose to glucose and play a key role in avoiding cellobiose inhibition and thus enhancing the hydrolysis rates of cellulose into glucose (David, 2008; Mehdi et al., 2010; Sunkyu et al., 2010; Baljit, 2014; Veeresh and Wu, 2014). *Aspergillus niger* produces an enzyme that plays a role in accelerating the conversion of cellobiose to glucose, the enzyme β -glucosidase (Juhasz et al., 2003). The combination of cellulase enzymes from *T. reesai* and *A. niger* can be expected to increase the change of cellulose into glucose which is a material for making bioethanol.

This study aims to determine the levels of bioethanol dragon fruit peel (*Hylocereus polyrhizus*) with the treatment of cellulase enzymes from *Trichoderma reesei* and *Aspergillus niger*.

Dragon fruit peels were obtained from a local market in Indonesia. For analytical purposes, the Dragon fruit peels were made powder. *Trichoderma reesei* and *Aspergillus niger* culture were obtained from The Food & Nutrition Culture Collection (FNCC), Food and Nutrition Center Universitas Gadjah Mada. The strains were maintained on PDA and incubated at room temperature for seven days (Safaria et al., 2013).

A total of 5 g of Dragon fruit peel powder were put into Erlenmeyer 250 mL and added 25 mL nutrition solutions which every 1000 mL contains 1.0 g yeast extract, 1.5 g peptone, 1.4 g $(\text{NH}_4)_2\text{SO}_4$; 2.0 g KH_2PO_4 , 0.005 g $\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$, 5 mL solution CMC 1%. The flasks were sterilized for 15 minutes at 121°C. Two milliliters of spores (107-108 spores/mL) were inoculated and incubates at room

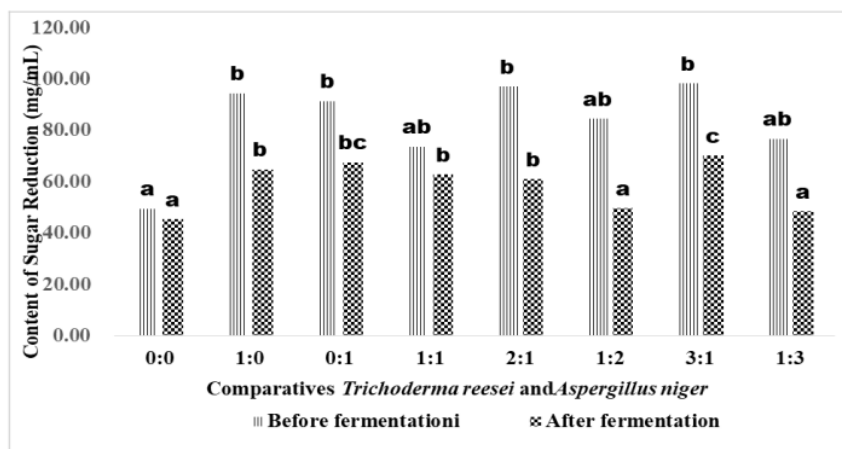


Figure 1. Reduced sugar levels before and after the treatment with *Saccharomyces cerevisiae*

temperature and static condition, for 6 days for *Trichoderma reesei* and 8 days for *Aspergillus niger* (Sri Winarsih et al., 2014).

Tween 80 0.1% solution was taken 100 mL and poured into the dragon fruit peel sample then stirred at 150 rpm for 120 minutes at room temperature. The solution was then centrifuged at 3000 rpm for 10 minutes. The supernatant obtained was used as a crude enzyme extract (Szendefy et al., 2006).

Dragon fruit peel porridge put into Erlenmeyer 100 mL then added the crude enzymes *T.reesei* and *A.niger* were added as much as 10% each according to the treatment with variations of 1: 0, 0: 1, 1: 1, 2: 1, 1: 2, 3: 1, 1: 3 using a measuring pipette aseptically and made three replications (21 Erlenmeyer), then stirred using a sterile glass stirrer. Erlenmeyer was covered with sterile cotton and coated with aluminum foil then incubated for 24 hours using an incubator at 37 °C. Hydrolysis results were measured for reducing sugar levels.

The culture of *Saccharomyces cerevisiae* JCM 3012 was obtained from The Food & Nutrition Culture Collection (FNCC), Food and Nutrition Centre, Universitas Gadjadara. The strain was maintained on YM to keep medium (yeast extract 3 g/L, malt extract 3 g/L, peptone 5 g/L, glucose 10 g/L) at 4 °C. To carry-out the tests *S. cerevisiae* was grown overnight at 30 °C on a rotary shaker (INNOVA 44, Incubator Shaker Series, New Brunswick Scientific) at 200 rpm, in tubes containing 20 ml YM medium.

Dragon fruit peel which produced on hydrolysis (21 Erlenmeyer treatment with crude enzyme mixture *A.niger* and *Trreesei*), each were added *Saccharomyces cerevisiae* at a dose of 10%, then each fermented for 96 hours for bioethanol production. The results of the fermentation

treatment were measured by reducing sugar levels with the DNS method to compare sugar levels before and after treatment with *S. cerevisiae* (Jackson and Jayanthi, 2014).

Table 1 shown that reducing sugar in the dragon fruit peel produced after treatment with *T. reesei* and *A.niger* were seen an increase because *T. reesei* and *A.niger* could produce cellulase enzymes, which can hydrolyze cellulose and hemicellulose into glucose. Enzymatic hydrolysis was regarded today as the most promising approach to liberating fermentable sugars in an energy-efficient way from the carbohydrates found in lignocellulosics in order to produce ethanol (Galbe and Zacchi, 2007).

Fig. 1 shown that reduced sugar decreased during the fermentation process with *S. cerevisiae*. The sugar content in the medium was continuously utilized by *S. cerevisiae* cells for the growth and formation of ethanol. The more reducing sugars used by *Saccharomyces cerevisiae* cells, the higher ethanol concentration produced and vice versa the less reducing sugars used, the lower the ethanol concentration. The increase in sugar concentration up to a certain level caused the fermentation rate to increase. However, the use of excessive sugar concentration will cause a steady fermentation rate, because the concentration of sugar used beyond the uptake capacity of the microbial cells. Generally, the maximum rate of ethanol production was achieved when using sugars at a concentration of 150 g/L. The initial sugar concentration also has been considered an important factor in ethanol production. (Zabed et al., 2016)

Fig 2. shown that the levels of bioethanol produced by fermentation are obtained by varying levels of bioethanol. The presence of fermented bioethanol was based on the opinions of Campbell, Reece, J.B., and Nitchel (2003), the results of *S.*

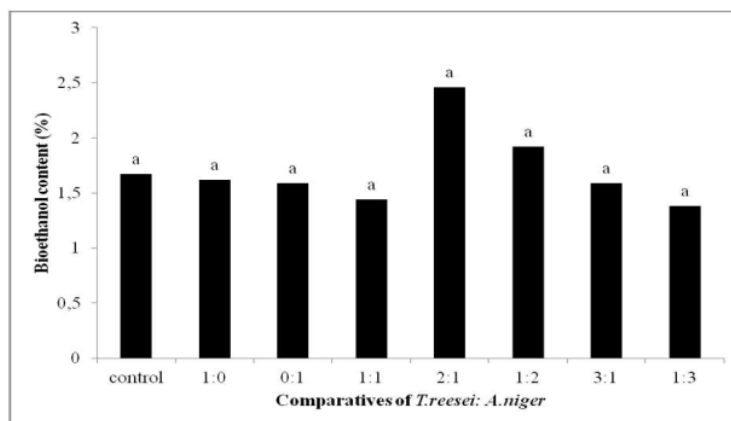


Figure. 2. Bioethanol levels of dragon fruit peel after fermentation using *Saccharomyces cerevisiae*. The DMRT test ($p > 0.05$) bioethanol levels were not differ significantly

cerevisiae metabolism in carbohydrate-based food sources such as sugar, starch, and cellulose are bioethanol. The presence of bioethanol indicates that the *S. cerevisiae* fermentation process is going well. According to the opinion (Wirahadikusumah, 2002) that the decomposition of carbohydrate or cellulose into pyruvate with the help of pyruvate decarboxylase enzyme which is reduced to bioethanol is through the event of glycolysis.

Enzymes from *Trichoderma reesei* and *Aspergillus niger* can increase reducing sugar levels 49.68 % in the treatment of *T. reesei*: *A. niger* (3: 1) and produce the highest bioethanol level, which is 2.46 % in the treatment of *T. reesei*: *A. niger* (2: 1)

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