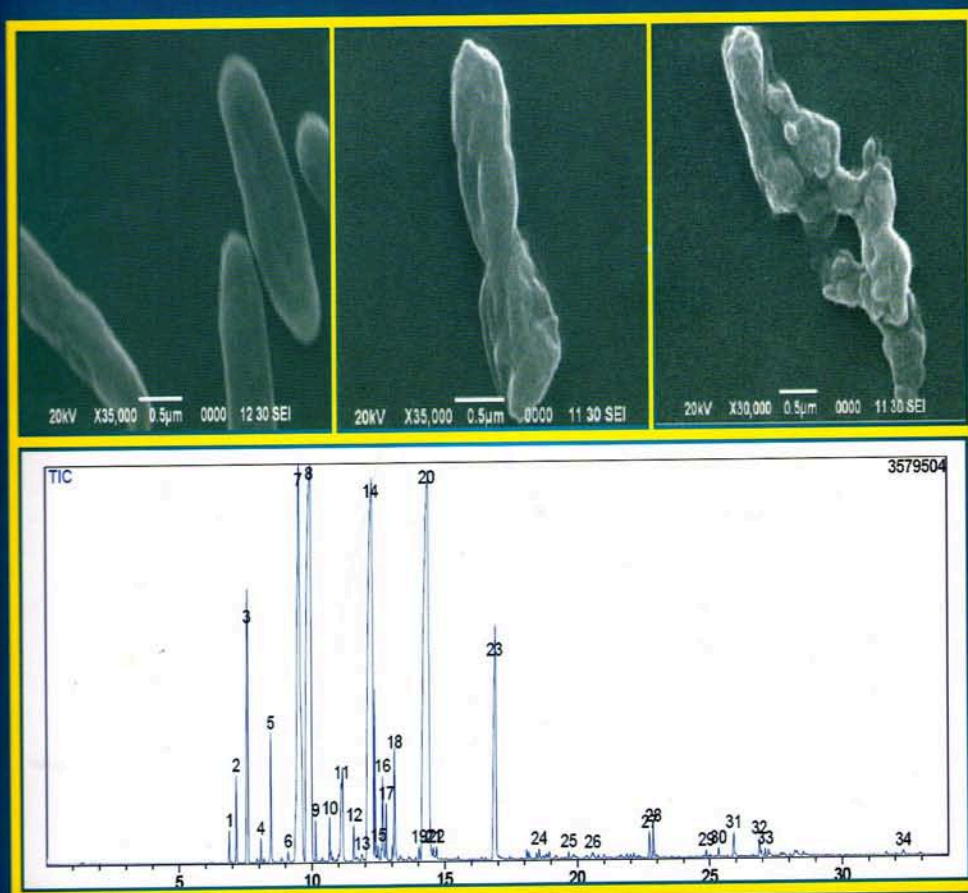


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## DAFTAR ISI

### MAKALAH HASIL RISET (ORIGINAL PAPERS)

#### **KERUSAKAN DINDING SEL *Escherichia coli* K1.1 OLEH MINYAK ATSIRI TEMU KUNCI (*Kaempferia pandurata*)**

[Cell Wall Disruption of *Escherichia coli* K1.1 by Temu Kunci (*Kaempferia pandurata*) Essential Oil]

Miksusanti, Betty Sri Laksmi Jennie, Bambang Ponco dan Gatot Trimulyadi.....1

#### **KERAGAMAN AKTINOMISETES KEPULAUAN WAIGEO, KABUPATEN RAJA AMPAT, PAPUA DAN POTENSINYA SEBAGAI PENDEGRADASI SELULOSA DAN PELARUT FOSFAT**

[Actinomycetes Diversity in Waigeo Island, Raja Ampat Regency, Papua and Their Potentials as Cellulose Degradation and Phosphate Solubilization]

ArifNurkanto.....9

#### **POTENSI IKAN MUJAIR (*Sarotherodon mossambica*) SEBAGAI BIOAKUMULATOR PENCEMARAN PESTISIDA PADA LINGKUNGAN PERTANIAN**

[The Potential of Mujair Fish (*Sarotherodon mossambica*) as Bioaccumulator of Pesticides Contamination in Agricultural Land]

Yulvian Sani dan Indraningsih.....19

#### **PEMBUATAN STARTER UNTUK EKSTRAKSI MINYAK KELAPA MURNI MENGGUNAKAN MIKROBA AMILOLITIK**

[Preparation of Starter for Extracting Virgin Coconut Oil by Using Amylolytic Microbes]

ElidarNaiola.....31

#### **RETRANSFORMATION AND EXPRESSION OF RECOMBINANT VIRAL PROTEIN OF JEMBRANA SU AND Tat (JSU AND JTat) IN pGEX SYSTEM**

[Retransformasi dan Ekspresi Protein Virus Rekombinan JSU dan JTat Penyakit Jembrana dalam Sistem pGex]

Endang T Margawati, Andi Utama and Indriawati.....39

#### **POPULASI POHON JENIS DIPTEROCARPACEAE DI TIGA TIPE HUTAN PAMAH KALIMANTAN**

[Tree Population of Dipterocarpaceae Species in Three Vegetation Types of Lowland Forests Kalimantan]

Herwint Simbolon.....45

#### **DAUR PATOLOGIS TEGAKAN HUTAN TANAMAN *Acacia mangium* Willd.**

[Pathological Rotation of *Acacia mangium* Willd. Forest Stand]

Simon Taka Nuhamara, Soetrisno Hadi, Endang Suhendang, Maggy T Suhartono, Wasrin Syafii dan Achmad.....59

#### **KEANEKARAGAMAN FLORA CAGAR ALAM NUSA BARONG, JEMBER - JAWA TIMUR**

[Floral Diversity of Nusa Barong Nature Reserve, Jember - East Java]

Tukirin Partomihardjo dan Ismail.....67

#### **KARAKTERISASI 17 FAMILI IKAN NILA (*Oreochromis niloticus*) GENERASI KE TIGA (G-3) BERDASARKAN METODE TRUSS MORFOMETRIKS**

[Characterization of 17 Families of Nile tilapia (*Oreochromis niloticus*) Third Generation (G-3) Based on Truss Morphometrics]

Nuryadi, Otong Zenal Arifin, Rudhy Gustiano dan Mulyasari.....81



<b>INDUKSI KALUS DAN REGENERASI TUNAS PULAI PANDAK (<i>Rauwolfia serpentina</i> L.)</b> <b>[Callus Induction and Shoot Regeneration of Pulai pandak (<i>Rauwolfia serpentina</i> L.)]</b> <i>Rossa Yunita dan Endang Gati Lestari</i> .....	<b>91</b>
<b>POTENSI ANTIBAKTERIA EKSTRAK DAN FRAKSI LIBO (<i>Piper mlnlatum</i> BL.)</b> <b>[Antibacterial Potential of Extract and Fraction of Libo (<i>Piper mlnlatum</i> BL.)]</b> <i>Sumarnie H Priyono</i> .....	<b>99</b>
<b>TOLERANSI SENGON BUTO (<i>Enterolobium cyclocarpum</i> Griseb) YANG DITANAM</b> <b>PADA MEDIA LIMBAH TAILING TERCEMAR SIANIDA DENGAN PERLAKUAN PUPUK</b> <b>[Tolerance of Sengon buto (<i>Enterolobium cyclocarpum</i> Griseb) Grown on Cyanide</b> <b>Contaminated Tailing Media with Fertilizer Application]</b> <i>Fauzia Syarif</i> .....	<b>105</b>
 <b><u>KOMUNIKASI PENDEK</u></b>	
<b>MENGESTIMASI NILAI KERUSAKAN TUMBUHAN INANG AKIBAT PEMARASITAN</b> <b>BENALU</b> <b>[Estimating the Destruction of Host Plant caused by Mistletoe Parasitizing]</b> <i>Sunaryo</i> .....	<b>111</b>

# RETRANSFORMATION AND EXPRESSION OF RECOMBINANT VIRAL PROTEIN OF JEMBRANA SU AND Tat (JSU AND JTat) IN pGEX SYSTEM [Retransformasi dan Ekspresi Protein Virus Rekombinan JSU dan JTat Penyakit Jembrana dalam Sistem pGex]

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## ABSTRAK

Genom virus penyakit Jembrana setidaknya memiliki 3 gen besar yang menyandi protein dan beberapa di antaranya diperlukan untuk replikasi virus. Protein JSU dan JTat diduga dapat menginduksi kekebalan yang protektif pada sapi Bali terhadap penyakit Jembrana sehingga keduanya sangat berpotensi untuk dipakai sebagai vaksin rekombinan. Penelitian ini dirancang untuk meretransformasi protein rekombinan JSU dan JTat ke dalam *Escherichia coli* menggunakan sistem pGEX. Konstruksi JSU dan JTat dalam pGEX dikoleksi plasmidnya dengan metode miniprep dan kemudian diretransformasikan ke dalam *E. coli* strain BL21 dan DH5a. JSU dan JTat hasil retransformasi diekspresikan pada medium LB untuk skala produksi kecil dengan sistem pGEX. Hasil penelitian ini menunjukkan bahwa kedua JSU dan JTat hasil retransformasi ke dalam *E. coli* strain BL21 terlihat tumbuh lebih baik pada medium LB jika dibandingkan retransformasi ke dalam *E. coli* strain DH5a. Hasil retransformasi JSU dan JTat dikarakterisasi dan diidentifikasi dengan *Western blotting* dan tampak menunjukkan ukuran protein yang benar, yaitu protein rekombinan JSU berukuran 60kDa dan JTat berukuran 36,7kDa. Protein rekombinan JSU muncul dengan pita tunggal dan lebih jelas jika dibandingkan dengan protein JTat. Konsentrasi protein JSU sedikit lebih rendah (1,883 mg ml<sup>-1</sup>) jika dibandingkan dengan JTat (1,981mg ml<sup>-1</sup>). Penelitian ini menunjukkan bahwa JSU pGEX masih tersimpan dan diekspresikan dengan baik, sementara JTat mungkin perlu dilakukan perakitan ulang untuk memantapkan ekspresinya.

**Kata kunci:** Jembrana, sapi Bali, protein rekombinan, JSU, JTat.

## ABSTRACT

The genome of Jembrana disease virus (JDV) has at least three major genes encoding for the viral proteins and some of them use it for its replication. Recombinant proteins of JSU and JTat are expected to induce a protective immune response against the virus and are, therefore potential for the development of a protective recombinant vaccine for Jembrana disease. This research was designed to retransform both JSU and JTat into *Escherichia coli* and expressing the proteins using pGEX system. Constructs of pGEX bearing JSU and JTat genes were collected by a miniprep method and then retransformed either into *E. coli* of BL21 or DH5a strains. The retransformed JSU and JTat were expressed in a small scale production of LB medium using pGEX system. The results showed that both retransformed JSU and JTat into *E. coli* strain BL21 appeared to be better growing on LB medium as compared to that of transformed constructs into strain DH5a. In western blotting assay, both retransformed JSU and JTat produced the right size of 60kDa for JSU recombinant protein and 36.7kDa for JTat recombinant protein. Recombinant protein JSU showed a sharper and a single band as compared to that JTat protein. Protein concentration of JSU was slightly lower (1.883mg ml<sup>-1</sup>) as compared to that JTat (1.981mg ml<sup>-1</sup>). This result suggests that JSU construct in pGEX is well-expressed in *E. coli* whereas JTat might need to be reconstructed to enhance its expression.

**Keywords:** Jembrana disease, Bali cattle, recombinant proteins, JSU, JTat.

## INTRODUCTION

Jembrana disease is an infectious, severe and acute viral disease in Bali cattle (Kertayadnya *et al.*, 1993). It was recognized initially in Jembrana districts in Bali, and was therefore termed as Jembrana disease virus (JDV). In Bali cattle, the JDV caused about 20% case fatality rate. The JDV was then identified as lentivirus of Retroviridae family (Chadwick *et al.*, 1995; Wilcox *et al.*, 1995). The prevention of Jembrana disease (JD) in Bali cattle is currently conducted by vaccination. Until recently, a crude vaccine is prepared from infected

organ of Bali cattle acutely affected by Jembrana disease. The crude vaccine is available for use to prevent the spread of the disease in Bali cattle and has been widely used in many parts of Indonesia. Some disadvantages of the crude vaccine are as follows. It still requires to sacrifice JDV-infected donor cattle in order to prepare the vaccine. As the vaccine is prepared from an infected organ, it is also difficult to instigate a proper quality control for the vaccine. In addition, the vaccine is still expensive to produce. The development



of a recombinant Jembrana vaccine is therefore expected to solve some of the problems associated with disadvantages of the current vaccine.

Jembrana disease viral genome consists of at least three major genes: *pol*, *gag* and *env*. Jembrana Surface Unit (JSU) protein is encoded by *env* gene, while *tat* gene that encodes for JTat protein, is a small accessory gene lies between *pol* and *env* genes. At least two proteins, JSU and JTat, are potential candidates for the development of a protective vaccine against Jembrana disease in Bali cattle (ACIAR, 2004). The JSU is the virus attachment protein which initiates the first interaction between the virus and the infected cells. Antibody capable of binding with the protein will be likely preventing the viral infection (Barnett *et al.*, 2001). Tat protein is a minor protein required by the virus to activate the transcription of the viral genome into mRNA which is necessary for the efficient replication of the virus in the infected host (Cota-Gomez *et al.*, 2002; Wu, 2004). Immune response against the protein is expected to be able to slow down or completely stop the viral replication.

The production of the recombinant proteins for JDV is possible as the genes encoding for the proteins have been well-characterized (Chadwick *et al.*, 1995) which in turn will enable the cloning of the genes in an appropriate expression systems such as pGEX. Cloning of JSU and JTat is normally performed using pGEX system in which the genes were fused with GST tag. Recombinant JSU protein has the molecular weight of  $\pm 34$ kDa and JTat has the molecular weight of  $\pm 10.7$ kDa, whereas the GST tag *per se* has the molecular weight of  $\pm 26$ kDa (Amershampharmacia biotech, 1997). Cloning of both JSU and JTat genes in pGEX system produces fused protein with a GST tag. The fused recombinant proteins, GST-JSU and GST-JTat are expected to have the molecular weights of  $\pm 60$ kDa and  $\pm 36.7$ kDa respectively. Both JSU and JTat recombinant proteins are normally expressed in the form of inclusion body which needs to be solubilized to obtain native proteins. Previous studies have shown that JSU and JTat expression did not result in the right size of molecular weight after characterization by SDS-PAGE and Western blotting (Margawati *et al.*, 2004). It appears that the constructs were not stable or not

well-maintained in glycerol. This study was therefore designed to re-transform both constructs of JSU and JTat into *Escherichia coli* and to express them in pGEX system.

## MATEWALAND METHODS

### Retransformation of Construct

#### Plasmid Isolation of JSU dan JTat

The origin of JSU and JTat pGEX constructs was provided by the virology laboratory of Murdoch University. Plasmid bearing *su* and *tat* genes were collected from JSU and JTat pGEX constructs by miniprep method. Colonies of the JSU and JTat constructs were cultured overnight in Luria Bertani (LB) medium (consists of bacto trypton, yeast extract; NaCl) supplemented with 100µg/ml ampicillin at shaking conditions of 200 rpm at 37°C. Bacterial *E. coli* pellets were collected by centrifugation at 4000 rpm in 10°C environment for 15 min. The collected pellets were lysed with three types buffer solutions, cell lysis buffer, neutralization buffer and centrifuged at 4000 rpm in 10°C environment for 15 min, and the supernatant was collected. Into supernatant was added isopropanol and incubated at -20°C for 2 hours, and was then centrifuged at 4000 rpm at 10°C for 15 minutes to collect pellets. The pellets were washed with 80% ethanol and again centrifuged with 4000 rpm at 10°C for 15 min to collect the pellets (bearing the plasmid). The pellets on the bottom of a micro tube were air dried, and were then diluted in TE buffer.

#### Competency of *Escherichia coli* strain BL21 and DH5a

Competent host cells of *E. coli* (BL21 and dh5oc) are needed for retransformation. A method of TSS (Transformation Storage Solution, consists of LB Medium, PEG6000, DMSO and 1M MgCl<sub>2</sub>) was used to prepare competent *E. coli* (Chung *et al.*, 1989). In prior to competing cells, *E. coli* colony was overnight cultured with shaking of 200 rpm at 37°C and the pellet was then collected by centrifugation of 4000 rpm 4°C for 5 minutes. The collected pellets (*E. coli*) were re-suspended by adding a cold TSS solution. This TS<sup>+</sup> contains *E. coli* competent and ready to be used.

#### Retransformation of JSU and JTat

An amount of 2µl TE buffer (containin

plasmid) was introduced into 100  $\mu$ l of competent cells in TSS by heat shock method for re-transformation. Into the retransformed *E. coli* was added 400  $\mu$ l LB medium without ampicillin and incubated at 37°C with shaking at 200 rpm for 1 hour. The substrate medium was centrifuged with 4000 rpm at 4°C for 1 minute to collect supernatant (SN). The SN was spread out and grown on LB agar plate containing ampicillin.

#### Expression of Protein Overnight Culture

Colonies (containing retransformed JSU and JTat) were overnight cultured into 2-5 ml LB medium containing 100  $\mu$ g/ml ampicillin and incubated at 37°C with shaking of 200 rpm.

#### Inoculation and IPTG Induction

Inoculation was performed with a ratio of 1:20 (overnight culture: LB medium) in 100ml LB medium and incubated at 37°C with shaking of 200 rpm for 3 hours, or up to achieve 0.8 OD (optical density). Induction of 0.1 mM IPTG was implemented after the culture OD of 0.8 was achieved and the culture was kept for another 1 hour. Induction of IPTG was intended to stimulate protein expression (Mills, 2001).

#### Harvesting and Lysis

*E. coli* was harvested after 4 hours in a total culture, including the IPTG induction, by centrifugation of 5000 rpm at 4°C for 15 minutes. Pellet was digested by freezing and thawing method without addition of cell digester (*e.g.* lysozyme) except supplementation of lysis buffer. The freezing and thawing were conducted for 5 cycles each of which consisted of sample exposure (containing pellet and lysis buffer) into liquid nitrogen (N<sub>2</sub>) for 30 seconds and then thawing into water bath at 42°C for 60 seconds. This substrate was then centrifuged with 5000 rpm at 4°C for 15 minutes to collect the pellet (inclusion body).

#### Solubilization of Inclusion Body

Prior to solubilization, the pellet was washed three times by adding washing buffer and centrifuged with 12.000 rpm at 4°C for 15 minutes to collect the pellet. The pellet was overnight solubilized by adding buffer and kept at 4°C (Singh and Panda, 2005).

#### Expression and Purification

The solubilized substrate (semi-purified protein) was purified by using glutathione sepharose 4B with a batch capture method. The purified protein

was analysed by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE), and directly transferred onto nitrocellulose membrane for western blotting (WB) assay. The JSU and JTat proteins were identified by monoclonal anti-GST antibody. The purified protein can be identified by checking their sizes, and using pGEX system JSU has  $\pm$  60kDa while JTat has  $\pm$  36.7kDa.

#### Quantification of Protein

The purified recombinant protein was quantified using a BCA™ Protein Assay Kit with BSA standard (Pierce).

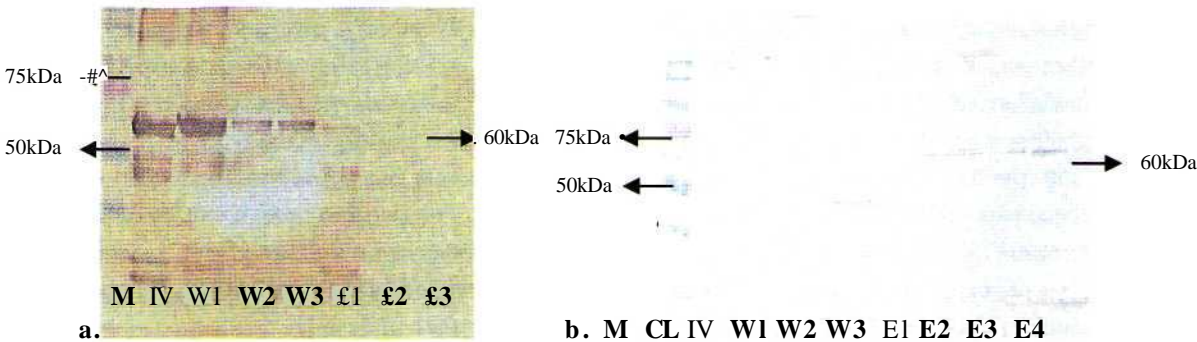
## RESULTS

#### Retransformation JSU and JTat

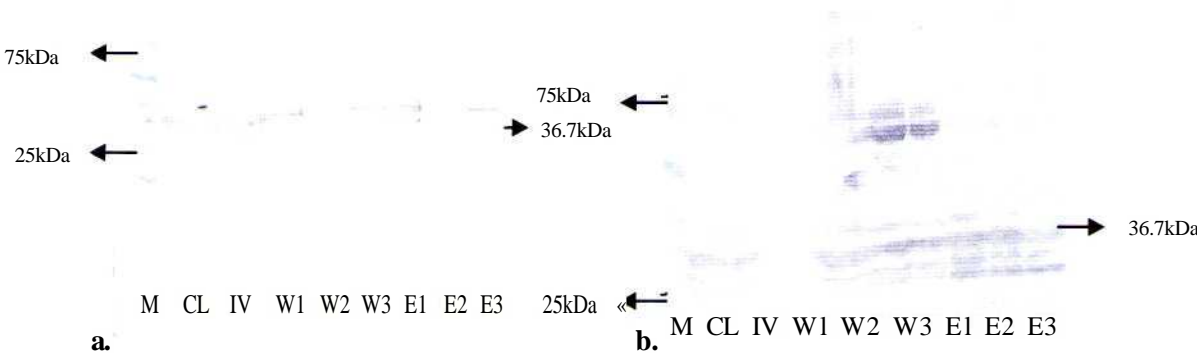
Plasmid containing both *Jembrana SU* and *tat* genes were transformed into *E. coli* strain BL and dH5a. The transformed *E. coli* cells were grown on LB agar plate containing ampicillin. Both transformed *E. coli* (strain BL and dH5oc) containing both JSU and JTat grew and formed colonies. However, the colonies derived from *E. coli* strain BL formed bigger colonies as compared to that colonies derived from dH5a on both JSU and JTat. Therefore, retransformed JSU and JTat derived from BL strain was used for protein expression.

#### Expression of JSU and JTat pGEX

The expression of retransformed JSU and JTat into *E. coli* strain BL was performed three times each of which was in a volume of 100 ml LB culture medium. The solubilized substrate (semi-purified protein) of both JSU and JTat was purified with glutathione sepharose 4B by a batch capture method. JSU and JTat proteins were analysed by SDS-PAGE and then identified by WB using monoclonal anti-GST antibodies. Result of the characterization of both JSU and JTat on WB showed on the right sizes on both retransformed constructs: JSU= $\pm$ 60kDa (Figure 1) and JTat= $\pm$ 36.7kDa (Figure 2). However, the band derived from the retransformed JSU resulted in a sharper and thicker band (Figure 1) as compared to that of JTat (Figure 2) on elution 1,2,3,4 (E1, E2, E3 and E4) of each construct. Moreover, WB of JTat showed very fine bands on E3 and E4.



**Figure 1.** Characterization and identification of retransformed JSU pGEX by western blotting: a. Trial 1 and b. Trial 2. (M= Marker; CL= Cell Lysate; IV= Inner Volume; W1, W2, W3= Washing1, 2, 3; E1, E2, W3, W4 = Elusion 1,2, 3, 4)



**Figure 2.** Characterization and identification of retransformed JTat pGEX by western blotting: a. Trial 1 and b. Trial 2. (M= Marker; CL= Cell Lysate; IV= Inner Volume; W1, W2, W3= Washing1, 2, 3; E1, E2, W3, W4 = Elusion 1,2,3,4)

Protein Concentration

In this study, purified protein was still bound to the glutathion sepharose 4B matrix up to elusion 4 for JSU trial 1 (Figure 1b), whereas the others were eluted up to elusion 3. Therefore, we considered to calculate the protein concentration until elusion 3 (E1, E2 and E3) of those retransformed JSU and JTat pGEX. The protein quantification was repeated twice for each retransformed construct. The average concentration of re-transformed JSU protein was a slightly lower (1.883mg ml<sup>-1</sup>) as compared to that of JTat (1.98 lmg ml<sup>-1</sup>), (Table 1). However, JSU recombinant protein appeared to be better-purified as only a single and sharp band was detected (Figure 1) as compared to JTat (Figure 2). The data on the protein concentration for both recombinant proteins in each elution (E) and each average concentration from two trials were shown at Table 1.

**Table 1.** Concentration of purified JSU and JTat proteins derived from retransformation into *E. coli* strain BL21

Trial	Elution (E)	Cone, (mg ml <sup>-1</sup> )	
		JSU	JTat
Trial 1 (T1)	E1	2,181	2,769 1,5142,835
	E2	1,905	
	E3	1,317	
Average		1,801	2,373
Trial 2 (T2)	E1	1,952	1,967
	E2	2,032	1,507
	E3	1,908	1,289
Average		1,964	1,588
Avearage (T1+T2)		1,883	1,981

DISSCUSION

Retransformation and Expression of JSU dan JTat pGEX

Plasmid JSU and JTat pGEX were transformed into host cells, *E. coli* strain BL21 resulting in bigger

colonies on LB medium agar plate as compared to that of transformed into DH5a. This result showed that *E. coli* strain BL21 is more suitable for use in the protein expression of both JSU and JTat in pGEX system. As reported by Inoue *et al.* (1990), *E. coli* strain DH5a is usually used for plasmid transformation whereas *E. coli* strain BL21 is normally used for protein expression. Similar finding was also reported by Ramos *et al.* (2004) that the *E. coli* strain DH5a was used for all routine cloning experiments while the *E. coli* strain BL21 (DE3) was used for recombinant protein expression. In our previous study, *E. coli* strain JM109 was used for cloning experiment of recombinant protein JTat. It showed a similar result to strain DH5a and grew small colonies in LB agar plate medium. Therefore, both strains DH5a and JM109 are suitable for cloning experiment.

Transformation of plasmid into DH5a or JM109 is mainly intended for cloning purposes that is to check whether or not the plasmid bears the desired gene insert upon screening. Once the cloned plasmid is detected to bear the desired gene insert, it is then retransformed into *E. coli* strain BL.

In this study, the retransformation was performed by using the supernatant (SN) derived from centrifugation after heat shocking and then the SN was grown on LB agar plate medium. As reported in our previous study (Margawati *et al.*, 2006), the supernatant collected after heat shocking of plasmid J-Tat pGEX resulted in a better growing of the construct on LB agar plate medium and even yielded a higher concentration of purified recombinant protein. Another simple protocol was reported by Tu *et al.* (2005) that plasmid transformation or mixture of plasmid and competent cell can be directly spread out and grown on the agar plate medium without centrifugation.

#### **Expression and Concentration of retransformed JSU and JTat**

In this study, constructs of JSU and JTat were re-transformed into *E. coli* BL21 for protein expression. As stated by Inoue *et al.* (1990) and Ramos *et al.* (2004), recombinant protein can be expressed in *E. coli* strain BL21, while *E. coli* strain DH5a is suitable for routinely cloning experiment. Cells were lysed by adding lysis buffer followed by freezing and thawing method.

Basically, the lysis of cells is performed by employing non-ionic detergents to disrupt the cellular membrane after the wall of cells was broken by lysozyme treatment. As reported in the recent finding, cell lysis can be conducted in a simpler method by insertion of a bacteriophage lamda lysis gene cluster into pET22b expression (Jang *et al.*, 2007) to make a simpler and a more efficient process of protein purification.

Expression of recombinant JSU and JTat proteins often resulted in the form of inclusion body which needs further solubilization. Such solubilization can be achieved by adding solubilization buffer to obtain soluble protein (Singh and Panda, 2005) or semi-purified protein. The semi-purified protein was then further purified by glutathione sepharose 4B with a batch capture method. The purity of the proteins was, then checked by SDS-PAGE and WB. Both JSU and JTat recombinant proteins appeared to be expressed properly using pGEX system which resulted in  $\pm 60$ kDa protein for JSU (Figure 1) and  $\pm 36.7$ kDa protein for JTat (Figure 2). The JSU appeared to be expressed better as it produced a sharper and a single band in SDS-PAGE as compared to that JTat. This sharp band of purified JSU was obtained from pooled elution 1, 2, 3 and even up obtained from elution 4 (Figure 1b). In contrast, JTat showed inconsistent band quality (sharp and thick) in every elution following analysis by SDS-PAGE and WB. Expression of JTat recombinant protein was not as clear as the JSU, and it appeared to be due to the quality of the construct. A good and stable JSU construct seems to be established using pGEX system before re-transformation. The JTat, however, appeared to unstable construct. As seen on the visualization of purified protein by western blotting, JSU produced a single band of  $\pm 60$ kDa (Figure 1) while JTat, produced multiple protein bands beside the JTat protein ( $36.7$ kDa *per se* (Figure 2). Consequently, this expression consistently affected the concentration of the purified recombinant protein (retransformed) of JTat in pGEX which showed a slightly higher protein concentration ( $1.981 \text{ mg ml}^{-1}$ ) as compared to that retransformed JSU ( $1.981 \text{ mg ml}^{-1}$ ) (Table 1). The maintenance of constructs, therefore, needs to be taken into account by keeping the constructs in the right lower temperature of  $-70$  up to  $-80^\circ\text{C}$  and retransformed

periodically (monthly or in every a 3-month) for refreshing their micro-environment and keeping their survival.

## SUMMARY

Retransformation of JSU and JTat was successfully conducted into to *E. coli* strain BL21 while transformation into *E. coli* strain DH5a appeared to be better for cloning purposes. Expression of JSU and JTat recombinant proteins in pGEX system resulted in the formation of inclusion body which needs for further solubilization. Both JSU and JTat were on the right sizes when detected by western blot assay which clearly showed 60kDa and 36.7kDa proteins, respectively for JSU and JTat. However, JSU seems to better expressed that Jtat as only a single band but a slightly lower concentration of purified protein as compared to that of JTat.

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