

The 3rd International Conference on Biological Science 2013 (The 3rd ICBS-2013)

MERCURY REDUCTASE ACTIVITY OF AN INDIGENOUS MERCURY RESISTANT BACTERIAL ISOLATE (*Bacillus* sp. S1) FROM KALIMAS-SURABAYA AS A POTENTIAL REDUCING AGENT FOR MERCURIAL ION (HG²⁺)

Enny Zulaika^{1**} and Langkah Sembiring²

¹ Jurusan Biologi-FMIPA, Institut Teknologi Sepuluh Nopember, Surabaya
² Laboratorium Mikrobiologi, Fakultas Biologi, Universitas Gadjah Mada, Yogyakarta
**enny@bio.its.ac.id

ABSTRACT

Mercury reductase activity and Hg²⁺ lowering capacity of a Mercury Resistant Bacteria (MRB) (*Bacillus* sp. S1) which was isolated from Kalimas River of Surabaya, Indonesia were studied. The activity was determined by Mercury Reductase Assay System (MRAS) in a solution mixture contained 50 mM PBS (pH ± 7.0), 0.5 mM EDTA, 200 μ M MgSO4, 0.1% (v/v) ß-merchaptoethanol, 200 μ M NADH2 and 25 mg/L HgCl₂ and one volume of crude extract incubated at room temperature for various interval period of time. Mercury reductase activity was measured spectrophotometrically at 340 nm. One unit of reductase activity was defined as one molar of oxidized NADH₂ produced per total cell per minute. Results of study showed that, the isolate could resist the concentration of HgCl₂ up to 11 mg/L. At 30 minute incubation period at room temperature, the highest mercury reductase activity and the Hg²⁺ lowering capacity was found to be 0.006 unit/109cell and 1.48 mg/L/109cell/minute, respectively with the reduction efficiency of Hg²⁺ to Hg⁰ of 0.18% per minute. Therefore, it can be concluded that the *Bacillus* sp. S1 isolate could be assumed to be exellent mercury bioremediation agent since it was found to be highly mercury resistant and very efficient to reduce cationic mercury (Hg²⁺) to elemental mercury (Hg⁰).

Key words: Kalimas-Surabaya, Bacillus sp. S1, Mercury resistant, Mercury reductase

INTRODUCTION

Kalimas River in Surabaya, East Java province of Indonesia has been contaminated by mercury as indicated by the measured levels of mercury of 6.3 mg/Kg in the samples taken from downstream sediments of the river (Zulaika *et al.*, 2012). The phenomenon was very worrying because mercury may cause neurotoxic effects on humans. However, in natural habitats, there are bacteria that might be able to resist mercury which are known as Mercury Resistant Bacteria (MRB). Such bacteria, possess enzymatic mechanism to reduce divalent mercury (Hg²⁺) into volatile mercury of Hg⁰ (Nascimento & Chartone-Souza, 2003) due to their possession of mercury resistant gene known as *mer-operon*, and *mer*A is one of *mer-operon* gene is located within plasmid or transposon (Davis *et al.*, 2005). Plasmid is likely to carry a certain gene or more that might code for some special characters in bacteria, such as resistance and viability to antibiotics and heavy metals (Nithya *et al.*, 2011).

Mercury reductase is an oxydoreductase which involves NADPH₂ or NADH₂ coenzyme in catalyzing electron replacement (Murray, 1996). As an electron source in oxidation-reduction process, NADH₂ or NADPH₂ are involved to change Hg²⁺ to be volatile Hg⁰ or *vice versa* (Zeroual *et al.*, 2003). The efficiency rate of changing Hg²⁺ to be volatile Hg⁰ by *Pseudomonas putida* SP-1 MRB was reported to be 89% (Zhang *et al.*, 2012) whereas *Hydrogenivirga* sp. 379 128 R1-1 was reported to reduce 150 imole Hg²⁺/10⁶ cell/hour (Freed-

ISSN 2413-0877 © 2015 The Authors.

Published by KnowledgeE Publishing Services This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0)

Selection and Peer-review under responsibility of the 3rd ICBS-2013

Doi http://dx.doi.org/10.18502/kls.v2i1.190

man *et al.*, 2012). A bacterial isolate belong to the genus *Bacillus* isolated from Kalimas-Surabaya was found to be dominant among the MRB isolates, and *Bacillus* sp. S1 was one of them which was highly resistant to HgCl₂ concentration of 11 mg/L (Zulaika., 2013). This research aimed at measuring mercury reductase activity as well as Hg²⁺ lowering capacity of *Bacillus* sp. S1 as an indigenous MRB isolate from Kalimas-Surabaya Indonesia.

MATERIALS AND METHODS

Preparation of enzyme extracts

After 12 hour of incubation periods of culture, the culture of *Bacillus* sp. S1 was harvested by centrifugation at 15000 rpm for 20 minute at 4°C. Crude cell extract was re-suspended in 30 ml Phospate Buffer Saline (PBS), pH \pm 7.0. Cells were then disrupted by sonication with ultrasonic processor at 600 watt and 50% amplitude for 60 seconds, and the resultant homogenate was centrifuged at 15000 rpm for 30 minute at 4°C. The supernatant was collected as crude enzyme extract and then removed into black and dark vial tube (Ogunseitan, 1998) to be used in further experiments.

Mercury Reductase Assay System (MRAS)

Mercury reductase activity was measured in a Mercury Reductase Assay System (MRAS) solution in dark tube contained 50 mM PBS (pH \pm 7.0), 0.5 mM EDTA, 200 ìM MgSO₄, 0.1% (v/v) ß-merchaptoethanol, 200 ìM NADH₂ and 25 mg/L HgCl₂ (Ogunseitan, 1998). One volume of crude extract was added into MRAS solution and incubated at room temperature for various interval duration of 30, 60, 90, 120 and 150 minutes. Mercury reductase activity was measured spectrophotometrically at 340 nm (Takeuchi *et al.*, 1999). A one unit of mercury reductase activity was defined as one molar of oxidized NADH₂ produced per total cell per minute in the assay condition (Zeroual *et al.*, 2003).

RESULTS AND DISCUSSION

The MRB isolate was identified to be member of the genus *Bacillus* and named as *Bacillus* sp. S1 based on key characters (Holt *et al.*, 1994) which were rod cell shape, gram positive, motile, forming endospore, aerobes, chemo-organotroph and catalase positive. (Table 1).

No	Characters	Genus Bacillus	Bacillus sp. S1
1	Road	+	+
2	Gram	+	+
3	Motility	+	+
4	Endospore	+	+
5	Aerobes/ facultative anaerobes	+/+	+/-
6	Chemoorganotroph	+	+
7	Catalase	+	+

Table 1. Identification of Bacillus sp. S1 characters based on key characters of the genus Bacillus

The genus *Bacillus* contains 18 sub-groups which form endospore, gram positive, and road cell shape (Holt *et al.* (1994). According to Maughan dan Auwera (2011), member of the genus *Bacillus* has abundant number in nature. Those strains are very adaptive to

environments by means of character variation. They might be antagonistic, synergistic, pathogenic or even beneficial. Certain *Bacillus* strains could produce antibiotic, insecticide, hence they could be used as bioremediation agent. Another *Bacillus* strain member may be tolerant to higher osmotic pressure. Their optimum growth temperature is around 30 - 45°C (mesophilic).

Bacillus sp. S1 isolate was found to be tolerant to antibiotic, such as ampicillin, tetracycline and chloramphenicol (Zulaika *et al.*, 2012), indicated that *Bacillus* sp. S1 contained a plasmid. In MRB, it has been clear that plasmid codes for one amongst genes coding for heavy metal resistance, and *mer*A is the gene which codes for mercury reductase and therefore they are resistant to mercury and able to reduce Hg²⁺ into Hg⁰ (Kiyono & Pan-Hou, 2006). Mercury reductase activity of *Bacillus* sp. S1 isolate in various incubation period is shown in Table 2.

Periods (minutes) *	Enzyme activity (unit/10 ⁹ cell)	Oxidized NADH₂ (μΜ/10 ⁹ cell /minute)	Reduction of Hg ²⁺ * (mg/L/10 ⁹ cell /minute)	% Reduction of Hg²+ /minute
30	0.0060 ± 0,0003	0.179 ± 0,028	$1.480^{a} \pm 0,19$	0.18
60	0.0033 ± 0,0002	0.146 ± 0,023	$0.487^{b} \pm 0.08$	0.12
90	0.0023 ± 0,0001	0.109 ± 0,018	$0.243^{\circ} \pm 0.04$	0.09
120	0.0019 ± 0,0003	0.100 ± 0,030	$0.167^{d} \pm 0,05$	0.08
150	0.0016 ± 0,0002	0.087 ± 0,025	0.117 ^e ± 0,03	0.07

Table 2. Mercury reductase enzyme activity, oxidized NADH and reduced Hg²⁺

) Incubation periods affected to Hg^{2} reduction. Numbers with different alphabet on beside shown significant differs (p < 0.05; a > b > c > d)

Based on Table 2, the highest mercury reductase activity appeared to be at 30 minute incubation period which was 0.0060 unit/10⁹cell either expressed as oxidized NADH₂ amount 0.221 μ M/10⁹sel/minute or as the amount of reduced Hg²⁺ 1.480 mg/L/10⁹sel/minute as shown in Figure 1. This might be due to the fact that the longer the incubation period, the lower substrate availability and the less intact enzyme quantity, and therefore autaomatically decreased the oxidized NADH₂ production as well as lowered the Hg²⁺ reduction.

Bacillus sp. S1 as MRB was isolated from downstream of Kalimas River of Surabaya with 6.3 mg/Kg level of mercury (Zulaika *et al.*, 2012) hence it was ecpected that it could adapt to survive in medium containing mercury. It was reported that indigenous MRB isolate from contaminated mercury area were found to be much more adaptive to Hg^{2+} than the isolate originated from incontaminated area due to its effectiveness in reducing Hg^{2+} (Ghoshal *et al.*, 2011). Therefore, the isolate of *Bacillus* sp. S1 could be expected to be very potential to be used as mercury bioremediation agent. This isolate was exellent in terms of mercury reductase enzyme productivity, and capacity to reduce Hg^{2+} to Hg^0 by means of oxidation of NADH₂. The isolate could resist the concentration of $HgCl_2$ up to 11 mg/L. The highest mercury reductase enzyme activity (0.006 unit/10⁹cell) and the Hg^{2+} lowering capacity of 1.48 mg/L/10⁹cell/minute with 0.18% per minute reduction efficiency of Hg^{2+} to Hg^0 was observed at 30 minute incubation period at room temperature.

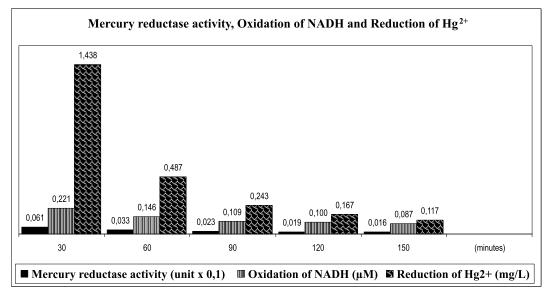


Figure 1. Mercury reductase activity (unit x 0,1), Oxidation of NADH₂ (μM) and Reduction of Hg²⁺ (mg/L/10⁹cell/minute) by *Bacillus* sp. S1 isolate.

CONCLUSION

Bacillus sp. S1 isolate could be assumed to be exellent mercury bioremediation agent since it was found to be highly mercury resistant. The isolate could resist the concentration of $HgCl_2$ up to 11 mg/L. At 30 minute incubation period at room temperature, the highest mercury reductase activity and the Hg²⁺ lowering capacity was found to be 0.006 unit/10⁹ cell and 1.48 mg/L/10⁹ cell/minute, respectively with the reduction efficiency of Hg²⁺ to Hg⁰ of 0.18% per minute.

REFERENCES

- Davis, I.J., A.P. Robert, D. Ready, H. Richards, M. Wilson, and P. Mullany. 2005. Linkage of a Novel Mercury Resistance Operon with Streptomycin Resistance on a Conjugative Plasmid in *Enterococcus faecium. Plasmid*, 54: 26 - 38.
- Freedman, Z., C. Zhu, and T. Barkay. 2012. Mercury Resistance, Mercuric Reductase Activities and Expression Among Chemotrophic Thermophilic Aquificae. *Applied Environmental Microbiology*, 10: 1-34.
- Ghoshal, S., P. Bhattacharya, and R. Chowdhury. 2011. De-mercurization of Wastewater by Bacillus cereus (JUBT1): Growth Kinetics, Biofilm Reactor Study and Field Emission Scanning Electron Microscopic Analysis. Journal of Hazardous Materials, 194: 355 -361.
- Holt, J.G., N.R. Krieg, H.A. Sneath, J.T. Staley, and S.T. Wiliam. 1994. *Bergey's Manual of Determinative Bacteriology* 9th Ed, William & Wilkins, Baltimore.
- Kiyono, M., and H. Pan-Hou. 2006. Genetic Enginering of Bacteria for Environmental Remediation of Mercury. *Journal of Health Science*, 52 (3): 199 204.
- Maughan, H., and G.V. Auwera. 2011. *Bacillus* Taxonomy in The Genomic Era Finds Phenotypes to be Essential Though Often Misleading. *Infection, Genetics and Evolution*, 11: 789 - 797.
- Murray, M. 1996. The Encyclopaedia of Nutritional Supplements. P. 44-53

- Nascimento, A.M.A., and E. Charton-Souza. 2003. Operon *mer*. Bacterial Resistance to Mercury and Potential for Bioremediation of Contamined Environments. *Journal Genetics and Molecular Research*, 2(1): 92 101.
- Nithya, C., B. Gnanalakshmi, and .K. Pandian. 2011. Assessment and Characterization of Heavy Metal Resistance in Palk Bay Sediment Bacteria. *Marine Environmental Research*, 71: 283 - 294.
- Ogunseitan, O.A. 1998. Protein Method for Investigating Mercuric Reductase Gene Expression in Aquatic Environments. *Applied Environmental Microbiology*, 64(2): 695-702.
- Takeuchi, F., K. Iwahori, T. Maeda, K. Kamimura, and T. Sugio. 1999. Isolation and Some Properties of the Ferrooxidans Strains with Differing Levels of Mercury Resistance from Natural Environments. *Journal of Bioscience and Bioenginnering*, 88(4): 387 - 392.
- Zeroual, Y. A., F.Z. Moutaouakkil, M. Dzairi, P.U. Talbi, K.L. Chung, and M. Blaghen. 2003. Purification and Characterization of Cytosolic Mercuric Reductase from *Klebsiella pneumonia*. *Annals of Microbiology*, 53: 149 -160.
- Zhang, W., L. Chen, and D. Liu. 2012. Characterization of a marine-isolated Mercuryresistant *Pseudomonas putida* strain SP1 and its potential application in marine Mercury reduction. *Applied Microbiology and Biotechnology*. 93: 1305-1314.
- Zulaika, E., L. Sembiring, and A. Soegianto. 2012. Characterization and dentification of Mercury-resistant Bacteria from Kalimas River Surabaya-Indonesia by Numerical Phenetic Taxonomy. *Journal Basic Applied Science Research*, 2(7): 7263 7269.
- Zulaika, E., 2013. *The Exploration of Indigenous Mercury-Resistant Bacteria in Kalimas, Surabaya as an Agent of Bioremediation of Mercury Pollution*. Disertasi. Fakultas Sains & Teknologi. Airlangga University, Surabaya.