

# Hyper-production of raw-starch-digesting enzyme by mutant fungal strain and optimisation of solid by-products

Sản xuất cao sản enzyme phân hủy tinh bột sống bởi chủng đột biến và môi trường tối ưu

Research article

Vu, Van Hanh<sup>1</sup>\*; Keun, Kim<sup>2</sup>

<sup>1</sup>Institute of Biotechnology, Vietnam Academy of Science and Technology, Hanoi 10600, Vietnam; <sup>2</sup>Department of Bioscience and Biotechnology, University of Suwon, Hwaseong 445-743, Korea

Selected fungal strain for production of raw-starch-digesting enzyme by solid state fermentation was improved by sequential exposures to  $\gamma$ -irradiation of Co<sup>60</sup>, ultraviolet and treatments with N-methyl-N'-nitrosoguanidine. Mutant *Aspergillus* sp. CXN2-3A was chosen and its production of raw-starch-digesting enzyme (RSDE) was improved 2 folds higher than that of wild type. Optimal condition for the production of the enzyme using wheat bran as the substrate was accomplished for the CXN2-3A. With the optimal fermentation condition and the solid medium supplemented with urea and NH<sub>4</sub>NO<sub>3</sub>, CoSO<sub>4</sub>, Tween 80, 1% glucose, CXN2-3A produced RSDE 19.23 folds higher than wild type cultured in pre-optimized condition and un-supplemented medium.

Chủng nấm chọn lọc sản xuất enzyme thủy phân tinh bột bằng cách lên men trạng thái rắn, chủng nấm được cải thiện bằng chiếu xạ tia cực tím, tia  $Co^{60}$  và các phương pháp xử lí với N-methyl-N'nitrosoguanidine. Mutant Aspergillus sp. CXN2-3A, đã được lựa chọn để sản xuất enzyme (RSDE) thủy phân tinh bột sống cải thiện cao hơn 2 lần so với chủng dại. Điều kiện tối ưu cho việc sản xuất các enzyme bằng cách sử dụng cám, lúa mì đã được thực hiện cho CXN2-3A. Với điều kiện lên men xốp tối ưu và bổ sung urê và NH<sub>4</sub>NO<sub>3</sub>, CoSO<sub>4</sub>, Tween 80, 1% glucose, CXN2-3A đã sản xuất RSDE cao gấp 19,23 lần so với kiểu dại ở cùng điều kiện.

**Keywords:** Aspergillus sp., N-methyl-N'-nitrosoguanidine, raw-starch-digesting enzyme, ultra violet,  $\gamma$ -irradiation of Co<sup>60</sup>

# 1. Introduction

Raw-starch-digesting enzyme (RSDE) composed of glucoamylase and other amylases is capable of hydrolyzing raw starch to glucose and other oligosaccharides at mild temperatures without requiring conventional two-step process of liquefaction and saccharification for ethanol fermentation (Kelly *et al.*, 1995; Yamamoto, 1994). The use of RSDE provides more advantages in conversion of biomass into sugars than single enzyme and acid hydrolysis since it gives higher quality products, energy efficiency, safer working environment and last longer equipment (Matsumoto *et al.*, 1982). RSDE was found in fungal strains such as *Aspergillus* sp., *A. usami, Rhizopus* sp., (Abe *et al.*, 1988; Abu *et al.*, 2005; Matsumoto et al., 1982). The improvement of microbial strains using different mutagenic agents for the over-production of industrial products has been considered in commercial fermentation process (Parekh *et al.*, 2000; Vu *et al.*, 2010). *Thermomyces lanuginosus* was treated with irradiation of ultraviolet (UV) and N-methyl-N'-nitrosoguanidine (NTG) to improve  $\alpha$ -amylase and glucoamylase (Rubinder *et al.*, 2002). *A. niger* was treated with  $\gamma$ -irradiation of Co<sup>60</sup> for improvement of glucoamylase production (Rajoka *et al.*, 2004). In this study, various mutagenic agents such as  $\gamma$ ray of Co<sup>60</sup>, UV, and NTG were combined to mutate fungal strains, then optimization of solid culture for hyperproduction of RSDE.

# 2. Materials and methods

#### 2.1 Strain improvement by mutations

Fungal strain C6.1 was found to be potent fungal strain for RSDE production and this strain was used for the strain improvement by mutation. The spores of C6.1 were harvested from 6 d-old spores grown on PDA plates and exposed to different doses (0.5 to 2.5 KGy with 0.5 KGy interval) of  $\gamma$ -ray. The survivors were grown by spreading 0.1 mL of treated spores on PDA plate containing 0.1% of raw starch-RBB and then incubated at 30°C for 4  $\div$  7 days. Based upon the clearing zones surrounding the colonies, y-ray mutant X2 was selected and was again treated with  $\gamma$ -ray of 2 KGy. According to Kuhad *et al.* (1994), the best  $\gamma$ -ray mutant X2 -2 was continuously exposed to UV irradiation at 30 W in a distance of 50 cm for  $5 \div 60$  min , where at 5 min interval the spores were taken out to test. The best mutant strain UV1 was sequentially treated with four cycles of NTG solution (100 µg/ ml in 0.2 M citrate buffer, pH 5) for 10 ÷ 60 min. After treatment with various mutagenic agents, many mutant strains were screened. Finally, one best mutant strain CXN2-3a was taken for further study.

#### 2.2 Determination of RSDE activity

The RSDE activity was determined as described in previous study (Vu *et al.*, 2010). One unit (U) of RSDE activity was defined as amount of enzyme in 1 mL, which liberates one  $\mu$ mol of glucose, per min, from raw rice starch (Miller, 1959).

#### 2.3 Preparation of seeding culture

In a 250 mL Erlenmeyer flask containing 10 g of rice bran powder and 90 mL distilled water, pH 3.5, was sterilized at 121°C for 20 min, and then cooled. To this, one plug (1 x 1 cm<sup>2</sup>) of 7 d-old conidia-mycelia were inoculated and incubated at 30°C in a shaker operated at 200 rpm for 1 day.

#### 2.4 Solid state of fermentation (SSF)

A 500 mL Erlenmeyer flask containing 20 g of rice bran with 33.3% (v/w) of moisture was plugged with cotton and sterilized for SSF.

# 2.5 Optimization of SSF

The SSF was carried out to study the effect of various parameters required for the optimum production of RSDE by selected mutant CXN2-3A. Substrate (wheat bran, rice husk, rice bran, saw dust), moisture (20, 30, 40, 50, 60, 70 and 80%, v/w), culture temperature (20, 25, 30, 35, 40, and  $45^{\circ C}$ ), initial pH of solid culture (3.0, 3.5, 4.0, 4.5, 5.0, 5.5, 6.0, 6.5, 7.0), culture time (2 ÷ 6 days), aeration area (15, 20, 25, 30, 35, 40, and 45 g of moistened wheat bran in a 500 mL Erlenmeyer flask), age of seeding culture (1 ÷ 4 days), inoculum size (10, 15, 20, 25, 30, 35, 40, 45, and 50%, v/w) were the parameters to be optimized.

The examined additives were carbon sources of glucose, maltose, rice starch, sucrose, and corn, each at 1%; nitrogen sources of urea, yeast extract, tryptone, tryptic soy, peptone, nutrient broth, skim milk, casamino acids, soytone, malt extract, NH<sub>4</sub>Cl, and NH<sub>4</sub>NO<sub>3</sub>, each at 1%; metal salts of CaCl<sub>2</sub>, MgCl<sub>2</sub>, MnCl<sub>2</sub>, CuSO<sub>4</sub>, CoSO<sub>4</sub>, FeSO<sub>4</sub>, ZnSO<sub>4</sub>, NaNO<sub>3</sub> and KCl, each at 2.5 mM; surfactant of Tween 20, Tween 80, Trixton-X100, each at 0.5% while SDS, and EDTA, each at 0.4 mM. SSF was carried out at 30°C for 3 days with wheat bran containing 33% of initial moisture, which was inoculated with 10% (v/w) of 1d-old seeding culture.

#### 2.6 Ethanol production by simultaneous saccharification and fermentation using RSDE and *Saccharomyces cerevisiae*

Uncooked rice winery waste rice-wine-cake slurry containing 25.73% (w/v) solid was used for simultaneous saccharification and fermentation (SSF). The SSF was conducted in a 100 liters fermentor containing 50 L slurry, pH 3.5, with crude enzyme of fungal strain of mutant CXN2-3a with 1.13 Unit of RSDE/ gds (gram dried substrate). To the slurry, yeast cells of *S. cerevisiae* KV25 was inoculated to make initial cell number of 2.5 x  $10^7/mL$ , incubated at 30°C for 5 days, and the ethanol formed was determined.

# 3. Results and discussion

#### 3.1 Screening and selection of a mutant strain

Selected fungal strain C6.1 was treated with two cycles of  $\gamma$ -irradiation. The best  $\gamma$ -ray-mutant strain X2-2 was then treated with UV irradiation. The best mutant UV strain was sequentially treated with four cycles of NTG. Finally, mutant strain CXN2-3A produced the highest activities of RSDE was selected. The mutant strain CXN2-3A exhibited 2.7 folds improved activity than that of wild type. The mutant CXN2-3A was identified as *Aspergillus* sp. with 99% of homology by sequencing the internal transcribed spacers.

Mutant strain CXN2-3A produced a high level of RSDE was obtained by mutagenic method and the strain was also stable for a long period to produce RSDE. In the past, the use of mutagenic agents such as  $\gamma$ -ray, UV, and NTG for hyper-production of industrial products has been reported but the repeated and sequential use of all of these mutagenic agents as shown in this study was not documented yet. A mutant Thermomyces lanuginosus obtained after treatment with three cycles of UV and NTG produced  $\alpha$ -amylase and glucoamylase 7- and 3-folds, respectively, higher than those of wild type (Rubinder et al., 2002). Mutant A. niger created by  $\gamma$ -irradiation produced glucoamylase 2.0- 2.5 folds higher than those of wild type (Rajoka et al., 2004). Vu et al. (2009) used repeated and sequential mutagenesis to mutate fungal strain, as a result, the mutant strain secreted more RSDE activity than that of wild type.

# **3.2 Optimal SSF-condition for the production of RSDE by mutant** *Aspergillus* sp. CXN2-3A

Substrate: Among several solid media, rice bran showed the highest production of RSDE (62.2 U/g) while ricehusk, and saw dust exhibited lower enzyme production. Of the various solid substrates used in this study, rice bran proved to be suitable for the colonization of *Aspergillus* sp. CXN2-3A as indication by the maximum visible growth on medium and the highest enzyme yield.

Initial moisture content: Moistened solid medium containing 50% moisture content yielded the highest level of RSDE (62.2 U/g), while those containing 20, 30, 40, 60, 70, and 80% produced lower activity. In SSF, moisture level plays an important role in biosynthesis and secretion of the enzymes. High moisture results in decreased substrate porosity as well as reduced oxygen penetration (Vu *et al.*, 2010) but low moisture leads to poor microbial growth and poor accessibility to nutrient (Pandey, 1992). *Aspergillus* sp. A3 at 80% of moisture using wheat bran (Ellaiah *et al.*, 2002)], *Rhizopus nigricans* PCSIR18 at 60 % of moisture (Mahmmod *et al.*, 1997).

The optimal temperature for the highest production of RSDE was at 30°C (Table 1). The optimal growth temperature varied with different fungi such as at 35°C for *A. niger* NRRL 3112 and *A. niger* NRRL 337 but at 30°C for *Aspergillus* sp. A3 (Ellaiah *et al.*, 2002). Incubation temperature at 30°C proved to be the best one for  $\alpha$ -amylase production by *A. oryzae* but higher temperature affected harmfully the enzyme production (Ramachandran *et al.*, 2004).

Among the physical parameters, the pH of the growth medium plays an important role by inducing morphological change in microorganism and in enzyme secretion. The high production of RSDE was observed at medium-pH of 4.5 (Table 1) and the production of RSDE was influenced by pH, which is very important in growth of microorganism and its metabolic activities. The metabolic activities of the microorganisms are very sensitive to changes in pH and also optimal pH varies with different microorganisms and enzymes. Thus, *A. niger* NCIM1245 produced maximum glucoamylase yields at pH 4.7 (Pandey & Radhakrishnan, 1993) but *Aspergillus* sp. A3 at pH 5 (Ellaiah *et al.*, 2002).

In the growth-related production of enzymes, after 3 days of incubation, the highest production (61.0 U/g) of RSDE was observed (Table 1). Incubation beyond 3 days was undesirable as this resulted in decreased enzyme production.

Aeration area (g solid medium / 250 mL Erlenmeyer flask): Amount of moistened rice bran to flask volume affects the porosity and aeration differently. It was proven that the high production of RSDE (65.0 U/g) was observed at (15 g/500 mL flask), therefore, this ratio was used for further study. Similar findings were reported by Bhatti (2007) and Ellaiah (2002), where ratio of substrate mass to flask volume of 1 : 50 was optimal for glucoamylase production by *Fusarium solani* and *Aspergillus* sp. A3.

Age of seeding culture: It was found that the seeding culture age of 2 days was suitable for inoculation and enzyme production but beyond 2 days was undesirable as these results in decreased enzyme yield. As a result, the highest production of RSDE (70.5 U/g) was observed (Table 1).

Inoculum size: The maximum production (78.5 U/g) was observed at inoculum size of 25% (v/w). This inoculum size was used for further study. Inoculum density is an important factor in an SSF process.

Effect of various medium-additives on the enzyme production: From the achieved data of enzyme production (Table 1), it was found that with the single addition of carbon source such as sucrose, maltose, rice starch and corn starch into the solid medium, there was a slight enhancement in the enzyme production of *Aspergillus* sp. CXN2-3A but with the addition of glucose to the medium, there was a significant enhancement of enzyme production (83.52 U/g). In other reports, glucoamylase production by fungi in wheat bran was enhanced by supplementation with sucrose (Anto *et al.*, 2006), fructose (Ellaiah *et al.*, 2002), starch (Bhatti *et al.*, 2007; Ramachandran *et al.*, 2004), but the improvement was not observed in maltose supplement (Ramachandran *et al.*, 2004).

It was found that the addition of urea or  $NH_4NO_3$  resulted in the highest RSDE production of 95.5 and 102.2 U/g, respectively, while other nitrogen sources showed a little enhancement or an inhibition of enzyme production (Table 1). The  $(NH_4)_2SO_4$  enhanced glucoamylase production by *A. awamori* (Anto *et al.*, 2006), yeast extract and peptone enhanced glucoamylase production (Bertolin *et al.*, 2003; Ramachandran *et al.*, 2004) urea, ammonium salts enhanced glucoamylase production (Bhatti *et al.*, 2007; Ellaiah *et al.*, 2002), (Ramachandran *et al.*, 2004).

The use of surfactants and fatty acids is well documented to increase the production of hydrolytic enzymes (Singh *et al.*, 1991). Surfactants have the potential to enhance microbial growth in SSF by promoting the penetration of water into the solid substrate matrix that leads to an increase in surface area (Asgher *et al.*, 2006). Herein, we found that Tween 80 enhanced the most highly the production of RSDE (85.7 U/g).

It is  $Cu^{2+}$  which is an essential metal ion to all organisms (Silva & Williams, 1993). An addition calcium ions onto medium enhanced  $\alpha$ -amylase activity (Nirmala & Muralikrishna, 2003), improved glucoamylase activity (Anto *et al.*, 2006). In the present study, it was shown that Co-SO<sub>4</sub> enhanced the most highly the RSDE production (88.52 U/g).

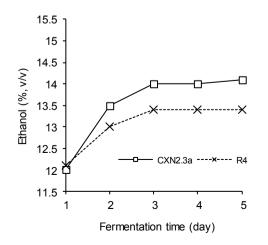
Production of RSDE at optimal condition with solid medium supplemented with additives: Optimal culture condition of solid rice bran for production of RSDE was accomplished, which were at 30°C, medium-pH of 4.5, moisture content of 50%, 2 d-old seeding-culture age, inoculum size of 25%, nitrogen additives of 1% urea and 1% NH<sub>4</sub>NO<sub>3</sub>, carbon additive of 1% glucose, metal additive of 2.5 mM CoSO<sub>4</sub>, surfactant additive of 0.05% Tween 80, substrate mass to Erlenmeyer flask volume of 3 : 100 (w/v). With this condition, the production of RSDE was 548.2 U/g. The results showed that the production of RSDE by mutant *Aspergillus* sp. CXN2-3A was improved 19.23 folds higher than that of wild type produced in pre-optimized condition and basal medium

without additives (Table 1). Also this 581.41 U/g of RSDE of *Aspergillus* sp. CXN2-3A was much higher than those of commercial enzyme preparation from *Rhizopus* sp. (102.3 U/g) and *A. usamii* (94.5 U/g).

Table 1. Effect of different factors on the RSDE production	Table 1.	. Effect of	different	factors on	the <b>RSDE</b>	production
---	----------	-------------	-----------	------------	-----------------	------------

$ \begin{array}{c c c c c c c c c c c c c c c c c c c $				<b>L</b>						
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	Moisture (%)	20	40	50	60	80				
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $		47.7	53.99	60.2	56.45	45.0				
pH of medium       3       4       5       6       7       8         U/g       36.3       60.4       58       53       50       47         Culture time (day)       2       3       4       5       6         U/g       20       61       57.83       52.5       43.41         Solid medium (g)/       15       20       360       40       50         Flask (250 mL)       U/g       65       62.2       45.5       38.2       37.2         Seeding age (day)       1       2       3       4	Temp. (°C)	20	25	30	35	40	45			
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	U/g	40	45	53.1	42	37.2	33.1			
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	pH of medium	3	4	5	6	7	8			
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Ū/g	36.3	60.4	58	53	50	47			
Solid medium (g)/ Flask (250 mL)15203604050Flask (250 mL)U/g6562.245.538.237.2Seeding age (day)1234U/g65.270.565.342Inoculumn size1020304050(%, v/w)U/g7573.278.568.362.5Carbon additiveNoneglucosecorn-su-malt-riceU/g7573.278.568.362.570.2U/g78.581.580.379.313.581.3Nitrogen additiveNoneNH4ClUreaMaltCaseinYeastTrypTrypticSoy- toneU/g78.274.595.578.974.478.578.576.580.5Nitrogen additiveNuneNH4NO3Pep-SkimNBViceViceU/g102.279.5776.597.597.550.570.2SurfactantNoneTweenTritonEDTASDS20.2U/g78.281.583.285.578.598.570.2Metal ions addi- U/gNoneMg2+Mn2+Cu2+Co2+Fe2+Zn2+K+Ca2+U/g78.274.583.285.290.356.6557.562.181.3ParameterWT inMT inMT inMT inMT inMT in </td <td>Culture time (day)</td> <td></td> <td></td> <td>4</td> <td></td> <td>6</td> <td></td> <td></td> <td></td> <td></td>	Culture time (day)			4		6				
Flask (250 mL)         U/g       65       62.2       45.5       38.2       37.2         Seeding age (day)       1       2       3       4       4         U/g       65.2       70.5       65.3       42       4         Inoculumn size       10       20       30       40       50         (%, v/w)       U/g       75       73.2       78.5       68.3       62.5         Carbon additive       None       glucose       corn- starch       su- crose       malt- ose       rice         U/g       78.5       81.5       80.3       79.3       13.5       81.3         Nitrogen additive       None       NH4Cl       Urea       Malt       Casein       Yeast       Tryp       Tryptic       Soy- extract         U/g       78.2       74.5       95.5       78.9       74.4       78.5       78.5       80.5         Nitrogen additive (continued)       NH4NO3       Pep-       Skim       NB       Sim       Soy- extract       Soy tone         U/g       102.2       79.5       776.5       97.5       Surfactant       None       Tween- -20       80       X100         U/g	U/g			57.83	52.5	43.41				
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Solid medium (g)/	15	20	360	40	50				
Seeding age (day)       1       2       3       4         U/g $65.2$ $70.5$ $65.3$ $42$ Inoculumn size       10       20       30       40       50         (%, v/w)       U/g $75$ $73.2$ $78.5$ $68.3$ $62.5$ Carbon additive       None       glucose       corn-       su-       malt-       rice         U/g $78.5$ $81.5$ $80.3$ $79.3$ $13.5$ $81.3$ Nitrogen additive       None       NH4Cl       Urea       Malt       Casein       Yeast       Tryp       Tryptic       Soy-         u/g $78.2$ $74.5$ $95.5$ $78.9$ $74.4$ $78.5$ $76.5$ $80.5$ Nitrogen additive       NH <sub>4</sub> NO <sub>3</sub> Pep-       Skim       NB           (continued)       tone       milk       U/g $102.2$ $79.5$ $776.5$ $97.5$ Suffactant       None       Tween-       Triton       EDTA       SDS $U/g$ $78.2$ $81.5$ $85.5$ $78.5$ $98.5$ $70.2$	Flask (250 mL)									
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $		65	62.2	45.5	38.2	37.2				
Inoculumn size         10         20         30         40         50           (%, v/w)         U/g         75         73.2         78.5         68.3         62.5           Carbon additive         None         glucose         corn-su-starch         malt-rice         rice           U/g         78.5         81.5         80.3         79.3         13.5         81.3           Nitrogen additive         None         NH4Cl         Urea         Malt         Casein         Yeast         Tryp         Tryptic         Soy-tone           U/g         78.2         74.5         95.5         78.9         74.4         78.5         78.5         80.5           Nitrogen additive         NH4NO3         Pep-         Skim         NB         vextract         extract         ton         soy         tone           (continued)         tone         milk         U/g         102.2         79.5         776.5         97.5         Surfactant         None         Tween         Triton         EDTA         SDS           U/g         78.2         81.5         85.5         78.5         98.5         70.2         Metal ions addi-         None         Mg2 <sup>2+</sup> Mu2 <sup>2+</sup> Co2 <sup>2+</sup>	Seeding age (day)	-		-	-					
	U/g				42					
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Inoculumn size	10	20	30	40	50				
Carbon additiveNoneglucosecorn- starchsu- crosemalt- osericeU/g78.581.580.379.313.581.3Nitrogen additiveNoneNH4ClUreaMaltCaseinYeastTrypTrypticSoy- extractU/g78.274.595.578.974.478.578.576.580.5Nitrogen additiveNH4NO3Pep-SkimNB(continued)tonemilkU/g102.279.5776.597.5SurfactantNoneTween-TritonEDTASDS-2080X100X100X100X100U/g78.281.585.578.598.570.2Metal ions addi-NoneMg <sup>2+</sup> Mn <sup>2+</sup> Cu <sup>2+</sup> Co <sup>2+</sup> Fe <sup>2+</sup> Zn2+K <sup>+</sup> Ca <sup>2+</sup> U/g78.274.583.285.290.356.6557.562.181.3ParameterWT inMT inMT inMT inbasMopt,50.550.5										
starch crose oseU/g78.581.580.379.313.581.3Nitrogen additiveNoneNH4ClUreaMaltCaseinYeastTrypTrypticSoy- toneU/g78.274.595.578.974.478.578.576.580.5Nitrogen additiveNH4NO3Pep-SkimNBKitrogen additiveNH4NO3Pep-SkimNB(continued)tonemilkU/g102.279.5776.597.5SUFACTARSDSU/g78.281.585.578.598.570.2Cu <sup>2+</sup> Ca <sup>2+</sup> Fe <sup>2+</sup> Zn2+K <sup>+</sup> Ca <sup>2+</sup> U/g78.274.583.285.290.356.6557.562.181.3ParameterWT in basMMT in basMMT in basMMT in opt,MTNone			73.2	78.5	68.3					
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Carbon additive	None	glucose	corn-	su-	malt-	rice			
Nitrogen additive         None         NH <sub>4</sub> Cl         Urea         Malt extract         Casein casein         Yeast yeast         Tryp ton         Tryptic soy         Soy- tone           U/g         78.2         74.5         95.5         78.9         74.4         78.5         78.5         76.5         80.5           Nitrogen additive (continued)         NH <sub>4</sub> NO <sub>3</sub> Pep-         Skim         NB         5         76.5         80.5           Surfactant         None         Tween         Tween-         Triton         EDTA         SDS           U/g         78.2         81.5         85.5         78.5         98.5         70.2           Metal ions addi- tives         None         Mg <sup>2+</sup> Mn <sup>2+</sup> Cu <sup>2+</sup> Co <sup>2+</sup> Fe <sup>2+</sup> Zn2+         K <sup>+</sup> Ca <sup>2+</sup> U/g         78.2         74.5         83.2         85.2         90.3         56.6         557.5         62.1         81.3           Parameter         WT in basM         MT in basM         MT in opt,         MT in         MT in										
extract       extract       ton       for       for       soy       tone         U/g       78.2       74.5       95.5       78.9       74.4       78.5       78.5       76.5       80.5         Nitrogen additive       NH4NO3       Pep-       Skim       NB       NB       continued       tone       milk       U/g       102.2       79.5       776.5       97.5       Surfactant       None       Tween       Tween-       Triton       EDTA       SDS         U/g       78.2       81.5       85.5       78.5       98.5       70.2       70.2         Metal ions addi-       None       Mg <sup>2+</sup> Mn <sup>2+</sup> Cu <sup>2+</sup> Co <sup>2+</sup> Fe <sup>2+</sup> Zn2+       K <sup>+</sup> Ca <sup>2+</sup> U/g       78.2       74.5       83.2       85.2       90.3       56.6       557.5       62.1       81.3         Parameter       WT in       MT in       MT in       MT in       MT in       basM       opt,	U/g					13.5				
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Nitrogen additive	None	NH <sub>4</sub> Cl	Urea	Malt	Casein	Yeast	Tryp	Tryptic	
Note of the system         NH4NO3         Pep- tone         Skim milk         NB           (continued)         102.2         79.5         776.5         97.5           Surfactant         None         Tween         Tween- result         Triton         EDTA         SDS           U/g         78.2         81.5         85.5         78.5         98.5         70.2           Metal ions addi- tives         None         Mg <sup>2+</sup> Mn <sup>2+</sup> Cu <sup>2+</sup> Co <sup>2+</sup> Fe <sup>2+</sup> Zn2+         K <sup>+</sup> Ca <sup>2+</sup> U/g         78.2         74.5         83.2         85.2         90.3         56.6         557.5         62.1         81.3           Parameter         WT in basM         MT in basM         MT in opt,         MT in         MT in										
$\begin{array}{c cccc} (continued) & tone & milk \\ U/g & 102.2 & 79.5 & 776.5 & 97.5 \\ \hline Surfactant & None & Tween & Triton & EDTA & SDS \\ & -20 & 80 & X100 \\ U/g & 78.2 & 81.5 & 85.5 & 78.5 & 98.5 & 70.2 \\ \hline Metal ions addi & None & Mg^{2+} & Mn^{2+} & Cu^{2+} & Co^{2+} & Fe^{2+} & Zn2+ & K^+ & Ca^{2+} \\ tives & & & & \\ U/g & 78.2 & 74.5 & 83.2 & 85.2 & 90.3 & 56.6 & 557.5 & 62.1 & 81.3 \\ \hline Parameter & WT in & MT in & MT in \\ & basM & basM & opt, \end{array}$	U/g					74.4	78.5	78.5	76.5	80.5
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$		NH <sub>4</sub> NO <sub>3</sub>	Pep-		NB					
Surfactant         None         Tween         Triton         EDTA         SDS $U/g$ 78.2         81.5         85.5         78.5         98.5         70.2           Metal ions addi- tives         None         Mg <sup>2+</sup> Mn <sup>2+</sup> Cu <sup>2+</sup> Co <sup>2+</sup> Fe <sup>2+</sup> Zn2+         K <sup>+</sup> Ca <sup>2+</sup> U/g         78.2         74.5         83.2         85.2         90.3         56.6         557.5         62.1         81.3           Parameter         WT in basM         MT in basM         MT in basM         MT in basM         MT in         MT in										
-20         80         X100           U/g         78.2         81.5         85.5         78.5         98.5         70.2           Metal ions addi- tives         None         Mg <sup>2+</sup> Mn <sup>2+</sup> Cu <sup>2+</sup> Co <sup>2+</sup> Fe <sup>2+</sup> Zn2+         K <sup>+</sup> Ca <sup>2+</sup> U/g         78.2         74.5         83.2         85.2         90.3         56.6         557.5         62.1         81.3           Parameter         WT in basM         MT in basM         MT in basM         MT in basM         MT in         MT in										
U/g         78.2         81.5         85.5         78.5         98.5         70.2           Metal ions addi- tives         None         Mg <sup>2+</sup> Mn <sup>2+</sup> Cu <sup>2+</sup> Co <sup>2+</sup> Fe <sup>2+</sup> Zn2+         K <sup>+</sup> Ca <sup>2+</sup> U/g         78.2         74.5         83.2         85.2         90.3         56.6         557.5         62.1         81.3           Parameter         WT in basM         MT in basM         MT in opt,         MT in         MT in	Surfactant	None				EDTA	SDS			
Metal ions addi- tives         None         Mg <sup>2+</sup> Mn <sup>2+</sup> Cu <sup>2+</sup> Co <sup>2+</sup> Fe <sup>2+</sup> Zn2+         K <sup>+</sup> Ca <sup>2+</sup> U/g         78.2         74.5         83.2         85.2         90.3         56.6         557.5         62.1         81.3           Parameter         WT in basM         MT in basM         MT in opt,         MT in         MT in	<b>T</b> T /	70.0				00 <b>5</b>	<b>7</b> 0 <b>0</b>			
tives         U/g         78.2         74.5         83.2         85.2         90.3         56.6         557.5         62.1         81.3           Parameter         WT in MT in basM basM opt,         MT in opt,         MT in basM opt,         MT in basM opt,         MT in basM opt,         MT in opt,									¥ 7 <sup>+</sup>	G 2+
U/g         78.2         74.5         83.2         85.2         90.3         56.6         557.5         62.1         81.3           Parameter         WT in basM         MT in basM         opt,         60.1         81.3		None	Mg	Mn	Cu	Co	Fe	Zn2+	K	Ca <sup>2</sup>
Parameter     WT in     MT in       basM     basM     opt,		70.0	<b>- - -</b>	02.2	05.0	00.2			(0.1	01.2
basM basM opt,	•				85.2	90.3	56.6	557.5	62.1	81.3
	Parameter	WT in	MT in	MT in						
U/g 28.5 58.7 548.2										
	U/g	28.5	58.7	548.2						

WT in basM, wild type of strain produced RSDE in basal medium; MT in basM, Mutant strain produced RSDE in basal medium; MT in opt, Mutant strain produced RSDE in optimized medium.



# Figure 1. SSF of RWC slurry treated with crude enzyme of fungal trains and *S. cerevisiae* KV-25

SSF of RWC in a 100 L fermentor using crude RSDE of mutant fungal strains of CXN2-3a, together with *S. cerevisiae* KV-25. *R4* (commercial enzyme) used as control

Ethanol production from raw starch using crude RSDE of mutant CXN2-3A by SSF: The ethanol production from winery waste rice-wine-cake (RWC) slurry by SSF was conducted in a 100 liters fermentor at 30°C. RSDE from *Aspergillus* sp CXN2-3A was excellent to treat raw-starch with high efficiency of ethanol conversion. The SSF using the enzyme of *Aspergillus* sp. CXN2-3A produced 15.82% (v/v) ethanol from 25.73% of solid, after 96 h (Fig. 1). Also the ethanol fermentation efficiency using

the enzyme of *Aspergillus* sp. CXN2-3A was  $92.7 \div 94.0\%$ , which is very high efficiency of ethanol production. The SSF process was carried out without cooking and economically more attractive.

### 4. Conclusion

It can be concluded that fungal strain CXN2-3A mutant is a potential microorganism for production of RSDE. The RSDE produced by the mutant strain can be applied successful ethanol production from slurry containing a high level of raw starch by SSF without application of liquefying enzyme and cooking. On the other hand, the results provided valuable information of using novel efficient mutation method to obtain mutation that produced a high level of RSDE.

# 5. References

- Abe, J.I., Bergmann, F.W., Obata, K., Hizukuri, S. 1988. Production of the raw starch digesting amylase of *Aspergillus* sp. K-27. Appl Microbiol Biotechnol 27: 447-450.
- [2] Abu, E.A., Ado, S.A., & James, D.B. 2005. Raw starch degrading amylase production by mixed culture of *Aspergillus niger* and *S. cerevisiae* grown on sorghum pomace. Afr. J. Bioethanol 4: 785-790.
- [3] Anto, H., Trivedi, U.B., Patel, K.C. 2006. Glucoamylase production by solid state fermentation using rice flask manufacturing waste products as substrate. Biores Technol 97: 1161-1166.
- [4] Asgher, M., Asad, M.J., & Legge, R.L. 2006. Enhanced lignin peroxides synthesis by *Phanerichaete chrysosporium* in solid state bioprocessing of a lignocellosis substrate. World J Microbiol. Biotechnol 22: 449-453.
- [5] Bertolin, T.E., Schmidell, W., Maiorano A.E., Casara, J., and Costa, J.A.V. 2003. Influence of carbon, nitrogen and phosphorous source on glucoamylase production by Aspergillus niger in solid state fermentation. Z. Naturforsch. 58c: 708-712.
- [6] Bhatti, H.N., Mohammad, H.R., Rakhshanda, N., Muhammad, A., Raheela, P., Abdul, J. 2007. Optimization of media for enhanced glucoamyalase production in solid state fermentation by *Fusarium solani*. Food technol Biotechnol 45: 51-56.
- [7] Ellaiah, P., Adinarayana, K., Bhavani, Y., Padmaja, P., Srinivasulu, B. 2002. Optimization of process parameters for glucoamylase production under solid state fermentation by a newly isolated *Aspergillus* species. Process Biochem 38: 615-620.
- [8] Kelly, C.T., McTigue, M.A., Doyle, E.M., Fogarty, W.M. (1995). The raw starch degrading alkaline amylase of *Bacillus* sp. IMD 370. J Ind Microbiol 15: 446-448.
- [9] Kuhad, R.C., Kumar, M., Singh, A. 1994. A hypercellulolytic mutant of *Fusarium oxysporum*. Lett Appl Microbiol 19: 397-400.

- [10] Mahmmod, A., Airengzeb, M., Baig, R., Ahmad, M.A. 1997. Production of amyloglucosidase by *Aspergillus niger* under different cultivation regimes Pak J Biochem Mol Biol 30: 49-54.
- [11] Matsumoto, N., Fukushi, O., Miyanaga, M., Kakihara, K., Nakajima, E., Yoshizumi, H. 1982. Industrialization of non-cooking system for alcoholic fermentation from grains. Agric Biol Chem 18: 1549-1558.
- [12] Miller, G.L. 1959. Use of dinitrosalicylic acid reagent for determination of reducing sugar. Anal Chem 31: 426-428.
- [13] Nirmala, M., Muralikrishna, G. 2003. Three amylases from malted finger millet (Ragi, Eleusine coracana, Indaf-15)-purification and partial characterization. Phytochemist 62: 21-30.
- [14] Pandey, A. 1992. Production of starch saccharifying enzyme (glucoamylase) in solid cultures. Starch 44: 75-77.
- [15] Pandey, A., Radhakrishnan, S. 1993. The production of glucoamylase by *Aspergillus niger* NCIM 1245. Process Biochem 38: 305-309.
- [16] Parekh, S., Vinci, V.A., Strobel, R.J. 2000. Improvement of microbial strains and fermentation processes. Appl Microbiol Biotechnol 54: 287-301.
- [17] Rajoka, M.I., Yasmin, A., Latif, F. 2004. Kinetics of enhanced ethanol productivity using raw starch hydrolyzing glucoamylase from *Aspergillus niger* mutant produced in solid state fermentation. Lett Appl Microbiol 39: 13-18.
- [18] Ramachandran, S., Patel, A.K., Nampoothiri, K.M., Francis, F., Nagy, V., Szakacs, G. 2004. Coconut oil cake -a potential raw material for the production of α-amylase. Biores Technol 93: 169-174
- [19] Rubinder, K., Chadha, B.S., Singh, N., Saini, H.S., Singh, S. 2002. Amylase hyperproduction by deregulated mutants of the thermophilic fungus *Thermomyces lanuginosus*. J Ind Microbiol Biotechnol 29: 70-74.
- [20] Silva, J., Williams, R. (Eds.). 1993. The biological chemistry of the elements. New York, USA: Clarendon Press.
- [21] Singh, A., Abidi, A.B., Darmwal, N.S., Agrawal, A.K. 1991. Influence of nutritional factors of cellulase production from natural lignocellulosic residues by *Aspergillus niger*. Agric Biol Res 7: 19-27.
- [22] Vu, V.H., Pham, T.A., Kim, K. 2010. Improvement of a fungal strain by repeated and sequential mutagenesis and optimization of solid-state fermentation for the hyper-production of raw-starch-digesting enzyme. J Microbiol Biotechnol 20(4): 718-726.
- [23] Yamamoto, S. 1994. Raw starch-digesting enzyme (maltooligosaccharide producing type) of *Zoogeloea ramigera*. J Appl Glycoscience 41: 283-289.