RESEARCH PAPER

Optimization of urease production by *Bacillus halodurans* **PO15: a mangrove bacterium from Poovar mangroves, India**

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Abstract

Mangrove ecosystems are one of the most versatile habitats for microorganisms with a high potential for producing a variety of extracellular hydrolytic enzymes. In this study, bacteria with urease activity, enzymes that catalyze the hydrolysis of urea into carbon dioxide and ammonia, were isolated from mangrove sediments of Poovar (Trivandrum, India). *Bacillus halodurans*, strain PO15, isolated in this study with high urease (UA) activity (28 U/ml) was subjected to optimization using a Box-Behnken experimental design. Incubation variables included incubation period, pH, inoculation percentage and temperature. Signifcant factors identifed based on the model were incubation period, pH, incubation temperature, and inoculum percentage; variations in these produced a tenfold increase in UA activity (295.80 U/ml). The specifc activity of the purifed UA enzyme was 62.34 U/mg and was found to be thermostable (active up to 60 °C). UA of *B. halodurans* PO15 has potential for microbial-induced biomineralization with a reduction of free Ca^{2+} to about 82.8% \pm 0.17%. The microbialinduced calcium precipitation (MICP) using the UA enzyme will potentially be benefcial in the process of biomineralization as well as for a variety of industrial applications.

Keywords Urease · Biomineralization · Optimization · Mangrove · Enzymes

Introduction

Microorganisms play a major role in ecosystem engineering through several biogeochemical process (Graham et al. [2016](#page-7-0)). These processes are mostly driven by microbial enzymes. Microbial enzymes have been well studied for their industrial applications, and environment management. Urease, secreted by bacteria, is one such enzyme that has numerous applications. Urease is extensively used as a diagnostic tool in the detection of urea in blood (Smith et al. [1993](#page-8-0)), in alcoholic beverages to remove urea, biosensors for

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detection of heavy metal ions, and biocalcifcation (Sarda et al. [2009](#page-8-1)). Microbial ureases can induce calcite precipitation through reaction of urea and free calcium ions, a function that has applications in civil and geotechnical engineering for enhancing the strength and stifness properties of soil through the process of biomineralization (Anitha et al. [2018](#page-7-1); Bibi et al. [2018](#page-7-2); Cheng and Cord-Ruwisch [2013;](#page-7-3) Ivanov and Chu [2008\)](#page-7-4).

Urease (EC 3.5.1.15) is a nickel-containing enzyme which catalyzes the formation of carbon dioxide and ammonia from urea (Cheng and Cord-Ruwisch [2013\)](#page-7-3) resulting in an increase in pH in the surrounding media (Mora and Arioli [2014\)](#page-7-5). Urease-producing microorganisms are relevant to human microbiota, which hydrolyses urea (Chen and Burne [2003;](#page-7-6) Morou-Bermudez and Burne [2000;](#page-7-7) Wegmann et al. [2013;](#page-8-2) Yatsunenko et al. [2012\)](#page-8-3). The production of urease enzyme by ureolytic bacteria in soil may be infuenced by a variety of factors. It is induced in the presence of urea and inhibited in the presence of ammonia and nitrogen compounds (Mobley et al. [1995](#page-7-8)). Several studies have reported the isolation of indigenous ureolytic bacteria, using enrichment cultures from soil, ground water and cement samples (Achal and Pan [2011](#page-6-0); Burbank et al. [2012](#page-7-9); Elmanama et al. [2013](#page-7-10); Hammes et al. [2003;](#page-7-11) Rivadeneyra et al. [1993](#page-8-4)). Urease has been identifed as a virulence factor for several microbial pathogens (Mora and Arioli [2014\)](#page-7-5). Its role in microbial infections was well established through studies on *Helicobacter pylori* (Mora and Arioli [2014](#page-7-5)). The urease activity of microorganisms, such as *Proteus mirabilis* and *Staphylococcus saprophyticus,* plays a vital role in urinary tract infections through struvite–carbonate–apatite urinary stone formation. Based on the urease properties on formation of carbonate apatite stones, this enzyme has an application in environmental engineering. In addition to the above, urease enzyme also has many medical applications, including use as new drug targets.

Despite their application in the felds of biomedical and environmental engineering, the production conditions of the urease enzyme remain poorly understood. Previous studies have been limited to one-factor optimization (Bakhtiari et al. [2006](#page-7-12); El-Bessoumy et al. 2009) and have not concluded suitable production parameters for enhancing urease activity. For industrial applications, it is important to determine the optimal conditions for enzyme production. Traditionally, factorial methods were used in kinetic studies on microbially induced carbon precipitation (MICP) but this method is costly and time-consuming. Methods such as response surface methodology (RSM) in the form of the Box-Behnken experimental design were used to determine the optimum levels of key conditions as determined by Plackett–Burman design (PBD) (Box and Behnken [1960](#page-7-13); Plackett and Burnam [1946](#page-7-14)). This method could overcome the challenges of the conventional optimization techniques, which are laborious and result in unreliable and inaccurate results. These statistical techniques could help in designing

Fig. 1 Representation of urease production using *Bacillus halodurans* isolate PO15 based on Box-Behnken experimental design

experiments, building models, evaluating the interactive efects of variables, and determining optimum conditions (Shivam et al. [2009](#page-8-5)). RSM is widely used in bioprocessing technology for optimization of fermentation media (Desai [2008;](#page-7-15) Rishad et al. [2016;](#page-7-16) Sunitha et al. [1999\)](#page-8-6). In a study by Khodadadi and Bilsel [\(2015\)](#page-7-17), a central composite facecentered (CCF) design was used to ft a second-order model to evaluate microbial urease efficacy in the biocementation process. Optimum conditions of enzyme-specifc rate and urea hydrolysis were found to be signifcant. RSM or statistical modelling has been employed in the optimization of several enzymes, due to their reliability (Ameri et al. [2019](#page-7-18); Nathan et al. [2018](#page-7-19); Raza et al. [2019;](#page-7-20) Vijayaraghavan and Vincent [2014](#page-8-7)). This approach resulted in greater enzyme production of the microbial strains and are essential for any industrial applications. In this paper, we focus on a ureaseproducing bacterium, *Bacillus halodurans* PO15, isolated from mangrove sediment. We used statistical models for optimizing urease production at diferent incubation periods, pH, inoculation percentage and incubation temperatures, to achieve maximum enzyme activity. The biomineralization ability of the urease was also evaluated. This is the frst report of urease optimization through statistical models.

Results and discussion

Screening of urease‑producing bacteria

Fifty-two bacterial cultures were isolated from Poovar mangrove ecosystem sediment samples. Of these, 21 isolates gave urease positive results on urease agar. These

bacterial strains were inoculated into urease broth and urease production was quantified, based on a spectrophotometric assay. The urease activity of these isolates ranged from 1.8 to 28 U/ml. The bacterial isolate with the highest activity was selected for the further experiments. The strains with high urease activity were identified as *B. halodurans* through 16S rRNA ribotyping. The urease-producing bacteria is ubiquitous in natural environments. However, other common urease-producing strains reported here, *Helicobacter pylori*, *Proteus vulgaris*, *Staphylococcus aureus*, and *Pseudomonas aeruginosa*, etc., are pathogenic or opportunistic pathogens to humans (Stabnikov et al. [2013\)](#page-8-8). Additionally, many other bacterial strains that are used in microbial-induced calcium precipitation (MICP) with urease production *Bacillus* sp. VS1 and *Bacillus* sp. were reported (El-Bessoumy et al. 2009; Stabnikov et al. 2013). VUK5 has been extensively used in MICP studies (Stabnikov et al. [2013](#page-8-8)). In biocementation studies, spore-forming strains of ureaseproducing bacteria, were found to be more compatible to environments with high salt concentrations (Bachmeier et al. [2002](#page-7-21); Stabnikov et al. [2013\)](#page-8-8). For this reason, the optimization experiment in the present study was conducted for the *B. halodurans,* isolate PO15 alone. The major concern for environmental applications is the selection of an avirulent bacterial that has no adverse effect on humans or animals. Though many high urease-producing bacteria have been reported, they are mostly associated with human pathogenesis and cannot be used for any in situ environmental applications. In the present study, the mangrove bacterium, *B. halodurans* strain PO15*,* has no reported virulence and so could be employed in the optimization analysis.

Statistical optimization for urease production

In the present study, the most promising urease-producing isolate (*B. halodurans,* isolate PO15) was selected, based on urease activity. Figure [1](#page-1-0) presents a schematic representation of urease production in the *B. halodurans* isolate P015, based on Box-Behnken experimental design. Initial urease activity of *B. halodurans* isolate PO15 was about 28 U/ml. This value was quite higher than the activity reported for *Bacillus thuringiensis* N2, a marine bacterium (3.53 U/ml) (El-Bessoumy et al. [2009](#page-7-22)). To determine the efect of diferent factors on urease production, culture conditions were optimized using a Box-Behnken experimental design. A maximum urease activity of 295.80 U/ ml, a tenfold increase from initial activity, was achieved using the design (Table [1](#page-2-0)). Signifcant factors identifed,

Fig. 2 Contour plot showing response of variable infuencing urease production using *Bacillus halodurans* isolate PO15 based on Box-Behnken experimental design. The interaction between variables are shown: **a** incubation period vs. pH; **b** incubation period vs. temperature; **c** incubation period vs. inoculum (%); **d** pH vs. temperature; **e** pH vs. inoculum (%); **f** temperature vs. inoculum (%)

based on the model, were the incubation period, pH, incubation temperature, and inoculum percentage. Though aeration was reported as a major factor in enzyme production, for urease production, oxygen concentration has no role except in MICP (Bakhtiari et al. [2006\)](#page-7-12). The design predicted from the experiment was found to be signifcant $(R²$ value of 0.9961). Equation [1](#page-3-0) represents the quadratic model regression equation describing the predicted model. Khodadadi and Bilsel [\(2015](#page-7-17)) reported that the conditions favouring the urease production of *S. pasteurii*, the amount of urea hydrolyzed, and the rate of hydrolysis all inhibited bacterial cell growth and the specifc hydrolysis rate of urea and vice versa. However, in this study, a better urease production using *B. halodurans* PO15 was achieved. However, it is difficult to draw a comparison with other studies as most of them reported on the urea hydrolysis rate rather than urease production.

The ftted model is represented as Eq. [1](#page-3-0)

UA activity (U∕ml)

- $= 355.315 + 80.2708 \times$ Incubation Period
	- + $(-35.8631) \times pH + (-7.31012)$
	- \times Temperature + 7.14286 \times Inoculum %
	- $+ 0.125 \times$ Incubation period
	- \times pH + (- 1.7125) \times Incubation period
	- \times Temperature + (-4.75) \times Incubation period
- \times Inoculum $\%$ + 1.25 \times pH \times Temperature
- + $(-2.64286) \times pH \times Inoculum %$
- (1) $+ 0.921429 \times$ Temperature \times Inoculum %.

The interaction between the various medium components and factors for achieving maximum urease production are shown in contour plots (Fig. [2](#page-3-1)). From the interaction between the variables, it was found that the pH of the production medium is a critical factor for urease production.

When the medium pH vs. incubation period was tested, maximum urease activity of \sim 140 U/ml was achieved on the 6th day of incubation. For temperature vs. incubation period, urease activity reached a maximum after the 5th day only. Inoculum percentage vs incubation period achieved a maximum activity of 167 U/ml after 5 days. This suggests that a minimum of 5 days of incubation with a pH 5–7 is ideal for urease production, irrespective of all tested inoculum percentages and incubation temperatures. In another study on urease optimization of *A. niger* PTCC 5011, using conventional one-factor method, a maximum activity of 2.44 U/ml was obtained (Bakhtiari et al. [2006](#page-7-12)). Signifcant factors identifed, based on the model, were incubation period, pH, incubation temperature, and inoculum percentage. The production of the enzyme depended on process variables such as nutrients, pH, temperature, incubation period, inoculum level and inducer concentration (Sharma et al. [2009](#page-8-9)). Optimization of medium by the classical methods involved changing one independent variable (i.e., nutrient, pH, temperature) while keeping all other variables constant. However, one-factor optimization is extremely time-consuming, expensive for a large number of variables (Okyay and Rodrigues [2014](#page-7-23)) and often results in wrong conclusions. Use of statistical-based approaches, such as response surface methodology (RSM), could overcome the limitations of the single-factor optimization process. Also, the RSM-based approach requires fewer trials to calculate the diferent variables and their interactions, compared to other optimization methods (Managamuri et al. [2017](#page-7-24); Peng et al. [2018](#page-7-25)).

Except for the orthogonal array design-based approach for urease production using *A. niger* (Bakhtiari et al. [2006\)](#page-7-12) and response surface methodology (RSM) (Khan et al. [2019](#page-7-26))*,* there are no statistical optimization-based reports available for other bacteria. This is the frst report on the statistical optimization of urease production of *B. halodurans*. From the ANOVA results, the model F value of 6.65 and the *P* values < 0.005 indicate that the model was significant (Supplementary Table S1). The statistical model was validated through detection of 295 U/ml urease activity with optimized factors. Recently, El-Bessoumy et al. ([2009\)](#page-7-22) reported extracellular urease production from *B. thuringiensis* N2, however, the enzyme activity was very low.

Purifcation and characterization of UA

A specifc activity of 62.34 U/mg was observed for purifed urease with 5.6-purifcation fold and a yield of 87%. The specifc activity of the isolate PO15 was higher than those derived from *Aspergillus creatinolyticus* (32.74 U/ mg), *Lactobacillus reuteri* (13.0 U/mg) (Kakimoto et al. [1989\)](#page-7-27), *A. niger* (0.325 U/mg) (Smith et al. [1993](#page-8-0)), and *R. oryzae* (0.18 U/mg) (Geweely [2006](#page-7-28)). Based on the LB plot, a V_{max} of 333.33 mmol L⁻¹ mg⁻¹ min⁻¹ with K_{m} values of 1.7 mmol/L was observed for UA. The LB plot showing the enzyme kinetics is shown in Fig. [3a](#page-5-0). The UA enzyme was tested for its thermostability and was found to be stable up to 60 $\mathrm{^{\circ}C}$ (Fig. [3b](#page-5-0)). In another study, the fungal urease of *Aspergillus* exhibited maximum production and urease activity at 35 $\mathrm{^{\circ}C}$ and the least activity was at 50 $\mathrm{^{\circ}C}$ (Khan et al. [2019](#page-7-26)). Others reported 35 \degree C and 40 \degree C as the optimum temperature for urease activity (Danial et al. [2015;](#page-7-29) Fathima and Jayalakshmi [2012\)](#page-7-30). In this study, however, a maximum urease activity at 35 °C was observed, the enzyme had no signifcant decrease in urease activity even at 60 °C. This clearly validates the thermostable property of the urease enzyme.

Similarly, the optimal pH for maximum urease activity was found to be pH 7 (Fig. [3](#page-5-0)c). There was an increase in urease activity with an increase of pH from 3.0–9.0 (Khan et al. [2019\)](#page-7-26). Some bacterial ureases exhibited high activity in alkaline conditions (pH of 9.0) (Phang et al. [2018\)](#page-7-31), while some had maximum urease activity at pH 8 (Danial et al. [2015](#page-7-29); Mirbod et al. [2002](#page-7-32)). Two fungal isolates of the genus *Aspergillus* had an optimum pH of 8.0 and 8.5 (Kappaun et al. [2018](#page-7-33)). In general, the fungal urease had their maximum activity in the basic medium, while bacterial urease tended to be more variable (Khan et al. [2019\)](#page-7-26). The higher thermostability favours the application of UA in a variety of industrial and environmental engineering applications.

Urease‑mediated calcium precipitation

To understand the rate at which $CO₂$ is trapped as carbonates, a calcium carbonate precipitation study was carried out. The relative reduction of free calcium in the media is shown in Fig. [3](#page-5-0)d. *B. halodurans* PO15 was able to achieve a reduction of $(82.8 \pm 0.17)\%$ free Ca²⁺. Maximum reduction was observed after 48 h incubation. For the bioremediation of $CO₂$, the microbial biomineralization ability is of great importance (Silva-Castro et al. [2015\)](#page-8-10). Application of urease derived from *Sporosarcina pasteurii* for processes of biomineralization and co-precipitation of $CaCO₃$ was reported by Whiffin et al. ([2007\)](#page-8-11) and Al-Thawadi ([2011](#page-7-34)). This process of urease-aided $CaCO₃$ mineralization has a great potential in environmental engineering applications as well as for remediation and cementation in in situ conditions (Krajewska [2018](#page-7-35)). Bibi et al. ([2018\)](#page-7-2) reported indigenous *Bacillus* bacteria with biomineralization capability that could enhance soil stabilization isolated from Qatari soil. A similar report observed that *B. licheniformis* was able to precipitate calcium carbonate by ureolysis (Helmi et al. [2016\)](#page-7-36). This precipitation process uses carbonate ions released during urea hydrolysis and a pH shift to highly alkaline condition. It was found that the ureolytic property of *Bacillus* sp. is high with respect to any other genus and that this might be due to their physiological ability to adapt to

Fig. 3 Characterization of UA enzyme from *B. halodurans* isolate PO15 and evaluation of its biomineralization ability. (**a**) LB plot showing enzyme kinetics; (**b**) enzyme activity at diferent temperature; (**c**) enzyme activity at diferent pH; (**d**) Relative reduction of free $Ca²⁺$ in the media using urease enzyme produced by the *Bacillus halodurans* isolate PO15 [values expressed as mean \pm S.D of triplicate experiment]

stressed conditions (Helmi et al. [2016](#page-7-36)). Moreover, this also facilitates bioremediation of toxic metals and radionuclides through solid-phase capture (Fujita et al. [2000](#page-7-37)).

Conclusion

This is the frst report on the statistical optimization of extracellular urease production using a mangrove bacterium. A maximum urease activity of 295 U/ml was achieved by *B. halodurans* PO15 strain during statistical optimization. There was a tenfold increase in enzyme activity and the purifed enzyme exhibited a high specifc activity of 62.34 U/mg. The thermostable urease was active up to 60 °C and exhibited maximum activity at pH 7. UA of *B. halodurans* PO15 has potential for microbial-induced biomineralization with a reduction of free Ca²⁺ to about $(82.8 \pm 0.17)\%$. The microbial-induced calcium precipitation (MICP) using the UA enzyme could be useful in many environmental engineering applications and this opens up new avenues for simultaneous carbon mitigation and biomineralization.

Materials and methods

Isolation and screening of urease‑producing bacteria

Five sediment samples were collected from diferent locations of the Poovar mangrove system (N. Lat. 8°18′32′′ to 8°18′6′′ and E. Long. 77°4′32′′ to 77°5′14′′), located in southern Kerala, India. The samples were collected using sterile cylindrical PVC cores with a diameter of 10 cm. The samples were stored in iceboxes and transported to the laboratory. Upon reaching the laboratory, the samples were serially diluted up to 10^{-6} in physiological saline (0.85% NaCl) and plated onto nutrient agar (NA) (Hi Media, Mumbai, India) supplemented with 5% NaCl (Sigma Aldrich, USA) for the isolation of distinct bacterial colonies. Isolated bacterial colonies were subjected to urease enzyme screening on Urea agar base agar plates. The colour change of media from orange–yellow to deep pink indicated urease production.

Bacterial culture

A positive bacterial strain with high urease enzyme activity was isolated. The strain was identifed as *B. halodurans,* strain PO15, based on the 16S rRNA ribotyping (Refer Nathan et al. [2018](#page-7-19) for details of isolation and characterization). In this study, *B. halodurans,* strain PO15 was used for statistical optimization. The bacterium was grown in nutrient broth with 5% NaCl and incubated at 37 °C for 24 h. The culture was centrifuged at 10,000 r/min at 4 °C for 10 min to obtain crude urease enzyme.

Urease activity

The urease activity was determined by spectrophotometric assay based on the Nesslerization reaction. Briefy, 1.7 ml 10 mmol/L urea was mixed with 0.2 ml of 0.05 mol/L Tris—HCl (pH 7.0) and 20 µl of the urease. The mixture was incubated at 37 ˚C for 10 min and the reaction was stopped by adding 1.5 mol/L Trichloro acetic acid (TCA) (Sigma, USA). The reaction mixture was again incubated after adding 0.5 ml of Nessler's reagent at 37˚C for 10 min and absorbance was read at 405 nm on double beam UV–Vis spectrophotometer (SYSTRONIC MAKE, MODEL 101). One unit of urease was defned as the amount of enzyme required to release one micromole of ammonia as determined from an ammonium chloride standard curve (Kayastha et al. [1995](#page-7-38)).

Statistical optimization and model validation

For enhancing the urease activity, the optimization of culture conditions was carried out based on the Box-Behnken design, using the Design Expert 9.0 software. The response surface methodology (RSM) helped develop the mathematical models for understanding the enzyme activity on independent variables (Cui and Zhao [2012](#page-7-39)). The factors that varied during optimization were incubation period, pH, inoculation percentage and incubation temperature. Each factor was studied at two different levels (-1) low and $+1$ high) (Box and Behnken [1960\)](#page-7-13). Forty-three experiments were carried out in triplicate in 250 ml Erlenmeyer fasks. Bacterial inocula were prepared with 0.5 OD McFarland Standards and added according to the inoculum percentage (3–10%). The response experimental values were derived from the mean \pm S.D of three independent experiments. The Box-Behnken design (BBD) is based on a second-order polynomial equation (Eq. [2](#page-6-1)). The statistical model obtained from the experiments was validated using the optimized fermentation conditions. The enzyme assay was performed and urease activity was calculated.

$$
Y = \beta_0 + \beta_1 A + \beta_2 B + \beta_3 C + \beta_1 \beta_1 A^2 + \beta_2 \beta_2 B^2
$$

+ $\beta_3 \beta_3 C^2 + \beta_1 \beta_2 AB + \beta_1 \beta_3 AC + \beta_2 \beta_3 BC,$ (2)

where Y represents the response urease activity in U/ml; A, B, and C-coded independent variables; β_1 , β_2 , and β_3 -linear coefficients; β_0 —intercept term; $\beta_1\beta_1$, $\beta_2\beta_2$, and $\beta_3\beta_3$ –quadratic coefficients; $\beta_1\beta_2$, $\beta_1\beta_3$, and $\beta_2\beta_3$ –interactive coefficients.

Purifcation and characterization of urease

For the purifcation process, crude enzyme was obtained after centrifugation of the bacterial culture at 10,000 r/min at 4 °C. 80% ammonium sulphate was added to the enzyme and the pellet was dialyzed against 250 mmol/L Tris HCl bufer, pH 8.3 at 4° C for 48 h. The lysate was purified using affinity column chromatography (Sepharose®4B-L-tyrosine-p-aminobenzene sulfonamide). The purifed enzyme fraction was dried in a vacuum desiccator, resuspended in 10 mmol/L phosphate bufer (pH 7.2), which was stored at 4° C for further experiments. The specifc activity and yield (%) of urease were calculated. The enzyme–substrate interaction was further studied using the Line weaver Burk (LB) plot. K_m and V_{max} values were derived from the LB plot. The optimum pH and temperature for the maximum urease activity was also evaluated.

Calcite precipitation experiment

The ability of the bacterial isolate to sequester atmospheric $CO₂$ was demonstrated using the calcium precipitation assay. For this, nutrient broth was fortified with $NaHCO₃$ (25.2 mmol/L) and CaCl₂ (25.2 mmol/L). The relation of UA with free Ca²⁺ reduction was evaluated over a time period of 48 h. The bacterial isolate was inoculated to the medium and incubated under static conditions at (35 ± 2) °C. The samples were retrieved at 12 h interval, were subjected to free Ca^{2+} analysis using an atomic absorption spectrophotometer (AAS) (Elico, India) after centrifuging the broth at 10,000 r/min for 5 min to obtain the supernatant. Free calcium reduction (%) was expressed as mean \pm S.D from the triplicate experiments performed.

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Author contributions VKN is involved in work design, execution and manuscript preparation. JV and PA involved in manuscript preparation.

Compliance with ethical standards

Conflict of interest The authors declare that they have no confict of interest.

Animal and human rights statement This article does not contain any studies with human participants or animals performed by any of the authors.

References

Achal V, Pan X (2011) Characterization of urease and carbonic anhydrase producing bacteria and their role in calcite precipitation. Curr Microbiol 62:894–902

- Al-Thawadi SM (2011) Ureolytic bacteria and calcium carbonate formation as a mechanism of strength enhancement of sand. J Adv Sci Eng Res 1:98–114
- Ameri A, Shakibaie M, Soleimani-Kermani M, Faramarzi MA, Doostmohammadi M, Forootanfar H (2019) Overproduction of thermoalkalophilic lipase secreted by *Bacillus atrophaeus* FSHM2 using UV-induced mutagenesis and statistical optimization of medium components. Prep Biochem Biotechnol 49:184–191
- Anitha V, Abinaya K, Prakash S, Seshagiri Rao A, Vanavil B (2018) *Bacillus cereus* KLUVAA mediated biocement production using hard water and urea. Chem Biochem Eng Q 32:257–266
- Bachmeier KL, Williams AE, Warmington JR, Bang SS (2002) Urease activity in microbiologically-induced calcite precipitation. J Biotechnol 93:171–181
- Bakhtiari MR, Faezi MG, Fallahpour M, Noohi A, Moazami N, Amidi Z (2006) Medium optimization by orthogonal array designs for urease production by *Aspergillus niger* PTCC5011. Process Biochem 41:547–551
- Bibi S, Oualha M, Ashfaq MY, Suleiman MT, Zouari N (2018) Isolation, diferentiation and biodiversity of ureolytic bacteria of Qatari soil and their potential in microbially induced calcite precipitation (MICP) for soil stabilization. RSC Adv 8:5854–5863
- Box GE, Behnken DW (1960) Some new three level designs for the study of quantitative variables. Technometrics 2:455–475
- Burbank MB, Weaver TJ, Williams BC, Crawford RL (2012) Urease activity of ureolytic bacteria isolated from six soils in which calcite was precipitated by indigenous bacteria. Geomicrobiol J 29:389–395
- Chen YM, Burne RA (2003) Identifcation and characterization of the nickel transport system for urease biogenesis in *Streptococcus salivaius* 57.I. J Bacteriol 185:6773–6779
- Cheng L, Cord-Ruwisch R (2013) Selective enrichment and production of highly urease active bacteria by non-sterile (open) chemostat culture. J Ind Microbiol Biotechnol 40:1095–1104
- Cui F, Zhao L (2012) Optimization of xylanase production from *Penicillium* sp.WX-Z1 by a two-step statistical strategy: Plackett-Burman and Box-Behnken experimental design. Int J Mol Sci 13:10630–10646
- Danial EN, Hamza AH, Mahmoud RH (2015) Characteristics of immobilized urease on grafted alginate bead systems. Braz Arch Biol Technol 58:147–153
- Desai KM (2008) Comparison of artifcial neural network (ANN) and response surface methodology (RSM) in fermentation media optimization: case study of fermentative production of scleroglucan. Biochem Eng J 41:266–273
- El-Bessoumy A, El-Sharouny EB, Olam Z, Mothana A (2009) Purifcation and characterization of marine *Bacillus thuringiensis* N2 urease. Egyptian J Biochem Mole Biol 27:61–78
- Elmanama AA, Alhour MT (2013) Isolation, characterization and application of calcite producing bacteria from urea rich soils. J Adv Sci Eng Res 3:377–399
- Fathima F, Jayalakshmi S (2012) Characterization of urease enzyme from marine bacterium *Klebsiella* species. Afr J Microbiolo Res 6:5914–5923
- Fujita Y, Ferris FG, Lawson RD, Colwell FS, Smith RW (2000) Subscribed content calcium carbonate precipitation by ureolytic subsurface bacteria. Geomicrobiol J 17:305–318
- Geweely NS (2006) Purifcation and characterization of intracellular urease enzyme isolated from *Rhizopus oryzae*. Biotechnol 5:358–364
- Graham EB, Knelman JE, Schindlbacher A, Siciliano S, Breulmann M, Yannarell A, Beman JM, Abell G, Philippot L, Prosser J, Foulquier A (2016) Microbes as engines of ecosystem function: when does community structure enhance predictions of ecosystem processes? Front Microbiol 7:214
- Hammes F, Boon N, de Villiers J, Verstraete W, Siciliano SD (2003) Strain-specifc ureolytic microbial calcium carbonate precipitation. Appl Environ Microbiol 69:4901–4909
- Helmi FM, Elmitwalli HR, Elnagdy SM, El-Hagrassy AF (2016) Calcium carbonate precipitation induced by ureolytic bacteria *Bacillus licheniformis*. Ecol Eng 90:367–371
- Ivanov V, Chu J (2008) Applications of microorganisms to geotechnical engineering for bioclogging and biocementation of soil in situ. Rev Environ Sci Biotechnol 7:139–153
- Kakimoto S, Sumino Y, Akiyama S, Nakao Y (1989) Purifcation and characterization of acid urease from *Lactobacillus reuteri*. Agric Biol Chem 53:1119–2112
- Kappaun K, Piovesan AR, Carlini CR, Ligabue-Braun R (2018) Ureases: historical aspects, catalytic, and non-catalytic properties a review. J Adv Res 13:3–17
- Kayastha AM, Das N, Malhotra OP (1995) Urease from the seeds of pigeonpea (*Cajanus cajan* L.). In: Biopolymers and bioproducts: structure, function and applications. Dokya Pub, Bangkok, pp 382–386
- Khan YM, Munir H, Anwar Z (2019) Optimization of process variables for enhanced production of urease by indigenous *Aspergillus niger* strains through response surface methodology. Biocatal Agric Biotechnol 11:101–202
- Khodadadi HT, Bilsel H (2015) Statistical modeling of environmental factors on microbial urea hydrolysis process for biocement production. Adv Mat Sci Eng 2015:340930
- Krajewska B (2018) Urease-aided calcium carbonate mineralization for engineering applications: a review. J Adv Res 13:59–67
- Managamuri U, Vijayalakshmi M, Ganduri VRK, Rajulapati SB, Bonigala B, Kalyani BS, Poda S (2017) Isolation, identifcation, optimization, and metabolite profling of *Streptomyces sparsus* VSM-30. 3 Biotech 7:217
- Mirbod F, Schaller RA, Cole GT (2002) Purifcation and characterization of urease isolated from the pathogenic fungus *Coccidioides immitis*. Med Mycol 40:35–44
- Mobley HL, Island MD, Hausinger RP (1995) Molecular biology of microbial ureases. Microbiol Mol Biol Rev 59:451–480
- Mora D, Arioli S (2014) Microbial urease in health and disease. PLoS Pathog 10:e1004472
- Morou-Bermudez E, Burne RA (2000) Genetic and physiologic characterization of urease of *Actinomyces naeslundii*. Infect Immun 67:504–512
- Nathan VK, Kanthimathinathan SR, Rani ME, Rathinasamy G, Kannan ND (2018) Biobleaching of waste paper using lignolytic enzyme from *Fusarium equiseti* VKF2: a mangrove isolate. Cellulose 25:4179–4192
- Okyay TO, Rodrigues DF (2014) Optimized carbonate micro-particle production by *Sporosarcina pasteurii* using response surface methodology. Ecol Eng 62:168–174
- Peng H, Tan J, Bilal M, Wang W, Hu H, Zhang X (2018) Enhanced biosynthesis of phenazine-1-carboxamide by *Pseudomonas chlororaphis* strains using statistical experimental designs. World J Microbiol Biotechnol 34:129
- Phang IRK, San Chan Y, Wong KS, Lau SY (2018) Isolation and characterization of urease-producing bacteria from tropical peat. Biocatal Agric Biotechnol 13:168–175
- Plackett RL, Burman JP (1946) The design of optimum multifactorial experiments. Biometrika 33:305–325
- Raza A, Bashir S, Tabassum R (2019) Statistical based experimental optimization for co-production of endo-glucanase and xylanase from *Bacillus sonorensis* BD92 with their application in biomass saccharifcation. Folia Microbiol 64:295–305
- Rishad KS, Rebello S, Nathan VK, Shabanamol S, Jisha MS (2016) Optimised production of chitinase from a novel mangrove isolate, *Bacillus pumilus* MCB-7 using response surface methodology. Biocatal Agric Biotechnol 5:143–149
- Rivadeneyra MA, Delgado R, Delgado G, Moral AD, Ferrer MR, Ramos-Cormenzana A (1993) Precipitation of carbonates by *Bacillus* sp. isolated from saline soils. Geomicrobiol J 11:175–184
- Sarda D, Choonia HS, Sarode DD, Lele SS (2009) Biocalcifcation by *Bacillus pasteurii* urease: a novel application. J Indust Microbiol Biotechnol 36:1111–1115
- Sharma A, Bardhan D, Patel R (2009) Optimization of physical parameters for lipase production from *Arthrobacter* sp. BGCC# 490. Indian J Biochem Biophys 46:178–183
- Shivam K, Chandra P, Tripathi M, Mishra SK (2009) Culture conditions for the production of α-galactosidase by *Aspergillus parasiticus* MTCC-2796: a novel source. Electron J Biotechnol 12:1–9
- Silva-Castro GA, Uad I, Gonzalez-Martinez A, Rivadeneyra A, Gonzalez-Lopez J, Rivadeneyra MA (2015) Bioprecipitation of calcium carbonate crystals by bacteria isolated from saline environments grown in culture media amended with seawater and real brine. BioMed Res Int 2015:816102
- Smith PT, King AD Jr, Goodman N (1993) Isolation and characterization of urease from *Aspergillus niger*. Microbiol 139:957–962
- Stabnikov V, Jian C, Ivanov V, Li Y (2013) Halotolerant, alkaliphilic urease-producing bacteria from diferent climate zones and their application for biocementation of sand. World J Microbiol Biotechnol 29:1453–1460
- Sunitha K, Lee JK, Oh TK (1999) Optimization of medium components for phytase production by *E. coli* using response surface methodology. Bioprocess Eng 21:477–481
- Vijayaraghavan P, Vincent SGP (2014) Statistical optimization of fbrinolytic enzyme production by *Pseudoalteromonas* sp. IND11 using cow dung substrate by response surface methodology. Springerplus 3:60
- Wegmann U, Louis P, Goesmann A, Henrissat B, Duncan SH, Flint HJ (2013) Complete genome of a new *Firmicutes* species belonging to the dominant human colonic microbiota ('*Ruminococcus bicirculans*') reveals two chromosomes and a selective capacity to utilize plant glucans. Environ Microbiol 16:2879–2890
- Whifn VS, van Paassen LA, Harkes MP (2007) Microbial carbonate precipitation as a soil improvement technique. Geomicrobiol J 24:417–423
- Yatsunenko T, Rey FE, Manary M, Trehan I, Dominguez-Bello MG, Contreras M, Magris M, Hidalgo G, Baldassano RN, Anokhin AP, Heath AC, Warner B, Reeder J, Kuczynski J, Caporaso JG, Lozupone CA, Lauber C, Clemente JC, Knights D, Knight R et al (2012) Human gut microbiome viewed across age and geography. Nature 486:222–227