

Full Length Research Article

Advancements in Life Sciences — International Quarterly Journal of Biological Sciences

ARTICLE INFO

Open Access



Date Received: 04/01/2025; Date Revised: 12/07/2025; Date Published Online: 28/12/2025;

Production and purification of Nattokinase from *Pseudomonas* aeruginosa P49

Shahad Abdullah Shwan*, Ali Jabbar Reshak

Authors' Affiliation:

Biotechnology Department, College of Science, Baghdad University - Iraq

Abstract

*Corresponding Author: Shahad Abdullah Shwan Email: shahadabdullahhhhhhh@gmail.com **Background:** Nattokinase (NK) is a profibrinolytic serine protease enzyme produced by many bacterial strains, such as *Pseudomonas aeruginosa*. Therefore, this study aimed to produce NK from local isolate of *P. aeruginosa* P49 and optimize its conditions for the production of enzyme.

How to Cite:

Shwan SA, Reshak AJ (2025). Production and purification of Nattokinase from *Pseudomonas aeruginosa* P49. Adv. Life. Sci. 12(4): 827**Methods:** 150 samples were obtained from clinical sources, during the period from August to November 2022, from different hospitals. All samples were subjected to different examinations, in addition to VITEK2 system, to confirm that these isolates were *P. aeruginosa*.

Keywords:

Nattokinase; *Pseudomonas aeruginosa*; Enzyme Purification; IEC; SF; VITEK2 system; Ion exchange Chromatography **Result:** A total of one hundred *P. aeruginosa* isolates were screened to choose the best NK-producing isolates using skim milk agar, then broth media, whereas *P. aeruginosa* P49 gave the highest enzymatic activity (337.9 U/mg protein). Optimal conditions for the formation of NK were estimated, and the results showed that the maximum production of NK was gained using peptone-yeast medium containing sucrose, peptone, and CaSO4.2H2O at pH 7.5 and 37°C for 24 hours of incubation, whereas the activity of NK increased to reach a yield of 1603 U/mg protein. The NK was purified from *P. aeruginosa* P49 utilizing ion exchange chromatography (IEC) after precipitation by ammonium sulfate (0-75%). The results for enzyme purification gave 96% of NK enzyme with a purification fold of 3.6, and the specific activity was 2190.7 U/mg protein.

Conclusion: This result suggests *P. aeruginosa* P49 is a good source of NK production.

Introduction

Nattokinase is a serine protease with potent profibrinolytic activity and can degrade fibrin very effectively. Nattokinase has been manufactured using various host strains and possesses the benefits of increased stability, low cost, and minimal adverse effects. Unlike the other fibrinolytic enzymes, including streptokinase, tissue plasminogen activator (t-PA), and urokinase, NK demonstrates potential as a therapeutic drug against cardiovascular diseases as well as a healthy food supplement [1].

Recently, Nattokinase (NK) has drawn increasing attention owing to its wide range of potential applications [2]. By lowering plasma fibrinogen and blood viscosity levels, NK aids in the prevention and management of thrombotic diseases. Research points out its benefits in the areas of food safety, low cost, and extended biological activity [3,4]. Unlike other fibrinolytic drugs, this enzyme possesses special advantages such as suitability for oral administration, extended preventive effects, and tolerance in the digestive system. Also, NK is absorbed via the gastrointestinal system and then facilitates the lysis of fibrin later on [5]. In addition to its thrombolytic effect, NK shows various other effects such as the prevention of atherosclerosis, enhanced microcirculation, blood pressure regulation, inhibition of abnormal retinal vasculature proliferation, anticancer activity, and ability to counteract inflammation and oxidative stress.

A significant amount of NK is also found to be produced by many marine organisms besides Bacillus and Pseudomonas species [1]. Pseudomonas aeruginosa are abundant organisms found in both human-made and natural environments, including those involving plants and animals [7]. Pseudomonads opportunistic pathogenic bacterial species that can cause infectious diseases in the eyes, skin, and lungs of people with burns, abrasions, cystic fibrosis (CF), and HIV/AIDS [8]. Although some species of P. aeruginosa are pathogenic, these bacteria also produce different beneficial enzymes [9], like NK [1].

In general, NK is formed via submerged fermentation (SF) and solid-state fermentation (SSF) processes, but the latter has been utilized to a lower extent for production of these enzymes [10]. The culture of microorganisms in a liquid nutrient broth that is high in oxygen concentration is referred to as SF [11].

Despite the fact that the microbial synthesis of NK has been the subject of much research, the yields, activities, and stabilities of NK remain the most important factors limiting its applicability in industrial settings [12]. Many parameters affecting enzyme production from P. aeruginosa have been investigated, including the best substrate, the pH of the medium, the

incubation time, salts, and the incubation temperature. All enzymes are grown in culture media that contain carbon sources, nitrogen sources, and other components that assist the growth of the enzymes. Many studies show that P. aeruginosa grows in response to carbon sources, with each carbon source having a unique impact on NK synthesis [13]. Also, NK is a neutral or weak alkaline enzyme since it is stable within the pH ranges of 6.0 and 12.0 and is quickly inactivated below pH 5 [12]. The optimum temperature for microbial growth is not necessarily the same as the optimum temperature for enzyme production, even between strains of the same species, due to the fact that temperature affects the cell's metabolic activities. Incubation temperature has a major effect on the metabolic activities of bacteria [14].

Hence, this study aimed to produce NK from local isolate of P. aeruginosa, optimize its production conditions, and purify it.

Methods

Collection of samples

A total of 150 samples were obtained from various clinical sources, including wounds and burns, during the period from August 2022 and November 2022, from patients attained Al-Kindi Hospital, Al-Yarmouk Hospital, Teaching Hospital of Baghdad Medical city, and other specific laboratories.

Identification of samples

All samples were subjected to different examinations, including microscopic examinations (gram stain) [15], cultural features (cetrimide agars and MacConkey agars, nutrient agars, brain heart agar, and blood agar base) [16], biochemical tests (Simmons citrate, catalase, and oxidase) [17] and growth at 4°C and 42°C, in order to isolate P. aeruginosa as well as VITEK 2 system was utilized [18] to confirm that these isolates were P. aeruginosa.

Screening the *Pseudomonas aeruginosa* isolate for NK production

Semi-quantitative screening (primary screening)

All Pseudomonas aeruginosa isolates were screened to choose the best isolates for NK production utilizing a plate assay via skim milk agar medium. A volume of one hundred fifty microliters of inoculum of bacteria consisting of 2.5×10⁷ cells/milliliter was transferred into a skim milk agar plate and then placed in an incubator at 37°C for 24 hrs. The presence of NK secretion might be seen as a colorless zone of hydrolysis around the well [19].

Quantitative screening (secondary screening) for enzyme production using SF medium

Five isolates of NK-produced Pseudomonas aeruginosa were screened to choose the best isolates for NK production. SF medium, composed of 2.5 g yeast extract, 5.0 g of glucose, 2.5 g of peptone, 0.5 mg KH₂PO₄, 0.1 mg MgSO₄, and 2.5 g NaCO₃. This medium was poured into a moistened Erlenmeyer flask measuring 250 milliliters. The flask was tightened with cotton after adjusting the pH to neutral, and contents of the flask were mixed thoroughly. Then the flask was placed in an autoclave at 121°C temperature and 15 psi pressure for 15 minutes. After sterilization, the flasks were cooled at room temperature [20]. SF was used for enzyme production, where the five isolates with maximal productivities according to casein hydrolysis were chosen and cultured on fermentation medium. Five bacterial suspensions (2.5×10⁷ cells/ml) were prepared from 24-hour cultures, each used to inoculate the fermentation broth with 0.75 microliter aliquots. The flasks were put in an incubator shaker for 24 hours at 37°C.

Extraction of NK enzyme

After an incubation period of twenty-four hours, enzyme from each flask was extracted by centrifuging the flasks for thirty minutes at 10,000 rpm. The supernatant was then filtered using filter papers of Whatman No. 1. It was determined that the supernatant may be used as a crude enzyme.

Determination of NK activities

The activity of NK enzyme was determined according to Senior (1999) method as shown below: 1.8 milliliter of casein solution was placed in a bath of water for 5 min at 37°C, then 0.2 milliliter of the crude enzyme was firstly put into the solution of substrates and then in an incubator for 30 min at 37°C [21]. After adding three millilitres of a TCA solution with a concentration of ten percent, the reaction was stopped, and the pellet was separated by centrifuging the mixture at ten thousand revolutions per minute for ten minutes. A blank was prepared by following the identical protocols as the test sample, with the exception that a 10% TCA solution was mixed with the solutions of caseins before the addition of 0.2 millilitres of enzyme. A UV-VIS spectrophotometer was used in order to determine the absorbances of the supernatants at 280 nm. The enzyme activities and concentrations of proteins were each measured twice, and the findings were reported based on the average of the two readings. According to the formula that follows, a unit of activity of the enzyme was the amount needed to raise the absorbance ratio at 280 nm by 0.01 per minute in the experimental conditions:

Enzyme activity
$$\left(\frac{\text{U}}{\text{milliliter}}\right) = \frac{\text{Absorbance at 280 nm}}{0.01 \times 30 \times 0.2}$$

Where, 0.01 = Constant, 30 = Reactive time in minutes, and 0.2 = Volume of enzyme

Determinations of concentrations of proteins

Method of Bradford (1976) was utilized for measuring concentration of proteins, in accordance with Al-Sa'ady (2020) [22,23].

Determinations of specific activities

As per the following equation, the specific activity of the enzyme was calculated [24]:

Specific activity
$$\left(\frac{U}{mg} \text{protein}\right) = \frac{\text{Enzyme activity } \left(\frac{U}{\text{milliliter}}\right)}{\text{Protein concentration } \left(\frac{mg}{\text{milliliter}}\right)}$$

Optimal conditions for the production of NK Optimal culture medium

Six different media were examined to see which ones had the most impact on NK production. These substrates included:

- Peptone 1 g/100 ml, yeast extract (0.25 g/100 ml), NaCl (1 g/100 ml).
- Peptone (1 g/100 ml), yeast extract (0.25 g/100 ml), Glucose (1 g/100 ml).
- Glucose (2 g/100 ml), soy bean (2 g/100 ml), NaCl 0.5 g/100 ml, KH₂PO₄ (0.1 g/100 ml), K₂HPO₄ (0.4 g/100 ml), MgSO₄ (0.05 g/100 ml).
- Shrimp shell 1 g/100 ml, KH₂PO₄ 0.1 g/100 ml, MgSO₄ 0.05 g/100 ml
- Nutrient broth (1 g/100 ml), shrimp shell (1 g/100
- Glucose (0.5 g/50 ml), peptone (0.25 g/50 ml), yeast extract (0.25 g/50 ml), KH₂PO₄ (0.05 g/50 ml), MgSO₄ (0.01 g/50 ml), and NaCO₃ (0.25 g/50 ml).

In each 250 milliliter flask, 100 milliliters of distilled water was added to each agricultural waste and sterilized using an autoclave for 15 min at 121°C. The chosen bacterial inoculum of 1 ml and having a bacterial concentration of 2.5×10⁷ cells/milliliter, was used to inoculate each flask. These flasks were then incubated at 37°C for 24 hours, and the enzyme activity and protein concentration were estimated.

Effect of carbon source

To determine the best carbon source for NK production from P. aeruginosa P49 in selected medium, different carbon sources were used including glucose, sucrose, maltose, fructose, starch, lactose and cellulose, whereas 0.5% of each source was added into 100 milliliter of selected medium and the medium flask was sterilized using autoclave at 121°C for 15 min. Then, each flask was inoculated with 1.0 milliliter of selected bacterial inoculum (2.5×10⁷cell/milliliter) and placed in an incubator for 24 hours at 37°C [25]. Specific activity of enzyme was also estimated.

Effect of nitrogen source

To determine the best nitrogen source for NK production from *P. aeruginosa* P49 in selected medium, different nitrogen sources were employed, including peptone, yeast extract, and meat extract. For this purpose, the following procedure was carried out: firstly, each nitrogen source was used separately at concentration 20 g/L then half amount of two-nitrogen sources together and finally equal amount of three different nitrogen sources were employed, as follows: 20 g/L of peptone, 20 g/L of yeast extract, 20 g/L of meat extract, mixture of 10 g/L of peptone and 10 g/L of yeast extract, mixture of 10 g/L of peptone and 10 g/L of meat extract, mixture of 10 g/L of meat extract and 10 g/L of yeast extract and mixture of 6.6 g/L of peptone, 6.6 g/L of yeast extract and 6.6 g/L of meat extract. All flasks were placed in an autoclave and inoculated with 1.0 milliliter of chosen inoculum of bacterial species (2.5×107cell/milliliter) and put in an incubator for 24 hrs at 37°C. Specific activity of enzyme was estimated.

Effect of salts

To determine the salt effect on NK production from P. aeruginosa P49 in the selected medium, different mineral salt solutions (MS) were used, including FeSO4.7H₂O, CuSO₄.5H₂O, MgSO₄.7H₂O, MnSO₄.7H₂O, CaSO₄.2H₂O, and ZnSO₄.7H₂O. For this purpose, the following procedure was carried out: firstly, each salt was used separately at a weight of 0.1 g/100 milliliter, and half amount of two salts was utilized together. Besides the estimation of enzyme's specific activity, all flasks were inoculated as per previously described method and incubated for 24 hours at 37°C [26].

Initial pH

The pH of the chosen medium was varied throughout a wide range, from 3 to 9, including 3, 3.5, 4, 4.5, 5, 5.5, 6, 6.5, 7, 7.5, 8, 8.5, and 9. Similar steps of bacterial inoculation of the flasks and incubation, along with the estimation of specific enzyme activity, were repeated as mentioned in the previous section after autoclaving [26].

Optimum temperature

To find the ideal temperature for the production of NK, the selected medium was subjected to different temperatures of 25, 30, 35, 37, and 40°C. Each flask was inoculated after autoclaving as previously explained with bacterial inoculum, incubated, and specific enzyme activity was calculated [26].

Optimum incubation period

The determination of optimum incubation duration for the production of NK was estimated by inoculating each medium as previously described, incubating along with the estimation of specific enzyme activity, but this time the incubator was operated at different periods of time, that is, 12, 24, 36, and 48 hours [26].

Purification of NK

Through the use of ammonium sulphate precipitation and ion exchange chromatography, NK was extracted from a local isolate of P. aeruginosa P49 and then purified.

Ammonium sulfate precipitation and ion exchange chromatography

At a temperature of 4°C, solid ammonium sulphate ranging from 0 to 75% was progressively added to the crude enzyme. The ingredients were then mixed carefully for 2 hours and 45 minutes. After that, the mixture was centrifuged at a speed of 10,000 revolutions per minute for twenty minutes, and the supernatant was thrown away. The precipitate was put in 10 millilitres of a buffer that contained 0.2 M potassium phosphate. It was possible to determine both the enzyme's activity and the concentration of proteins.

Based on the method suggested by Schütte (1985), Ion exchange chromatography (DEAE-Cellulose) was utilized for NK purification [27]. During the stage that included ammonium sulphate precipitation and by utilizing a clean Pasteur pipette that moved in circular motions on the wall of column, a total of 13 millilitres of the concentrated enzyme was placed onto a DEAE-Cellulose column that measured 23 by 1.7 centimeters. The column was cleaned with an equilibrating solution of Tris-HCl with a pH of 8, a concentration of 0.2 M, and a flow rate of 30 milliliters/hr, with 3 milliliters being used for each fraction till the optical density at 280 nm read zero. The elution was carried out at a flow rate of 30 milliliters/hr using the gradient concentrations of NaCl ranging from 0.0 M to 1.0 M that were formed in a Tris-HCl buffer.

The protein fractions of both the washed and eluted fractions were determined at a wavelength of 280 nm. The portions of the protein peaks that had enzymatic activity were collected, and the sections of the protein peaks that did not include enzymatic activity were tested for NK activity. It was possible to determine both the enzyme activity and the concentration of the protein.

Results

Isolations and Identifications of P. aeruginosa

All samples were subjected to various examinations, including microscopic examinations, cultural characteristics, and biochemical, in order to isolate P. aeruginosa. All the essential tests and examinations for initial diagnoses of the well-grown bacterial isolates were performed as displayed in Table 1.

Examination	Characteristics	P. aeruginosa				
Microscopic	Shape of cells	Rods				
	Gram staining	Negative				
Cultural Characteristics	Cetrimide agar	Growing with green, fluorescent light, large colonies, and a grape-like odor				
	MacConkey agar	Elevated colonies of light non-lactose Fermenters				
	Nutrient agar/ Brain hart agar Develop with a green, fluorescent light, larg colonies, and a grape-like odor					
	Hemolysis	Positive (+ve) - Beta				
Biochemical	Simmons citrate	Positive (+ve)				
	Catalase	Positive (+ve)				
	Oxidase	Positive (+ve)				
	Growth at 42℃	Positive (+ve)				

Table 1: Results of examinations for identification of *P. aeruginosa* isolates from various clinical specimens.

Screening Program for NK (NK) Production

The findings of primary screening displayed that only five isolates of *P. aeruginosa* from 150 isolates were the better isolates for NK production compared with the other isolates, as shown in Figure 1A.

Secondary Screening

The five *P. aeruginosa* isolates (P11, P23, P42, P49, and P104) with the highest hydrolysis zone in the main screening were screened once again for their enzymatic activities utilizing the technique of digesting caseins. This was done for the purpose of further detection and the careful selection of an effective bacterial isolate to generate NK. *P. aeruginosa* P49 had the greatest enzymatic activity out of the five isolates tested. Its NK-specific activity in crude supernatant was 337.9 U/mg protein, while for supplementary strains, it varied from 222.9 to 314.1 U/mg protein, as displayed in Figure 1B.

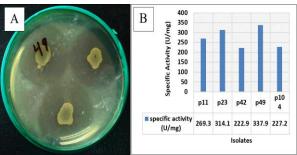


Figure 1: A) Proteolytic activity of *P. aeruginosa* P49 on skim milk agar; **B)** Quantitative screening of *P. aeruginosa* isolates for NK production utilizing SF medium.

Optimum Conditions for NK Production

The various bioprocess parameters that impact NK synthesis by *P. aeruginosa* P49 under SF were optimized for maximal production of enzymes, and the findings that were obtained were among the most fascinating ones. These results are explained in more detail below:

Optimum Culture Media

Six types of culture media were examined for their efficiency in NK production for *P. aeruginosa* P49. Of all media, the first culture medium of *P. aeruginosa* P49 showed the most enhanced NK production with a NK specific activity value of 521.12 U/mg protein,

whereas for second culture media this value was 105.40 U/mg, for third culture media it was 50.25 U/mg, for the fourth culture media it was 302.80 U/mg, for the fifth culture media it was 111.75 U/mg, and for the sixth culture media it was 227.20 U/mg, as shown in Figure 2A.

Optimum Carbon Source

A variety of carbon sources, including lactose, starch, fructose, maltose, sucrose, glucose, and cellulose, were examined at a concentration of 0.5% to see how well they contribute to the production of NK. The results for NK production from *P. aeruginosa* P49 showed that sucrose was the best carbon source with high specific activity (759.88 U/mg protein) compared with other sources, as shown in Figure 2B.

Optimum Nitrogen Source

Numerous kinds of nitrogen sources were examined for their efficiency in NK production. The result demonstrated that the peptone was the best source for NK production from *P. aeruginosa* P49. The specific activity of NK using the peptone was 1023.7 U/mg, while the specific activity of the enzyme was lower when using the nitrogen sources, as shown in Figure 2C.

Effect of salts

Numerous kinds of salts were examined for their efficiency in NK production by *P. aeruginosa* P49. These mineral salt solutions (MS) included: (FeSO₄.7H₂O, CuSO₄.5H₂O, MgSO₄.7H₂O, MnSO₄.7H₂O, CaSO₄.2H₂O, and ZnSO₄.7H₂O). These salts were added separately to the selected medium. The optimum salts for NK production were CaSO₄.2H₂O, which gave a high specific activity of NK (1424 U/mg), as observed in Figure 2D.

Effect of Initial pH

To study the impacts of the initial pH on NK production by *P. aeruginosa* P49, the selected medium was adjusted to different pH values. The optimum pH for NK production was pH 7.5 with high specific activity of NK (1579U/mg protein), while pH 3, 3.5, 4, and 9 didn't give any result for NK production, as shown in Figure 2E.

Purification Steps	Volume (milliliter)	enzyme Activity (U/milliliter)	Protein Concentration (Mg/milliliter)	Specific activity (U/mg)	Total activity (U)	Purification Fold	Yield (%)
Crude enzyme	100	11.7	0.007	1598.8	1117	1	100
Ammonium sulfate precipitation (0-75%)	13	68.16	0.024	2840	954.2	1.78	86
Ion exchange chromatography (DEAE-Cellulose)	64	34.23	0.006	5705	2190.7	3.6	96

Table 2: The purification step of NK from P. aeruginosa P49 isolate.

Optimum temperature

Different temperatures were examined to determine the appropriate temperature for NK production by P. aeruginosa P49. These temperatures were 25, 30, 35, 37, and 40°C. The optimum temperature for NK production was 37°C, with a higher specific activity of NK (1580U/mg protein) as compared with other temperatures, and mentioned in Figure 2F.

Optimum Incubation Period

After 24 hours of incubation, the specific activity reached a maximum value of 1603 U/mg protein. The findings of the experiment were shown in Figure 2G, which showed that the specific activity was 514.5 U/mg after 12 hours. In contrast, the specific activity was 1058 U/mg after 36 hours of incubation, but it had dropped to 1010 U/mg at 48 hours of incubation, as shown in Figure 2G.

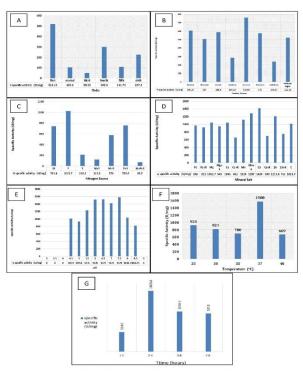


Figure 2: Effect of different A) media; B) Carbon sources; C) Nitrogen sources (M: Meat; P: Peptone; Y: Yeast); D: Salts (C: Control); E: pH; F: Temperature; G: Incubation period on NK production from P. aeruginosa P49 under the optimum conditions.

Purification of NK

The crude extract was purified by precipitation with ammonium sulfate. For the purpose of this

investigation, the ammonium sulfate saturation ratios ranging from 0 to 75% were used for the NK precipitation. After that, the second step for NK purification, which was ion exchange chromatography technique (DEAE-cellulose), was utilized after precipitation by ammonium sulfate (0-70%). The findings in Figure 3 showed one peak of protein in the wash stage. These results suggested that NK carried a positive charge comparable to the charge of the resin under the experimental circumstances, and then the fractions that had activity were collected. As indicated in Table 2, the yield reached 96% when this step's specific activity was 2190.7U/mg protein with a purification fold of 3.6. In contrast, the elution step shown in Figure 3 displayed five peaks of protein when it was eluted by gradient salt; nonetheless, there was no NK activity present. All steps of purification are illustrated in Table 2.

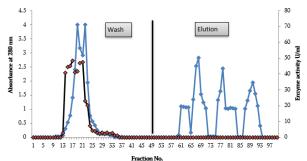


Figure 3: Ion exchange chromatography for NK purification from P. aeruginosa P49 by using DEAE-Cellulose column (23×1.7 cm) equilibrated with Tris-HCl buffer (0.02 M, pH 8.0), eluted with Tris-HCl buffer and NaCl gradient 0 M-1.0 M at a flow rate of 30 milliliter/hour.

Discussion

After all these examinations, 150 isolates of P. aeruginosa were obtained. All isolates of P. aeruginosa were subjected to VITEK2 system to confirm their species.

Pseudomonas aeruginosa is a leading cause of illness and death in hospitalized patients owing to its high level of antibiotic resistance [28]. This study asserts the same evidence as MKK (2019), who indicated the isolation frequency of the most common bacterium in diverse clinical specimens to be *P. aeruginosa* [29]. Similarly, 35.1% (40 out of 114) of patient samples proved to have P. aeruginosa through the documentation of Shatti (2022) [30]. In the same way, Shilba (2015) mentioned the infection of 37.9% of the patients admitted to Karbala hospitals with this

microbe [31]. A considerable difference in protease vields between strains existed. This difference may result from the strain-specific ability of P. aeruginosa to produce varied protease enzymes, e.g., Nattokinase (NK) [32]. Genetic diversity [33] and varied sources of isolation, yet from the same species, may also account for such differences [34].

Choubey (2016) conducted research and revealed 18 different bacterial strains with the ability to produce NK through the formation of haloes in skim milk agar plates [35]. Chandrasekaran et al. (2015) showed in their work that strains of Pseudomonas aeruginosa grown on corresponding media showed very high enzyme production ability [36]. Such variation in the activity of NK in different strains of P. aeruginosa can be the result of the intrinsic variation in the origin and type of bacteria and genetic content [37]. Based on these experimental results, a nutrient-rich medium combining peptone and yeast extract was selected to optimize NK yield. Among these components, peptone proved most effective for enzyme production, owing to its rich composition of proteins, essential vitamins, and other bioactive compounds that support microbial metabolism [38].

Experimental evidence showed that supplementation with sucrose provided maximum enzyme biosynthesis, and the said carbohydrate was used specifically based on its ability to activate NK production. Earlier work has shown that composition of the growth medium significantly affects exoenzyme secretion and carbon and nitrogen substrates are especially important regulatory elements in this respect [25,39].

A study performed by Ibrahim (2015) proved that different organic compounds of nitrogen, such as beef extract, casamino acids, peptone, and skim milk, could promote the synthesis of protease in different species of bacteria with varying efficacy depending upon the microbe strain [40]. In addition to this, peptone is also a highly nutrient-dense supplement with rich amounts of essential micronutrients (vital elements and vitamins) and protein derivatives (amino acids and peptides) important for proper microbe growth and metabolism [41].

As compared to the other studied cations, calcium ions exhibited the most noteworthy enhancement in the synthesis of protease. The majority of the other ions that were examined resulted in a considerable decrease in the enzyme production. It has been shown that calcium ions may stimulate the synthesis of proteases in a variety of species. In addition, the presence of Ba ions increased the amount of protease that Bacillus sp. produced. It has been found that the active shape of the enzyme was maintained at high temperature despite the presence of these metal ions, which prevented the enzyme from being denatured by

heat. Bacillus sp. had their protease production severely inhibited when the medium was supplemented with zinc, copper, iron, and cobalt, and this was most noticeable at high cation concentrations [41]. The addition of MgSO₄ and CaCl₂ led to an increase in the amount of NK produced by B. subtilis MX-6 [42].

In general, the impact of pH on enzyme production may be related to the fact that it plays a role in the solubility of the medium's nutritious elements, has an influence on the ionization of the substrate, and makes the substrate available to the microbe. In addition, pH has an influence on the stability of the enzyme. On the other hand, the fermentation process and the pace of enzyme production are often affected by the pH of the culture medium. The pH value of the medium influences cellular physiology to a significant extent via alteration of plasma membrane conformation. These conformation changes then modify membraneassociated ribosomes and their attendant protein biosynthesis activities [43].

Chandrasekaran et al. (2015) documented the ideal condition being pH 7 at 1514 U/ml and 2.8 mg/ml for the highest NK activity of P. aeruginosa. Equally, Ju et al. (2019) also discovered the ideal condition to be pH 7.0 when they registered the highest NK production of 3284 ± 58 IU/milliliter from their *P. aeruginosa* P49 strain [36,44].

Temperature is a vital factor that needs to be properly controlled in biological processes since its optimum range usually differs greatly between different organisms [45]. For example, Chandrasekaran et al. (2015) recorded the highest NK activity from P. aeruginosa at 25°C with 1514 U/ml and 2.8 mg/ml. Against this background, Ju et al. (2019) documented Bacillus subtilis producing its highest NK production of 3284 ± 58 U/mg at 30° C [36,44]. High temperatures are also recognized to disrupt or destroy the structure of proteins and reduce their protease activity [46]. Aside from its direct effect on enzymes, temperature also affects the availability of oxygen and the mass transfer rate in the fermentation broth and the metabolic activity rates in microorganisms [46].

The variations in yield of NK according to the temperature may be due to changes in the culture conditions that occur during these intervals, such as a decrease in oxygen and nutrients and an accumulation of toxic metabolites that prevent the development of bacteria. According to the findings of certain investigations, the manufacturing of the enzyme began in the earliest stages of the development of the microorganism [45].

In agreement with the results of this study, previous studies indicated that the optimal incubation period is 24 hours for the production of NK from *Bacillus* spp. and *P. aeruginosa* [36,44].

The simplicity of the charge differences-dependent separation principle, potential of reactivation for repetitive utilization, large capacity, simple handling, high power of resolution, and good separation are only a few of the numerous benefits offered by DEAE-Cellulose resin [47].

Author Contributions

Shahad Abdullah Shwan directed the research work, collection of data, and preliminary drafting of the manuscript.

Ali Jabbar Reshak supervised the research, helped with the interpretation of data, and critically reviewed the manuscript.

The final version of the manuscript was read and approved by both authors.

Competing Interest

The authors declare no conflict of interest concerning the publication of this manuscript.

References

- Cai D, Zhu C, Chen S. Microbial production of nattokinase: current progress, challenge and prospect. World Journal of Microbiology and Biotechnology, (2017); 33(2017): 1–7.
- Wu H, Zhang Q, Xu P, Chen J, Duan L, et al. Nattokinase Promotes Post-stroke Neurogenesis and Cognition Recovery via Increasing Circulating Irisin. Journal of Agricultural and Food Chemistry, (2023); 71(30): 11418-11428.
- Yoo HJ, Kim M, Kim M, Lee A, Jin C, et al. The effects of nattokinase supplementation on collagen-epinephrine closure time, prothrombin time and activated partial thromboplastin time in nondiabetic and hypercholesterolemic subjects. Food & Function, (2019); 10(5): 2888-2893.
- Zhou X, Liu L, Zeng X. Research progress on the utilisation of embedding technology and suitable delivery systems for improving the bioavailability of nattokinase: a review. Food Structure, (2021); 30(2021): 100219.
- Pagnoncelli M, Fernandes M, Rodrigues C, Soccol C (2017) Nattokinases. Current developments in biotechnology and bioengineering: Elsevier. pp. 509-526.
- Yuan L, Liangqi C, Xiyu T, Jinyao L. Biotechnology, bioengineering and applications of Bacillus nattokinase. Biomolecules, (2022); 12(7): 980.
- Jayaprakashvel M, Sami M, Subramani R (2020) Antibiofilm, antifouling, and anticorrosive biomaterials and nanomaterials for marine applications. Nanostructures for Antimicrobial and Antibiofilm Applications: Springer. pp. 233-272.
- Wood SJ, Kuzel TM, Shafikhani SH. Pseudomonas aeruginosa: Infections, Animal Modeling, and Therapeutics. Cells, (2023);
- Morihara K, Homma J (2018) Pseudomonas proteases. Bacterial enzymes and virulence: CRC Press. pp. 41-80.
- Gimenes NC, Silveira E, Tambourgi EB. An overview of proteases: production, downstream processes and industrial applications. Separation & Purification Reviews, (2021); 50(3): 223 - 243.
- Vantamuri AB, Manawadi SI, Guruvin SK, Holeyannavar VM, Shettar DS. Production of laccase by Ganoderma sp. in submerged fermentation. Journal of Advanced Research, (2019); 10(04): 67-71.

- 12. Li D, Hou L, Hu M, Gao Y, Tian Z, et al. Recent advances in nattokinase-enriched fermented soybean Foods: A review. Foods, (2022); 11(13): 1867.
- Ciurko D, Chebbi A, Kruszelnicki M, Czapor-Irzabek H, Urbanek AK, et al. Production and characterization of lipopeptide biosurfactant from a new strain of Pseudomonas antarctica 28E using crude glycerol as a carbon source. The Royal society of chemistry, (2023); 13(34): 24129-24139.
- 14. Kumar SS, Haridas M, Abdulhameed S. A novel fibrinolytic enzyme from marine Pseudomonas aeruginosa KU1 and its rapid in vivo thrombolysis with little haemolysis. International Journal of Biological Macromolecules, (2020); 162(2020): 470-
- Granato PA, Granato PA. Laboratory Manual and Workbook in Microbiology: Applications to Patient Care. 2011. McGraw-Hill
- Bergey DH. Bergey's manual of determinative bacteriology. 1994; 1-1063. Lippincott Williams & Wilkins
- Mackie TJ. Mackie & McCartney practical medical microbiology. 1996; 1-978. Churchill Livingstone
- Putra ARS, Effendi MH, Kurniawan F. Investigation of extended spectrum beta-lactamase (ESBL) producing Escherichia coli by VITEK-2 on dairy cows in Surabaya, Indonesia. Biochemical and Cellular Archives, (2020); 20(2): 6773-6777.
- Olajuyigbe FM, Ajele JO. Production dynamics of extracellular protease from Bacillus species. African Journal of Biotechnology, (2005); 4(8): 776-779.
- Mukhtar H, Haq I. Comparative evaluation of Agroindustrial byproducts for the production of alkaline protease by wild and mutant strains of Bacillus subtilis in submerged and solid state fermentation. The Scientific World Journal, (2013); 2013(1):
- 21. Senior BW. Investigation of the types and characteristics of the proteolytic enzymes formed by diverse strains of Proteus species. Journal of Medical Microbiology, (1999); 48(7): 623-628.
- 22. Bradford MM. A rapid and sensitive method for the quantitation of microgram quantities of protein utilizing the principle of protein-dye binding. Analytical Biochemistry, (1976); 72(1-2): 248-254.
- AL-Sa'ady AJ, Hilal MH. Determination of the optimum conditions for extracting polyphenol ox-idase and laccase enzymes from malva parviflora and their role in the decolorization of some dyes. Iraqi Journal of Science, (2020); 61(2): 306-313.
- Whitaker JR, Bernhard RA. Experiments for: an introduction to enzymology. 1972. Whiber Press
- Caraveo L, Medina H, Rodríguez-Buenfil I, Montalvo-Romero C, Evangelista-Martínez Z. A simple plate-assay for screening extracellular naringinase produced by streptomycetes. Journal Microbiol Methods, (2014); 102(2014): 8-11.
- 26. Lakshmi BKM, Sri PR, Devi KA, Hemalatha KPJ. Media optimization of protease production by Bacillus licheniformis and partial characterization of Alkaline protease. International Journal of Current Microbiology and Applied Sciences, (2014); 3(5): 650-659.
- 27. Schütte H, Hummel W, Tsai H, Kula MR. L-Leucine dehydrogenase from Bacillus cereus: production, large-scale purification and protein characterization. Applied Microbiology and Biotechnology, (1985); 22(1985): 306-317.
- Hatite Al-Daraghi WA, Abdulkadhim Al-Badrwi MS. Molecular Detection for Nosocomial Pseudomonas aeruginosa and its Relationship with multidrug Resistance, Isolated from Hospitals Environment. Medico-Legal Update, (2020); 20(1): 631-636.
- 29. MKK F, MA R, Rashid SS, MHM N. Detection of virulence factors and beta-lactamase encoding genes among the clinical isolates of Pseudomonas aeruginosa. Journal of International Pharmaceutical Research, (2019); 45(2019): 190-202.
- Shatti HH, Al-Saeed WM, Nader MI. Effect Biofilm Formation in Pseudomonas aeruginosa Resistance to Antibiotic. Mustansiriya Medical Journal, (2022); 21(1): 13-17.
- Shilba AA, Al-Azzawi RH, Al-Awadi SJ. Dissemination of Carbapenem Resistant Pseudomonas aeruginosa among Burn

- Patients in Karbala Province\Iraq. Iraqi Journal of Science, (2015); 56(3A): 1850–1857.
- Shawkat MS, Kareem AA. Primary and Secondary Screening of Pseudomonas aeruginosa for Protease Production. Iraqi Journal of Biotechnology, (2022); 21(2): 505-510.
- Zhao K, Liu H, Song W, Wu J, Gao C, et al. Combinatorial mutagenesis of Bacillus amyloliquefaciens for efficient production of protease. Systems Microbiology and Biomanufacturing, (2023); 3(3): 457–468.
- 34. Naveed M, Nadeem F, Mehmood T, Bilal M, Anwar Z, *et al.* Protease—a versatile and ecofriendly biocatalyst with multi-industrial applications: an updated review. Catalysis Letters, (2021); 151(2021): 307–323.
- Choubey D, Dhusia K, Gupta N, Viswan NA, Mandal S. Identification and characterization of Nattokinase producing bacteria and optimization of enzyme production. (2016).
- Chandrasekaran SD, Vaithilingam M, Shanker R, Kumar S, Thiyur S, *et al.* Exploring the in vitro thrombolytic activity of nattokinase from a new strain *Pseudomonas aeruginosa* CMSS. Jundishapur Journal of Microbiology, (2015); 8(10): e23567.
- 37. Shafique S, Bajwa R, Shafique S. Screening of *Aspergillus niger* and A. flavus strains for extra cellular alpha-amylase activity. Pakistan Journal of Botany, (2009); 41(2): 897–905.
- 38. Espoui AH, Larimi SG, Darzi GN. Optimization of protease production process using bran waste using Bacillus licheniformis. Korean Journal of Chemical Engineering, (2022); 39(3): 674–683.
- 39. Al-Sa'ady AJR, Aziz GM. Optimization of Lovastatin Production from A Local Isolate of Aspergillus terreus A50 in Solid State Fermentation by Classical and Statistical Methods. Iraqi Journal of Science, (2020); 61(10): 2525–2539.
- Ibrahim ASS, Al-Salamah AA, Elbadawi YB, El-Tayeb MA, Ibrahim SSS. Production of extracellular alkaline protease by new halotolerant alkaliphilic Bacillus sp. NPST-AK15 isolated from hyper saline soda lakes. Electronic Journal of Biotechnology, (2015); 18(3): 236–243.
- 41. Singh S, Bajaj BK. Bioprocess optimization for production of thermoalkali-stable protease from *Bacillus subtilis* K-1 under solid-state fermentation. Preparative Biochemistry and Biotechnology, (2016); 46(7): 717–724.
- 42. Man LL, Xiang DJ, Zhang CL. Strain screening from traditional fermented soybean foods and induction of nattokinase production in *Bacillus subtilis* MX-6. Probiotics Antimicrob Proteins, (2019); 11(2019): 283–294.
- Hussein SI, Aziz GM, Shanshal RM, Ghani AL. Determination the optimum conditions of laccase production from local isolate of *Streptomyces sp.* using solid state fermination. The Iraqi Journal of Agricultural Sciences, (2018); 49(4): 685-693.
- 44. Ju S, Cao Z, Wong C, Liu Y, Foda MF, *et al.* Isolation and optimal fermentation condition of the *Bacillus subtilis* subsp. natto strain WTC016 for nattokinase production. Fermentation, (2019); 5(4): 92.
- Sharma KM, Kumar R, Panwar S, Kumar A. Microbial alkaline proteases: Optimization of production parameters and their properties. Journal of Genetic Engineering and Biotechnology, (2017); 15(1): 115–126.
- Son HF, Cho IJ, Joo S, Seo H, Sagong HY, et al. Rational protein engineering of thermo-stable PETase from Ideonella sakaiensis for highly efficient PET degradation. Acs Catalysis, (2019); 9(4): 3519–3526.
- Abdalah ME, Al-Saady AJR, ALBahrani MHA, Al-Obaidi MJ, Abd Al-Hussein MY. Cleavage of mucin and salivary agglutinin by partial purified protease from a native strain of *Streptococcus mutans* AN67. Journal of Global Pharma Technology, (2018); 10(06): 404–410.



This work is licensed under a Creative Commons Attribution-NonCommercial 4.0 International License. To read the copy of this

license please visit: $\frac{https://creative commons.org/licenses/by-nc/4.0/}{}$