

Environmental Parameters Affecting Urease Production and Ammonification in *Phaseolus vulgaris*-Nodulating Rhizobia and *Vigna radiata*-Nodulating Rhizobia

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Abstract: Twenty three isolates of symbiotic nitrogen-fixing bacteria that nodulate common bean (*Phaseolus vulgaris*) or mungbean (*Vigna radiata*) were examined for their urease activity and ammonification. All isolates were observed to produce extracellular urease that ranged between 5.458 ± 1.021 to 28.326 ± 5.618 unit/ml supernatant. The ammonia produced from ammonification of peptone ranged between 70 ± 15.564 to 266.354 ± 23.833 μM . A different trend between urease production of common bean-nodulating rhizobia and mungbean-nodulating rhizobia was observed. In the case of 2 isolates of common bean-nodulating rhizobia, hydrolysis of urea slightly increased during 2 days of culture and the most increase was found on the third day. After the third day, urease activity was markedly decreased. Maximum production of urease was detected during the mid-log to late-log phases of growth. The production of urease by 2 isolates of mungbean-nodulating rhizobia was not significantly different during 2 days of culture, then it was continuously increased until the last day of cultivation in which cells were in their stationary phases. Similar profiles were observed from ammonification of 3 isolates of common bean-nodulating rhizobia and 3 isolates of mungbean-nodulating rhizobia. Ammonification, as measured by the amount of ammonia produced, increased along with growth until cells reached the stationary phases. Effects of factors on production of urease and ammonification of the selected isolates were studied. Effects of these factors varied in different isolates. In summary, Ni^{2+} , Co^{2+} and NaNO_3 decreased urease production in common bean-nodulating rhizobia. While, yeast extract and NH_4Cl decreased urease production in mungbean-nodulating rhizobia. Two heavy metals, Co^{2+} and Zn^{2+} , had negative effects on ammonification in almost all isolates tested.

Key words: Ammonification • Rhizobia • urease

INTRODUCTION

The specific groups of bacteria, collectively known as rhizobia, induce the formation of root or stem nodules of leguminous plants and establish a nitrogen-fixing symbiosis [1, 2]. Nitrogen fixation that is catalyzed by nitrogenase enzyme, is a reduction of nitrogen gas into ammonia [3]. This process can provide a source of assimilable nitrogen for bacteria and plants. Besides nitrogen fixation, ammonia can be obtained from the other metabolic processes such as urease production and ammonification. Urease is a nickel-containing, multi subunit enzyme that catalyzes the hydrolysis of urea to ammonia and carbonic acid [4]. Urease expression in bacteria is often regulated in response to environmental parameters such as nitrogen availability and pH [5]. Bacteria perform ammonification with the production of ammonia from nitrogen compounds such as nitrate [6], casamino acids and trypticase peptone [7].

Several factors were found to affect urease production in bacteria. Addition of nickel to a nickel-free culture led to recovery of urease activity of the cyanobacterium *Anabaena cylindrica* [8]. Supplementation of nickel in medium could result in up to 3.5-fold-increased expression of the urease subunit proteins UreA and UreB of *Helicobacter pylori* [9]. Acetohydroxamic acid (100 $\mu\text{g/ml}$), hydroxyurea (85 $\mu\text{g/ml}$), flurofamide (0.05 $\mu\text{g/ml}$) and EDTA (8 mM) inhibit enzyme activity of *Campylobacter pylori* by 50% [10]. Temperature and urea were found to affect urease gene expression in *Proteus mirabilis* [11]. The effects of some factors on ammonification by bacteria have been investigated; trichloroethylene (TCE) concentration has no effect on arginine ammonification potential of soil microbial populations [12]. Also, it was found that the effects of aldrin on ammonification of peptone are minor and irregular [13].

Table 1: Urease activity and ammonification of rhizobia used in this study

| Host plants | Rhizobial isolates | Urease activity | | Amount of ammonia (μM) produced from ammonification |
|---|--------------------|--|-------------------------------------|---|
| | | Extracellular urease (U/ml supernatant) | Intracellular urease (U/mg cell) | |
| <i>Phaseolus vulgaris</i> (common bean) | PV001 | 10.634 \pm 1.574 | 0.365 \pm 0.093 | 119.219 \pm 23.202 |
| | PV002 | 14.869 \pm 2.367 | 0.544 \pm 0.006 | 139.688 \pm 27.194 |
| | PV006 | 12.799 \pm 0.454 | 0.996 \pm 0.034 | 147.083 \pm 32.991 |
| | PV008 | 26.327 \pm 0.100 | 0.775 \pm 0.063 | 169.688 \pm 72.138 |
| | PV009 | 17.504 \pm 4.908 | 0.554 \pm 0.070 | 187.396 \pm 31.961 |
| | PV010 | 8.846 \pm 0.918 | 0.302 \pm 0.046 | 161.146 \pm 6.5120 |
| | PV011 | 19.762 \pm 1.835 | 0.224 \pm 0.036 | 82.083 \pm 10.723 |
| | PV012 | 19.763 \pm 1.956 | 0.291 \pm 0.036 | 113.229 \pm 16.514 |
| | PV013 | 28.326 \pm 5.618 | 0.376 \pm 0.206 | 162.084 \pm 56.047 |
| | PV014 | 27.949 \pm 2.834 | 0.426 \pm 0.029 | 70.104 \pm 15.564 |
| | PV015 | 11.528 \pm 6.057 | 0.255 \pm 0.017 | 115.105 \pm 26.475 |
| | PV016 | 22.515 \pm 3.693 | 0.412 \pm 0.077 | 187.500 \pm 65.188 |
| | PV017 | 21.221 \pm 1.200 | 0.598 \pm 0.102 | 163.230 \pm 23.396 |
| | PV018 | 23.809 \pm 1.765 | 0.567 \pm 0.055 | 113.229 \pm 80.931 |
| | DASA 06008 | 12.846 \pm 1.230 | 0.440 \pm 0.036 | 99.167 \pm 63.987 |
| | DASA 06006 | 14.822 \pm 1.474 | 0.529 \pm 0.017 | 173.959 \pm 22.418 |
| <i>Vigna radiata</i> (mungbean) | DASA 02007 | 6.775 \pm 2.473 | 0.351 \pm 0.042 | 106.979 \pm 35.842 |
| | DASA 02008 | 7.482 \pm 1.156 | 0.468 \pm 0.083 | 199.167 \pm 36.406 |
| | DASA 02009 | 11.010 \pm 1.717 | 0.481 \pm 0.205 | 252.709 \pm 29.359 |
| | DASA 02010 | 17.269 \pm 5.364 | 0.357 \pm 0.087 | 223.750 \pm 64.939 |
| | DASA 02011 | 5.458 \pm 1.021 | 0.199 \pm 0.009 | 218.230 \pm 21.940 |
| | DASA 02068 | 12.704 \pm 2.136 | 0.617 \pm 0.021 | 233.750 \pm 59.722 |
| | DASA 02077 | 6.214 \pm 1.114 | 0.349 \pm 0.022 | 266.354 \pm 23.833 |

The values shown are the mean values of triplicates \pm standard deviations

In this study, we investigated urease production and ammonification by symbiotic nitrogen-fixing bacteria that nodulate common bean (*Phaseolus vulgaris*) or mungbean (*Vigna radiata*). Sixteen isolates of common bean-nodulating rhizobia and 7 isolates of mungbean-nodulating rhizobia were screened for their urease activity and ammonification.

MATERIALS AND METHODS

Rhizobia and Culture Conditions: The 23 rhizobial isolates used in this study and their host plants are listed in Table 1. Nodulation tests and random amplified polymorphic DNA (RAPD) analysis of mungbean-nodulating rhizobia were carried out in the previous study [14]. These strains nodulated mungbean and generated particular RAPD profiles, suggesting that they were individual strains. Nodulation tests of common bean-nodulating rhizobia were carried out in this study. Yeast-Mannitol (YM) medium [15] was used for growth and maintenance unless otherwise indicated.

Measurement of Urease Activity: Bacteria were grown in Christensen's urea broth [16]. Bacterial cultures were incubated at 28°C with shaking at 150 rpm for 3-5 days until cultures reached the mid-log phases. Then the cultures were centrifuged at 10,000 rpm for 10 min. Cell pellets were sonicated for 10 min and resuspended in distilled water. The sonicated cell pellets were used to measure intracellular and/or surfaced-associated urease activity and the supernatants were used to measure extracellular urease activity. Urease activity was measured as described by Askin and Kizilkaya [17] with some modifications. A 3.0 ml of citrate buffer (pH 6.7) and 5.0 ml of 10% urea substrate solution were added to 1.0 ml of supernatant or sonicated cell suspension and the sample was then incubated for 3 h at 37°C. Then 5.0 ml of sodium phenolate (12.5% (w/v) phenol + 5.4% (w/v) NaOH) and 1.0 ml of 0.9% sodium hypochloride were added to the sample. The absorbance at 578 nm of the reaction mixture was measured. Controls without addition of urea were also included. Ammonia concentrations without and with addition of urea were used to determine the released ammonia resulting from urease activity.

The concentration of ammonia was determined by comparison with a standard curve. A standard curve was constructed by preparing reaction mixtures in which ammonium chloride at varied concentration (equivalent to 0-12 mg/ml ammonia) was used in instead of the supernatant or sonicated cell suspension. One unit of urease activity is equal to the amount of urease that hydrolyzes 0.5 μmol of urea and liberates 1.0 μmol of ammonia in 1 min at 37°C [18]. Based on the maximum activity, 2 isolates specific to each host plant were selected for further studies.

Measurement of Ammonification: Bacteria were propagated in peptone water and incubated at 28°C with shaking at 150 rpm for 3-5 days until cultures reached the mid-log phases. A 0.1 ml of Nessler's reagent was added into 2 ml of the culture [19]. The absorbance at 395 nm was immediately measured [20]. Ammonium chloride in the range of 0-500 μM was used in the reaction to prepare a standard curve. The concentration of ammonia produced was determined by comparison with a standard curve. Based on the maximum amount of ammonia produced, 3 isolates specific to each host plant were selected for further studies.

Urease Production and Ammonification During Growth of the Selected Rhizobia: To examine urease production and ammonification during growth of the selected rhizobia, cells were grown in YM broth at 28°C with shaking at 150 rpm for 3-5 days and used as inocula. The inocula were added into Christensen's urea broth (for measurement of urease activity) and peptone water (for measurement of ammonification) to obtain the initial cell number 1.00×10^5 Colony Forming Unit (CFU)/ml. Cultures were grown at 28°C with shaking at 150 rpm for 5 days. At a 1-day interval, the cell numbers in media were measured by the standard plate count method on YM agar containing 25 $\mu\text{g/ml}$ congo red, urease activity and ammonification were determined as described above.

Effects of Environmental Parameters on Urease Production, Ammonification and Growth of the Selected Rhizobia: Factors tested for urease production were inoculated into Christensen's urea broth including yeast extract 1 g/l; peptone 5 g/l; NH_4Cl 10 mM; NaNO_3 10 mM; NiSO_4 0.1 mM and CoCl_2 0.1 mM. Factors tested for ammonification were added to peptone water

including CuSO_4 0.2 mM; CuSO_4 2.0 mM; ZnSO_4 0.2 mM and ZnSO_4 2.0 mM. The inocula were prepared and the cultures were grown as described above. The cell numbers, urease activity and ammonification were measured on the day that each isolate had maximum urease activity or ammonification.

Statistical Analysis: The standard deviation was calculated using Microsoft Excel 2003 software.

RESULTS AND DISCUSSION

Measurement of Urease Activity: Urease activity was measured in cell-free supernatants (for extracellular urease) and disrupted cell pellets (for intracellular and/or surfaced-associated urease) (Table 1). Among the 23 rhizobia tested, all of them produced extracellular but almost no intracellular urease. Extracellular urease activity ranged between 5.458 ± 1.021 to 28.326 ± 5.618 U/ml supernatant, while intracellular urease activity ranged between 0.199 ± 0.009 to 0.996 ± 0.034 U/mg cell. Some other bacteria such as *Lactobacillus* sp. and *Helicobacter pylori* were found to produce extracellular urease [21, 22]. Five isolates of common bean-nodulating rhizobia (PV008, PV013, PV014, PV016 and PV018) produced the highest urease activity, which was not significantly different. Two isolates specific to each host plant were selected based on the maximum unit of extracellular urease/ml supernatant. Common bean-nodulating rhizobia (PV013 and PV014) as well as mungbean-nodulating rhizobia (DASA 02010 and DASA 02068) were selected for further studies.

Measurement of Ammonification: Ammonification was measured from the amount of ammonia produced from peptone. The results are shown in Table 1. The ammonia produced from rhizobial isolates ranged between 70 ± 15.564 to 266.354 ± 23.833 μM . One isolate of common bean-nodulating rhizobia (PV016) and 4 isolates of mungbean-nodulating rhizobia (DASA 02009, DASA 02010, DASA 02068 and DASA 02077) produced the highest amount of ammonia, which was not significantly different. Common bean-nodulating rhizobia (PV009, PV016 and DASA 06006) as well as mungbean-nodulating rhizobia (DASA 02009, DASA 02068 and DASA 02077) were selected for further studies.

The values shown are the mean values of 3 replicates. Error bars indicate standard deviations

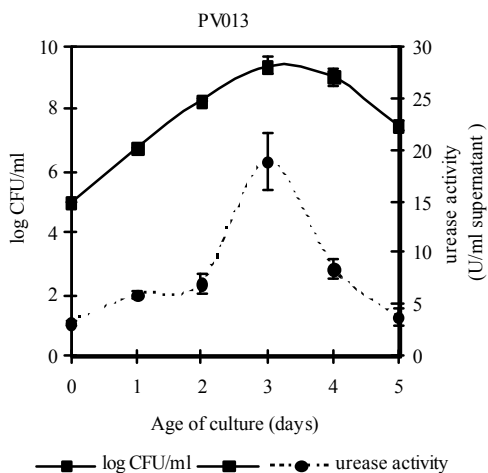


Fig. 1: Growth curve and urease production of the isolate PV013

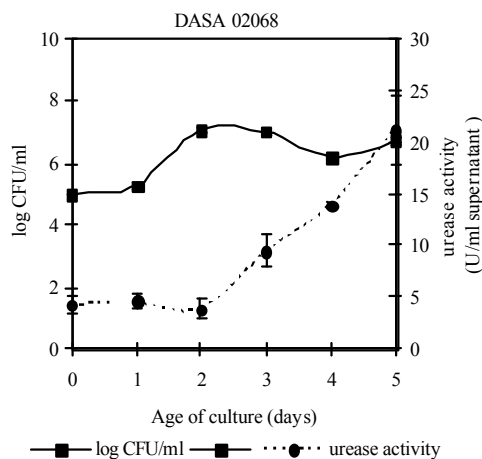


Fig. 4: Growth curve and urease production of the isolate DASA 02068

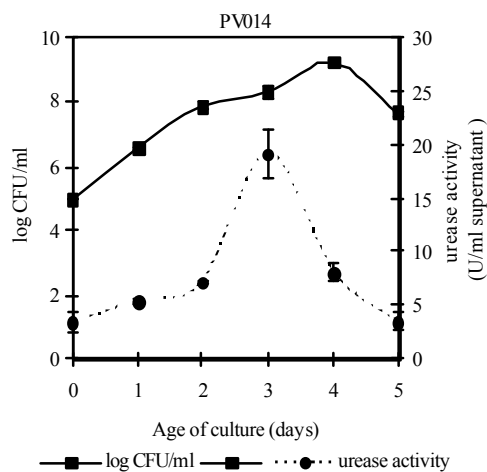


Fig. 2: Growth curve and urease production of the isolate PV014

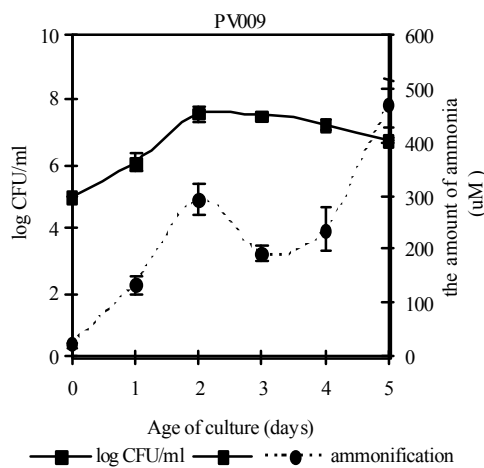


Fig. 5: Growth curve and ammonification of the isolate PV009

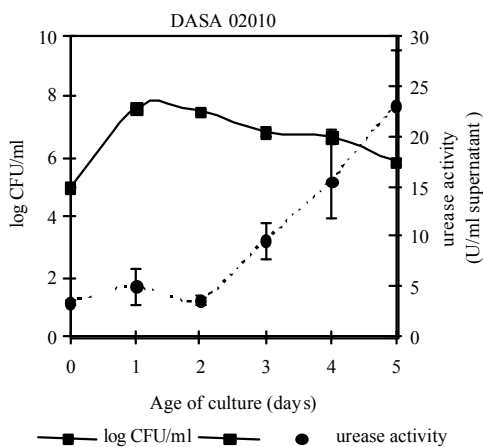


Fig. 3: Growth curve and urease production of the isolate DASA 02010

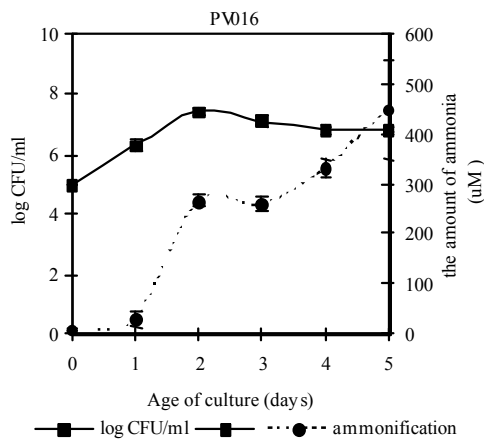


Fig. 6: Growth curve and ammonification of the isolate PV016

The values shown are the mean values of 3 replicates. Error bars indicate standard deviations

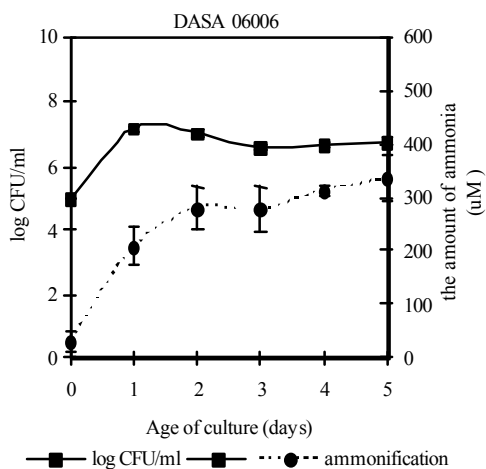


Fig. 7: Growth curve and ammonification of the isolate DASA 06006

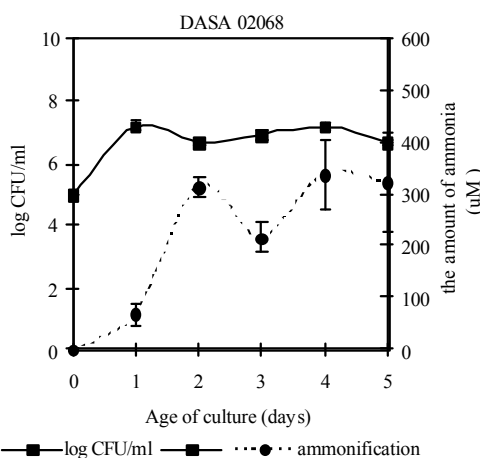


Fig. 9: Growth curve and ammonification of the isolate DASA 02068

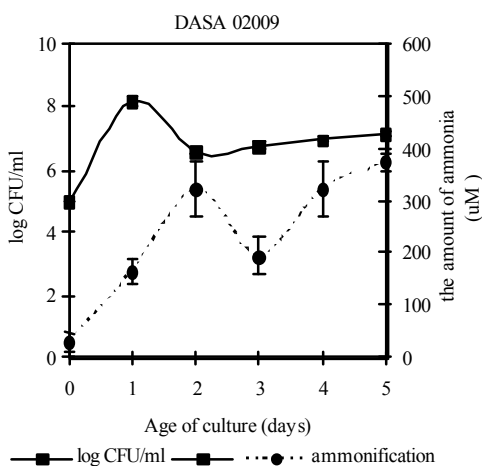


Fig. 8: Growth curve and ammonification of the isolate DASA 02009

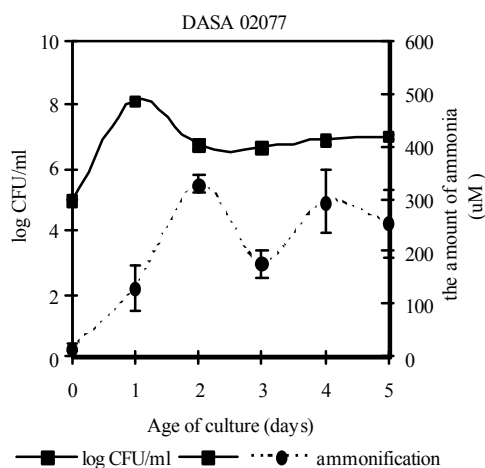


Fig. 10: Growth curve and ammonification of the isolate DASA 02077

Urease Production and Ammonification During Growth of the Selected Rhizobia: As only extracellular urease was detected in all rhizobia tested, it was measured during growth of the selected rhizobia (Figure 1-4). In case of common bean-nodulating rhizobia (PV013 and PV014), hydrolysis of urea slightly increased during 2 days of culture and the maximum increase was found on the third day. After the third day, urease activity decreased markedly. Maximum production of urease was detected during the mid-log to late-log phases of growth. The production of urease by mungbean-nodulating rhizobia was not significantly different, during the first 2 days of culture, (then it continuously increased until the last day of cultivation in which cells were in their stationary phases). In the previous

study, a parallel difference between the two strains with regard to the onset of ureolysis was observed, hydrolysis beginning at the onset of growth in the case of *Actinobacillus* and only at the end of the exponential growth phase in the case of *Lactobacillus* [21]. Figure 5-10 present the growth curves and ammonification of the isolates PV009, PV016, DASA 06006, DASA 02009, DASA 02068 and DASA 02077, respectively. Similar profiles were observed from ammonification of common bean-nodulating rhizobia and mungbean-nodulating rhizobia. The amount of ammonia increased along with growth until cells reached the stationary phases, after that it still maintained or decreased during the stationary phases, then increased again at the end of the stationary phases.

The values shown are the mean values of 3 replicates. Error bars indicate standard deviations

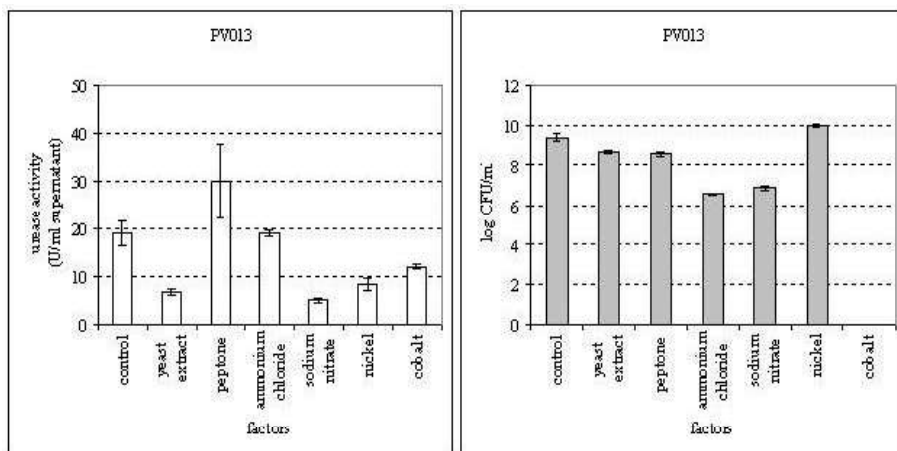


Fig. 11: The influence of environmental factors on urease activity and growth of the isolate PV013

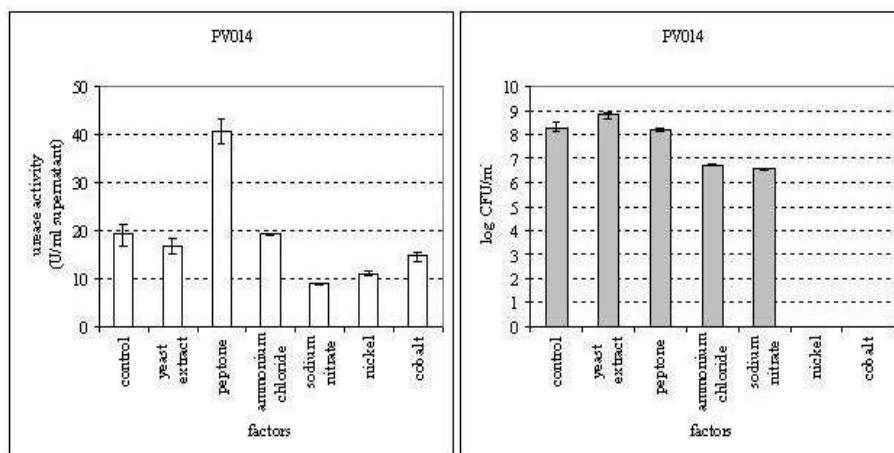


Fig. 12: The influence of environmental factors on urease activity and growth of the isolate PV014

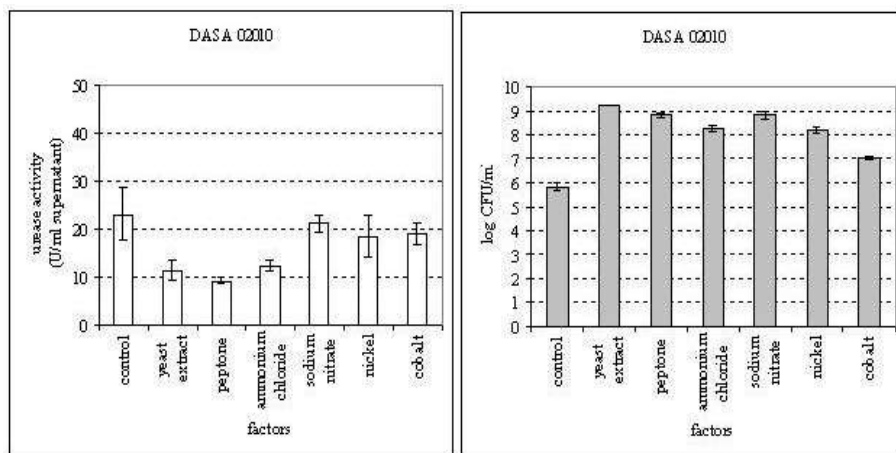


Fig. 13: The influence of environmental factors on urease activity and growth of the isolate DASA 02010

The values shown are the mean values of 3 replicates. Error bars indicate standard deviations

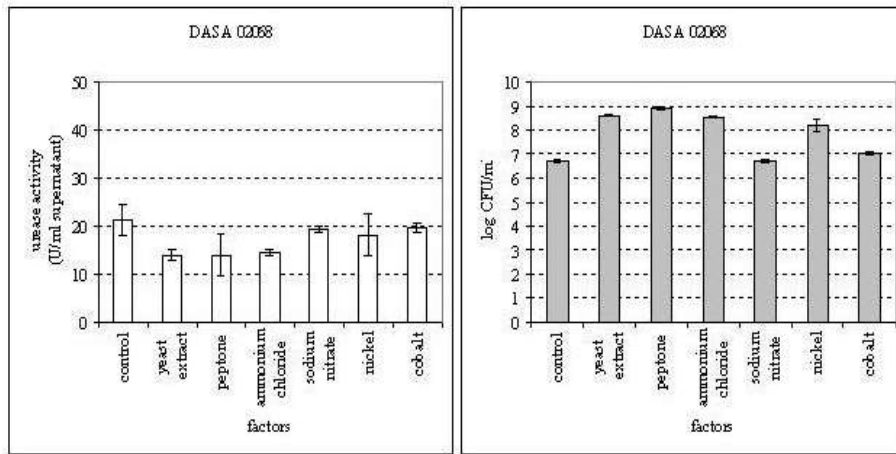


Fig. 14: The influence of environmental factors on urease activity and growth of the isolate DASA 02068

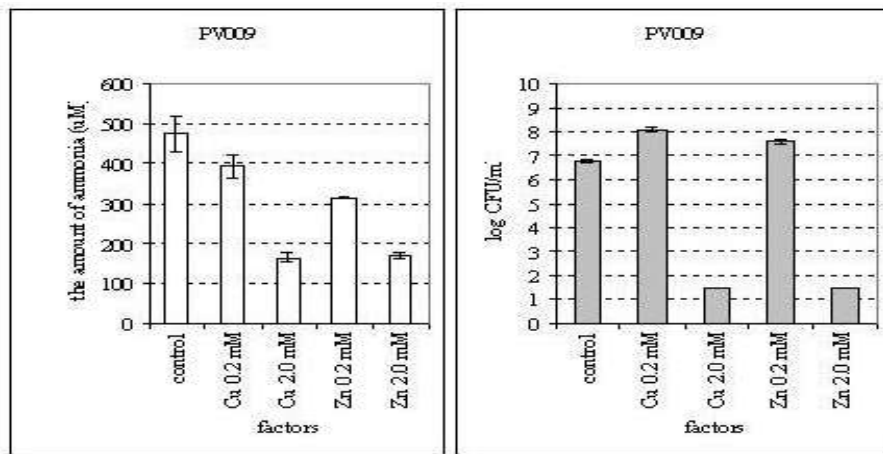


Fig. 15: The influence of environmental factors on ammonification and growth of the isolate PV009

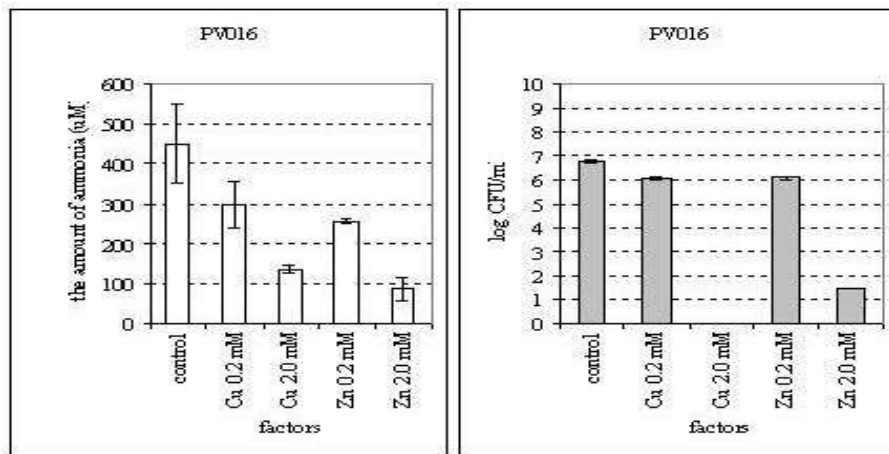


Fig. 16: The influence of environmental factors on ammonification and growth of the isolate PV016

The values shown are the mean values of 3 replicates. Error bars indicate standard deviations

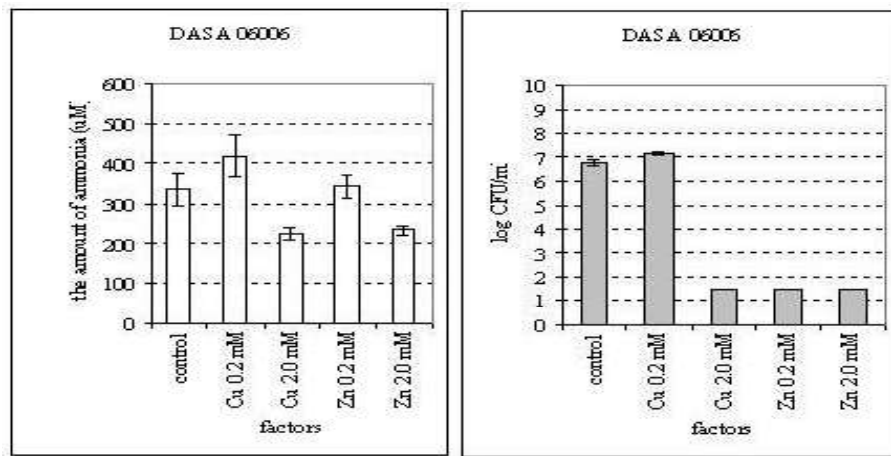


Fig. 17: The influence of environmental factors on ammonification and growth of the isolate DASA 06006

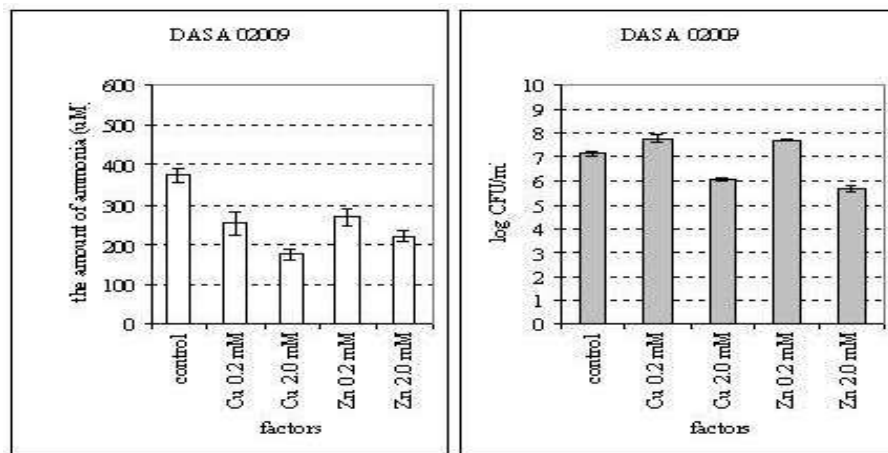


Fig. 18: The influence of environmental factors on ammonification and growth of the isolate DASA 02009.

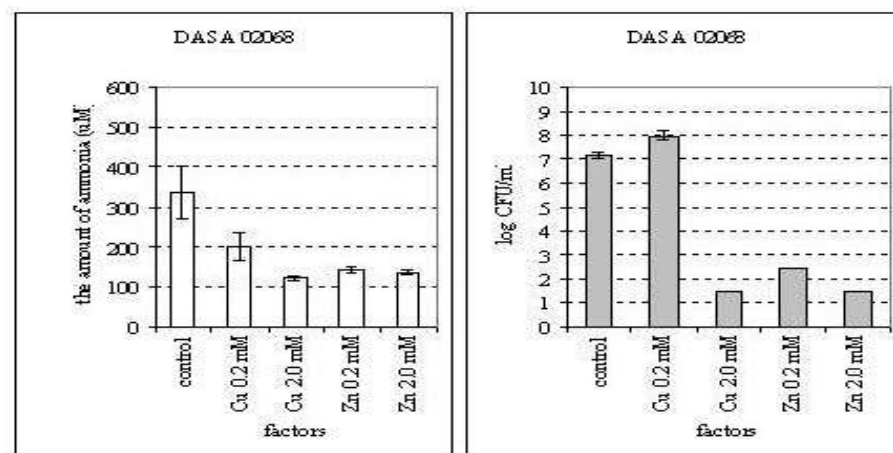


Fig. 19: The influence of environmental factors on ammonification and growth of the isolate DASA 02068

The values shown are the mean values of 3 replicates. Error bars indicate standard deviations

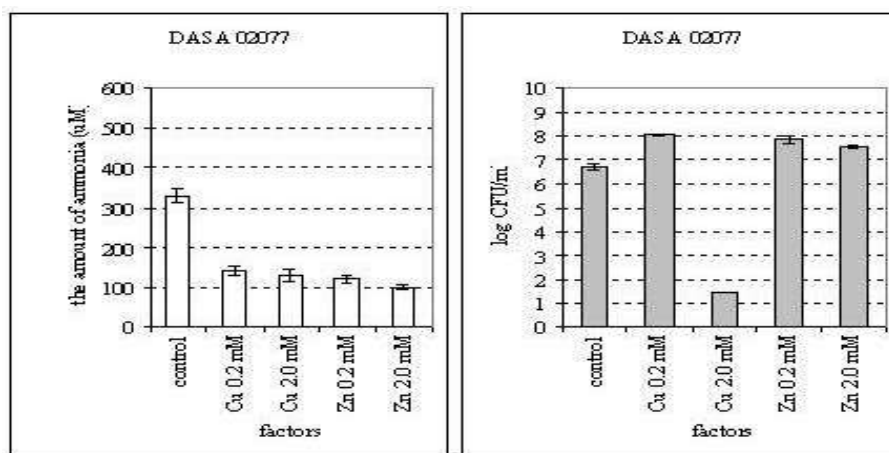


Fig. 20: The influence of environmental factors on ammonification and growth of the isolate DASA 02077

Effects of Environmental Parameters on Urease Production, Ammonification and Growth of the Selected Rhizobia:

Previous studies showed that nitrogen availability is found as one of environmental parameters that regulate urease expression in bacteria [5] and urease is a nickel-containing enzyme [4]. The urease activity of the cyanobacterium, *Anabaena cylindrica*, was also found to be nickel-dependent and the addition of nickel to a nickel-free culture led to recovery of urease activity [8]. The cobalt salt has been reported as a reversible inhibitor of intracellular urease produced by *Rhizopus oryzae* [23]. Figures 11-14 present the effects of these factors on urease production and growth of the isolates PV013, PV014, DASA 02010 and DASA 02068, respectively. It was found that effects of these factors varied in different isolates. As compared with controls, Ni²⁺, Co²⁺ and NaNO₃ decreased urease production in both isolates of common bean-nodulating rhizobia (PV013 and PV014). The greatest decrease of urease production was found with NaNO₃. Peptone was found to stimulate urease production in both common bean-nodulating isolates. In contrast, it decreased urease production in mungbean-nodulating isolate DASA 02010 and no significant difference was found with mungbean-nodulating isolate DASA 02068. Yeast extract decreased urease production of all isolates except PV014. The variation of these factors might be due to different nitrogen catabolic pathways in each isolate [24]. Heavy metals were found to inhibit growth of some isolates, leading to a decrease in urease production. Both common bean-nodulating isolates could not tolerate Co²⁺. The growth of a common bean-nodulating

isolate, PV014, was inhibited by Ni²⁺. Although Ni²⁺ did not have a growth-inhibitory effect on another isolate (PV013), it had a negative effect on urease production. In contrast, Ni²⁺ and Co²⁺ did not have a growth-inhibitory effect on both mungbean-nodulating isolates (DASA 02010 and DASA 02068). The urease production of these isolates was not affected by Ni²⁺ and Co²⁺. Regardless of a growth-inhibitory effect, it was found that 0.1 mM Ni²⁺ decreased urease production in the isolate PV013, but not in the other isolates (DASA 02010 and DASA 02068). Even though Ni²⁺ is the cofactor of the urease enzyme such as in *Helicobacter pylori* [25], it is toxic for some bacterial cells such as *Escherichia coli* [26]. Uncontrolled acquisition of transition metals like nickel may lead to toxicity, as they may participate in the generation of toxic oxygen radicals or block incorporation of cofactors into enzymes [27]. Bacterial survival in the presence of toxic compounds is dependent on resistance and tolerance mechanisms [26].

Factors tested for ammonification include 1) CuSO₄ 0.2 mM; 2) CuSO₄ 2.0 mM; 3) ZnSO₄ 0.2 mM; and 4) ZnSO₄ 2.0 mM. The effect of these heavy metals on ammonification of soil has been reported. Under aerobic incubation conditions, 100 ppm Cu²⁺ was found to increase ammonification of soil but 1,000 ppm Cu²⁺ had no effect. The 100 and 10,000 ppm Zn²⁺ levels reduced ammonification whilst 1,000 ppm Zn²⁺ did not affect ammonification [28]. Figure 15-20 present the effects of environmental factors on ammonification and growth of the isolates PV009, PV016, DASA 06006, DASA 02009, DASA 02068 and DASA 02077, respectively.

The Cu and Zn heavy metals, added as sulphate, at varied concentrations (0.2 mM and 2.0 mM) had a negative effect on ammonification as these heavy metals are toxic to cells. Regardless of a growth-inhibitory effect, 0.2 mM Cu²⁺ reduced ammonification of the isolates PV009, DASA 02009, DASA 02068 and DASA 02077 (by 17% to 57%). The 0.2 mM Zn²⁺ reduced ammonification of the isolates PV009, DASA 02009 and DASA 02077 (by 28% to 63%).

In conclusion, common bean-nodulating rhizobia and mungbean-nodulating rhizobia produced extracellular urease, with maximum production achieved during the mid-log phases to stationary phases of growth. Ammonification of peptone varied among these rhizobia. The effect of other nitrogen sources (yeast extract, peptone, NH₄Cl and NaNO₃) and heavy metals (Cu²⁺ and Zn²⁺) on urease production and cell growth varied in different isolates whilst Co²⁺ and Zn²⁺ had negative effects on ammonification and cell growth of almost all isolates tested.

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REFERENCES

- Hartmann, A. and N. Amarger, 1991. Genotypic diversity of an indigenous *Rhizobium meliloti* field population assessed by plasmid profiles, DNA fingerprinting and insertion sequence typing. *Can. J. Microbiol.*, 37: 600-608.
- Pongsilp, N. and N. Boonkerd, 2007. Research techniques for estimating phenotypic and genotypic diversity of root- and stem-nodule bacteria. *Suranaree J. Sci. Technol.*, 14: 297-308.
- Kaminski, P.A., J. Batut and P. Boistard, 1998. A survey of symbiotic nitrogen fixation by rhizobia. In *The Rhizobiaceae: Molecular biology of model plant-associated bacteria*, Eds., Spaink, H.P., A. Kondorosi and P.J.J. Hooykaas. Kluwer Academic Publishers, pp: 431-460.
- Morou-Bermudez, E. and R.A. Burne, 1999. Genetic and physiologic characterization of urease of *Actinomyces naeslundii*. *Infect. Immun.*, 67: 504 - 512.
- Mobley, H.L., M.D. Island and R.P. Hausinger, 1995. Molecular biology of microbial ureases. *Microbiol. Rev.*, 59: 451-480.
- Hoffmann, T., N. Frankenberg, M. Marino and D. Jahn, 1998. Ammonification in *Bacillus subtilis* utilizing dissimilatory nitrite reductase is dependent on *resDE*. *J. Bacteriol.*, 180: 186-189.
- McSweeney, C.S., B. Palmer, R. Bunch and D.O. Krause, 1999. Isolation and characterization of proteolytic ruminal bacteria from sheep and goats fed the tannin-containing shrub legume *Calliandra calothyrsus*. *Appl. Envir. Microbiol.*, 65: 3075-3083.
- Mackerras, A.H. and G.D. Smith, 1986. Urease activity of the cyanobacterium *Anabaena cylindrica*. *J. Gen. Microbiol.*, 132: 2749-2752.
- Van Vliet, A.H.M., E.J. Kuipers, B. Waidner, B.J. Davies, N. De Vries, C.W. Penn, C.M.J.E. Vandenbroucke-Grauls, M. Kist, S. Bereswill and J.G. Kusters, 2001. Nickel-responsive induction of urease expression in *Helicobacter pylori* is mediated at the transcriptional level. *Infect. Immun.*, 69: 4891-4897.
- Mobley, H.L., M.J. Cortesia, L.E. Rosenthal and B.D. Jones, 1988. Characterization of urease from *Campylobacter pylori*. *J. Clin. Microbiol.*, 26: 831-836.
- Poore, C.A. and H.L.T. Mobley, 2003. Differential regulation of the *Proteus mirabilis* urease gene cluster by *UreR* and *H-NS*. *Microbiol.*, 149: 3383-3394.
- Fuller, M.E. and K.M. Scow, 1997. Impact of trichloroethylene and toluene on nitrogen cycling in soil. *Appl. Envir. Microbiol.*, 63: 4015-4019.
- Fletcher, D.W. and W.B. Bollen, 1954. The effects of aldrin on soil microorganisms and some of their activities related to soil fertility. *Appl. Envir. Microbiol.*, 2: 349-354.
- Pongsilp, N. and N. Nuntagij, 2007. Selection and characterization of mungbean root nodule bacteria based on their growth and symbiotic ability in alkaline conditions. *Suranaree J. Sci. Technol.*, 14: 277-286.
- Keele, Jr. B.B., P.B. Hamilton and G.H. Elkan, 1969. Glucose catabolism in *Rhizobium japonicum*. *J. Bacteriol.*, 97: 1184-1191.
- Christensen, W.B., 1946. Urea decomposition as a means of differentiating *Proteus* and *Paracolon* cultures from each other and from *Salmonella* and *Shigella* types. *J. Bacteriol.*, 52: 461-466.
- Askin, T. and R. Kizilkaya, 2006. Assessing spatial variability of soil enzyme activities in pasture topsoils using geostatistics. *Europ. J. Soil Biol.*, 42: 230-237.

18. Starnes, S.R., J.W. Spears, M.A. Froetschel, W.J. Croom, Jr., 1984. Influence of monensin and lasalocid on mineral metabolism and ruminal urease activity in steers. *J. Nutr.*, 114: 518-525.
19. Cappucino, J.C. and N. Sherman, 1992. *Microbiology: a Laboratory Manual*. Benjamin/ Cummings Publishing Company, pp: 125-179.
20. Liotenberg, S., D. Campbell, R. Rippka, J. Houmard and N. Tandeau De Marsac, 1996. Effect of the nitrogen source on phycobiliprotein synthesis and cell reserves in a chromatically adapting filamentous cyanobacterium. *Microbiol.*, 142: 611-622.
21. Moreau, M.C., R. Ducluzeau and P. Raibaud, 1976. Hydrolysis of urea in the gastrointestinal tract of "monoxenic" rats: effect of immunization with strains of ureolytic bacteria. *Infect. Immun.*, 13: 9-15.
22. Marcus, E.A. and D.R. Scott, 2001. Cell lysis is responsible for the appearance of extracellular urease in *Helicobacter pylori*. *Helicobacter*, 6: 93-99.
23. Geweely, N.S.I., 2006. Purification and characterization of intracellular urease enzyme isolated from *Rhizopus oryzae*. *Biotechnol.*, 5(3): 358-364.
24. Levican, G., J.A. Ugalde, N. Ehrenfeld, A. Mass and P. Parada, 2008. Comparative genomic analysis of carbon and nitrogen assimilation mechanisms in three indigenous bioleaching bacteria: predictions and validations. *BMC Genomics.*, 9: 581.
25. Schauer, K., C. Muller, M. Carrière, A. Labigne, C. Cavazza and H. De Reuse, 2010. The *Helicobacter pylori* GroES cochaperonin HspA functions as a specialized nickel chaperone and sequestration protein through its unique C-terminal extension. *J. Bacteriol.*, 192: 1231-1237.
26. Perrin, C., R. Briandet, G. Jubelin, P. Lejeune, M. Mandrand-Berthelot, A. Rodrigue and C. Dorel, 2009. Nickel promotes biofilm formation by *Escherichia coli* K-12 strains that produce curli. *Appl. Envir. Microbiol.*, 75: 1723-1733.
27. Ernst, F.D., J. Stoof, W.M. Horrevoets, E.J. Kuipers, J.G. Kusters and A.H.M. Van Vliet, 2006. NikR mediates nickel-responsive transcriptional repression of the *Helicobacter pylori* outer membrane proteins FecA3 (HP1400) and FrpB4 (HP1512). *Infect. Immun.*, 74: 6821-6828.
28. Premi, P.R. and A.H. Cornfield, 1969. Effects of addition of copper, manganese and chromium compounds on ammonification and nitrification during incubation of soil. *Plant Soil*, 31: 345-352.