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## Isolation, Identification And Characterization Of Keratinase Producing Bacteria From Poultry Farms.

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### ABSTRACT

To meet consumer demand, the global poultry business is constantly expanding; hence it is crucial to ensure the biosafety of the poultry farms. In poultry farms, the main sources of bacterial development, increased antibiotic resistance, and environmental dispersion may be poultry feed, litter, and water. The focus of this research is on the identification and characterization of bacterial pathogens found in commercial poultry farms, drinking water samples, feed samples, and litter samples. The 100 samples were gathered from chicken farms in different districts of Telangana (Jagtial, Sircilla, Karimnagar, Nizamabad) and transported to the lab for additional examination. Serial dilution was used to analyse the samples, which were then examined for colony morphology, microscopic features, and the biochemical traits of the bacteria *Escherichia coli*, *Pseudomonas* spp., *Proteus* spp., *Salmonella* spp., *Shigella* spp., *Bacillus* spp., *staphylococcus* spp., and *Streptococcus* spp. The results of this research indicate that the main sources of bacterial infections are chicken feed, drinking water samples and poultry litter.

**Keywords:** Keratin, Keratinase, Microbes, Poultry, biochemical test.

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## INTRODUCTION

Humans are at risk of contracting food-borne illnesses because of contaminated poultry products caused by bacterial infections. Food-related illnesses caused by germs like *Salmonella* spp. and *E. coli* are spread through poultry meat [1]. The elements that affect the contagious transmission of bacterial infections must be taken into consideration to prevent them from emerging from poultry habitats. Poultry feed (starters, growers, finishers, and layers) and drinking water from poultry farms are contaminated during processing [2], storage while adding the raw material ingredients, and ultimately during exposure of the product to the environmental microbes [3,4]. The primary worry of industrialized nations is the safety of drinking water and chicken feed in poultry farms. All birds' basic needs are food and water, and contaminated feed results in significant economic loss [5]. The abundance of different bacteria in chicken diets depends on several variables, including the nutrient makeup of the feed, pH, oxygen tension, and water activity. According to Aisha Ali [6] and [7], microbial proliferation can reduce feed quality by replacing the nutrients in the feed, and damage done during feed processing or the creation of toxins has a detrimental influence on animal health [8]. The safe use of poultry products increases the length of time that must pass before microbiological contaminants are assessed and exposed, which causes the customer undue harm [9]. According to Osei et al. [10], the material used to create poultry beds, feather waste, manure, and leftover feed are the main sources of litter utilized by poultry. Bacterial infections can spread through leftover poultry.

The expanding population and the commerce of animals are under increasing health and ecological hazard [11]. Numerous illnesses in poultry are caused by bacterial infections such *Escherichia coli*, *Staphylococcus*, and *Bacillus* species. Although research is being done to repurpose chicken waste material for agricultural and feed additives for other animals [12], efficient poultry waste disposal practices are not currently in practice in developing countries [13,14]. The use of antibiotics in animal feed at low doses for extended periods of time may contribute to microbial resistance. Due to the risk of resistance genes propagating to the environment, it is possible for resistant bacteria to enter sewage systems through chicken waste [15,16]. As a result, it was suggested that the current research work isolate, characterize, and identify the isolated pathogenic bacteria linked to poultry as a source of the potential contamination in poultry farms [17]. Keratinases (EC.3.4.99.11) belong to the group of serine proteases capable of degrading keratin. It is an extracellular enzyme produced in a medium containing keratinous substrate such as feathers and hair. Keratinases have applications in traditional industrial sectors including feed, detergent, medicine, cosmetics and leather manufacturers, they also find application in more recent fields such as prion degradation for treatment of the dreaded mad cow disease, biodegradable plastic manufacture and feather meal production. Hence present study focuses on the production of enzyme keratinase. Because of the numerous potential uses of keratinases, study was undertaken to screen a bacterium that produces active keratinase.

## MATERIALS AND METHODS

### Sample Collections

Six chicken farms in the Tamil Nadu state of India's Jagtial, Sircilla, Karimnagar, Nizamabad districts provided samples of poultry litter, poultry water, and commercial feed. Litter samples from chicken farms were chosen at random, and feed and water samples were gathered directly from the chicken farms' feeding operations. In total, 100 samples were acquired from each farm, including at least 6 litter, 6 feed and 6 water samples. The samples were taken in a sterile bag and container, and they were clearly labelled.

### Analysis of Samples

The Mathan et al. [18] protocol was followed for the isolation and subsequent identification of the bacterial pathogens. 1 g of poultry litter, 1 g of chicken feed, and 1 ml of water sample were inoculated into a 100 ml nutritional broth and then incubated for an additional 24 hours at 37°C. All the samples that were obtained underwent a total bacterial count on nutrient agar plates before being serially diluted at concentrations of 1 in 10. After 24 hours of incubation, the colony formation is expressed in CFU/mg using a colony counter [19].

## Characterization pf bacterial isolates

### Gram Staining and Microscopy

The isolated bacterial cultures were characterized by morphological and biochemical activity analysis. The cultures were examined via gram staining and microscopy. Further the gram negative and positive isolated bacterial cultures were further examined under microscope. After 24 hours of incubation, the colonial morphology of a bacterial culture including its size, shape, colour, and texture was inoculated into different media [20,21].

### Biochemical tests

Bio-chemical tests such as the catalase test, oxidase test, citrate utilisation test, Methyl red test, Indole test, Voges-proskauer test, triple sugar iron test, and carbohydrate fermentation tests were carried out using isolated bacterial strains that displayed distinctive colony morphology [22,23]. The selected bacterial cultures were allowed for large scale growth and culture was centrifuged to collect the supernatant. The keratinase enzyme is an extracellular enzyme where bacteria excrete outside of plasma membrane. The collected supernatant was used for keratinase enzyme activity via keratinase assay.

### Keratinase Assay

The isolated potent strain (KP9) was transferred to 25 ml of the seed medium containing (g/L) peptone, 5; yeast extract, 1.5; beef extract, 1.5 and sodium chloride, 5; (pH 7 ± 0.2) and incubated at 37°C on rotary shaker at 180 rpm for 24 h. This was used as the inoculum for the keratinase enzyme production process. Submerged fermentation process was carried out by inoculating pure culture of potent isolate into the production medium (Suntomsuk and Suntomsuk, 2003). The production medium contains (g/L) Feather meal, 10; Yeast extract, 0.1; MgSO<sub>4</sub>, 0.1; NH<sub>4</sub>Cl, 0.5; K<sub>2</sub>HPO<sub>4</sub>, 0.3; KH<sub>2</sub>PO<sub>4</sub>, 0.3; NaCl, 0.5. Initial pH of the medium was adjusted to 7.2 ± 0.2 with Tris-HCl buffer. The medium was sterilized in an autoclave for 15 min at 121°C. The production medium was inoculated with 5% (v/v) of 24 h old inoculum culture of potent isolate containing approximately 2×10<sup>6</sup> cells/ml. The flasks were incubated on a rotary shaker at 37°C and 180 rpm for 4 days. The 10 ml fermented broth was centrifuged at 10,000 rpm for 20 min, and the supernatant was used as crude enzyme. The keratinase enzyme activity was assayed by taking 1.0 ml of crude enzyme properly diluted in Phosphate buffer (0.05 M of pH 7.0) was incubated with 1 ml of keratin solution at 50 °C in a water bath for 10 min, and the reaction was ceased by the addition 2.0 ml 0.4 M Trichloroacetic acid (TCA). The resulted precipitate was removed by centrifugation at 10000 rpm for 20 min. The absorbance of the supernatant was determined at 280 nm against a control. The control was prepared by incubating the enzyme solution with 2.0 ml TCA without the addition of keratin solution. One unit (U/ml) of keratinase enzyme activity was defined as an increase of corrected absorbance of 280 nm with the control for 0.01 per minute under the conditions described above.

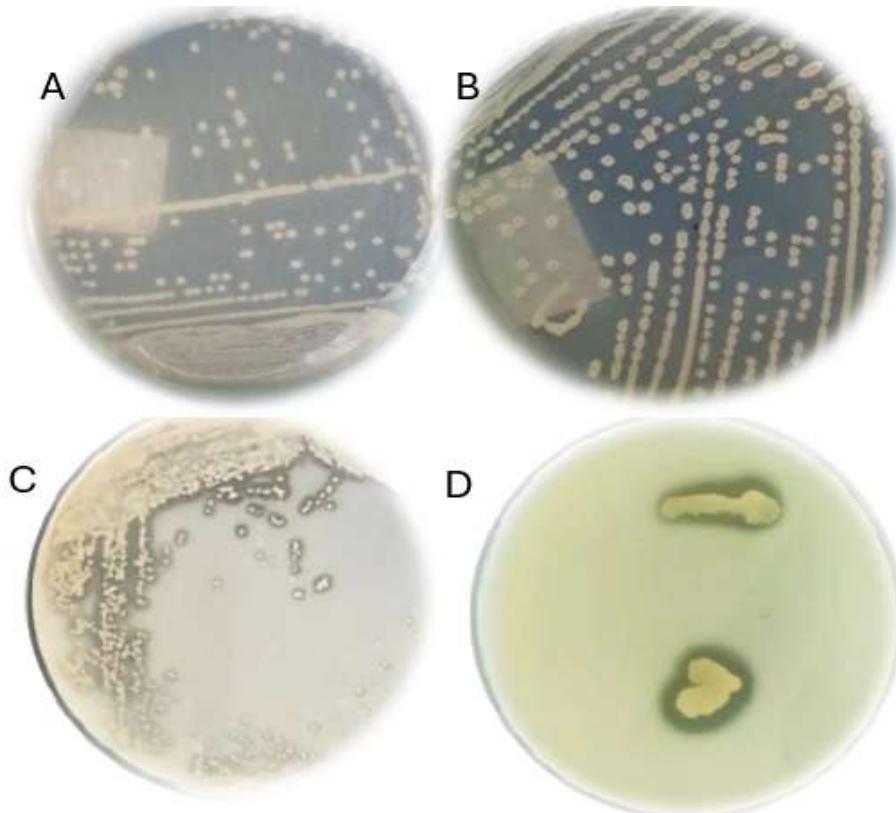
## RESULTS AND DISCUSSION

**Table 1: Represent the total plate count of bacterial isolates for the obtained samples (commercial Feed, poultry litter and water).**

| S. No | CFU X Dilution factor |
|-------|-----------------------|
| 1     | 3.9x10 <sup>9</sup>   |
| 2     | 3.5x10 <sup>9</sup>   |
| 3     | 5.5x10 <sup>8</sup>   |
| 4     | 2.5x10 <sup>8</sup>   |
| 5     | 3.2x10 <sup>10</sup>  |
| 6     | 1.5x10 <sup>8</sup>   |
| 7     | 3.5x10 <sup>9</sup>   |
| 8     | 4.9x10 <sup>8</sup>   |
| 9     | 3.2x10 <sup>9</sup>   |
| 10    | 3.8x10 <sup>9</sup>   |

The goal of the current research was to identify microbes with the potential to keratin hydrolysis (keratinase activity) from poultry feed, litter, and water from poultry feeding operations. Using varied and selective culture media, 6 samples from each farm were evaluated. There were 5 different poultry farms involved. Here in the present study

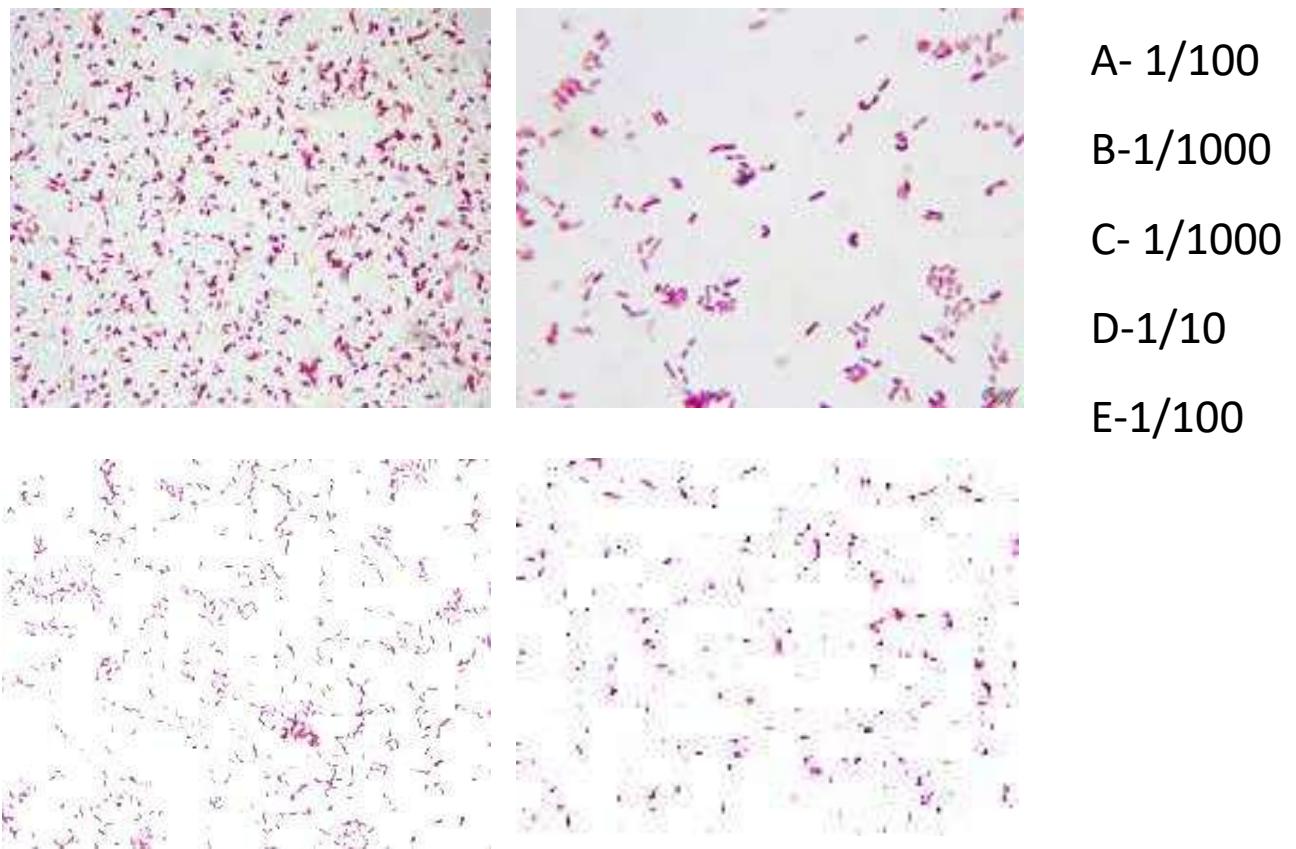
**Bacterial Isolates' Characterization:** The collected bacterial isolates were subjected to various culture medium and biochemical characteristics (Table 2), which were chosen based on colony morphology, growth, and number of biochemical tests.



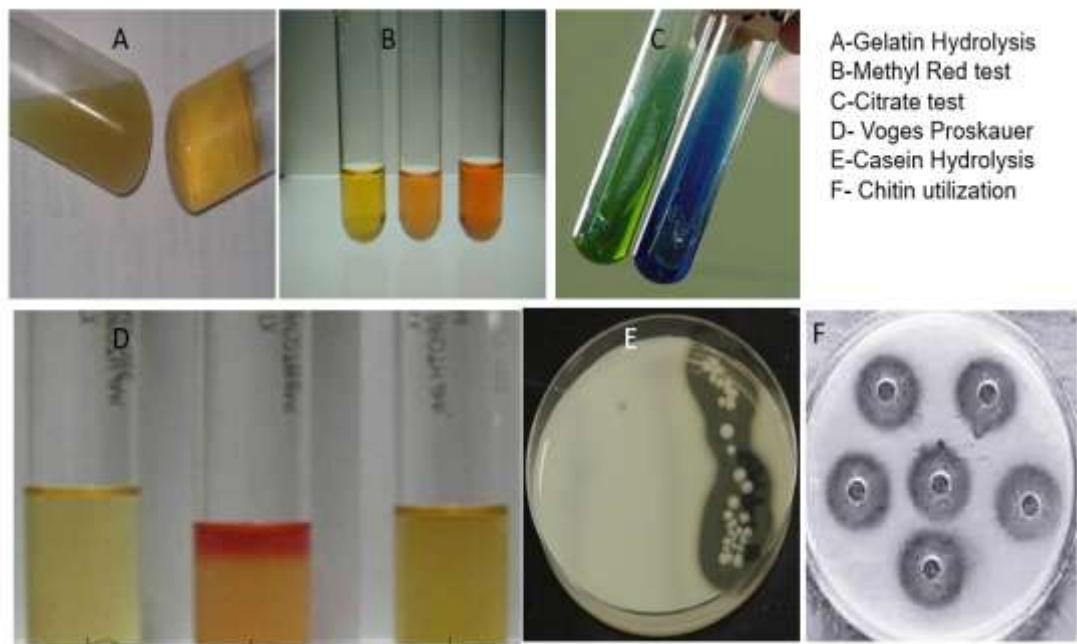
**Figure 3: Isolated bacterial species from the poultry environment samples and setting. The key organism reported *Bacillus licheniformis* (A), *Bacillus substilis* (B), *Shigella spp* (C) and *Proteus spp* (D).**

**Table 2: Morphological identification of bacterial isolates**

| <b>Morphological characterization of isolated bacteria</b> | <b>Gram Staining</b> | <b>Shape</b>      | <b>organism</b>           |
|--|----------------------|-------------------|---------------------------|
| Green pigment-producing colonies                           | Gram -ve             | Rod               | <i>Pseudomonas spp</i>    |
| Pink, convex with smooth-edged colonies                    | Gram -ve             | Rod               | <i>Escherichia coli</i>   |
| Colonies with irregular margins, flat and white colour     | Gram +ve             | Rod               | <i>Bacillus spp</i>       |
| Colonies were white and swampy                             | Gram -ve             | Rod               | <i>Proteus spp</i>        |
| Creamy, round, convex smooth Edged colonies                | Gram -ve             | Rod               | <i>Salmonella spp</i> ,   |
| Shiny, spherical colonies on blood agar                    | Gram +ve             | Cocci in chains   | <i>Streptococcus spp</i>  |
| Creamy and convex with Smooth edged colonies               | Gram +ve             | Cocci in clusters | <i>Staphylococcus spp</i> |



**Figure 2: Gram staining and microscopy of isolated bacterial culture isolated samples from poultry environmental setting.**



**Figure 1: Biochemical analysis of isolated bacterial culture isolated samples from poultry environmental setting.**

**Table 3: Bacterial isolates distribution as sample wise poultry feed, poultry litter and water samples collected from the poultry setting.**

| <b>S No</b> | <b>Sample</b>  | <b>Bacterial species reported</b>   |
|-------------|----------------|---|
| 1.          | Poultry feed   | Pseudomonas spp, E. coli, Salmonella spp, staphylococcus spp, Vibrio spp, Shigella spp        |
| 2.          | Poultry litter | Salmonella spp, Pseudomonas spp, E. coli, Streptococcus spp, Bacillus substilis, Shigella spp |
| 3.          | Water sample   | Pseudomonas spp, Shigella spp. Salmonella spp Vibrio spp                                      |

**Table 4: Effect of inoculum size on enzyme activity.**

| <b>S No</b> | <b>Incubation period (h)</b> | <b>Enzyme activity (U/ml)</b> |
|-------------|------------------------------|-------------------------------|
| 1.          | 24                           | 166 ± 2.10                    |
| 2.          | 48                           | 257 ± 2.14                    |
| 3.          | 60                           | 102 ± 11.6                    |
| 4.          | 72                           | 65 ± 1.3                      |

### CONCLUSION

The objective of the present study was to isolate the keratinase producing bacteria from chicken feathers, since it is rich in keratin. The production of enzyme keratinase and optimization of various parameters for the enzyme production were studied. This novel keratinolytic *Bacillus licheniformis* could be a potential candidate for the degradation of feather keratin and in dehairing process in leather industry. The enzyme activity of *Bacillus licheniformis* was reported 257 U/ml. This isolate could be used to produce keratinase for biotechnological applications and effectively used in the large-scale production of enzyme for commercial purposes.

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### REFERENCES

- [1] Ghosh S, Sarkar T, Chakraborty R. Formation and development of biofilm-an alarming concern in food safety perspectives. *Biocatalysis and Agricultural Biotechnology*. 2021;38:102210.
- [2] Farid N, Waheed A, Motwani S. Synthetic and natural antimicrobials as a control against food borne pathogens: A review. *Heliyon*; 2023.
- [3] Onyeze RC, Onah Gloria T, Eluke Obinne C. Bacterial contaminants associated with commercial poultry feeds in Enugu Nigeria. *International Journal of Life Sciences Biotechnology and Pharma Research*. 2013;2:3.
- [4] Rujuta Kedar K, Yashvi Udaybhai R, Devikaben BV, Dhruv GP. Isolation and Identification of pathogenic bacterial contaminants associated with poultry. *International Journal of Current Microbiology and Applied Sciences*. 2021;10(05):294-305.
- [5] Khan MAS, Rahman SR. Use of phages to treat antimicrobial-resistant *Salmonella* infections in poultry. *Veterinary Sciences*. 2022;9(8):438.
- [6] Aisha Ali Rabiu D, Kamalu A, Alhassan M, Manjur S. Isolation and identification of microbial contaminants associated with commercial poultry feeds. *Journal of Applied and Advanced Research*. 2018;3(5):142-147.
- [7] Dasari Venkateswarlu, Arjun P, Samiraj R. Detection of  $\beta$ -lactamase in clinical samples of Prakasam District, Andhra Pradesh. *Research Journal of Pharmacy and Technology*. 2021;14(6):2995-2998.
- [8] Islam MS, Khanam S, Krishna M., Moni M. Isolation, characterization and identification of bacterial isolates from the poultry environment at Rajshahi Metropolis, Bangladesh. *J. Entomol. Zool. Stud.* 2017; 5(4):918-926.
- [9] Nagarajan S, Arjun P, Raaman N, Anup S, Elizabeth Sobhia M, Mohan Das T. Stereoselective synthesis of sugar-based  $\beta$ -lactum derivatives: Docking studies and its biological evaluation. *Tetrahedron*. 2012;68(4):3037-3045.

[10] Osei FB, Boamah VE, Agyare C, Abaidoo RC. Physicochemical properties and microbial quality of water used in selected poultry farms in the ashanti region of ghana. Open Microbiology Journal. 2019; 13(1).

[11] Nagarajan S, Mohan Das T, Arjun P, Raaman N. Design, synthesis and gelatin studies of 4,6-O-butylidene-  $\alpha$ , $\beta$ - unsaturated- $\beta$ -C-glycosidic ketones: application to plant tissue culture. Journal of Materials Chemistry. 2009;26: 4587-4596.

[12] Krishnani KK, Boddu VM, Chadha NK, Chakraborty P, Kumar J, Krishna G, Pathak H. Metallic and non-metallic nanoparticles from plant, animal, and fisheries wastes: potential and valorization for application in agriculture. Environmental Science and Pollution Research. 2022;29(54):81130-81165.

[13] Kelleher BP, Leahy JJ, Henihan A, Dwyer TF, Sutton D, Leahy MJ. Advances in poultry litter disposal technology - A review. Elsevier. 2002;83:27-36.

[14] Adegunloye DV. Microorganisms associated with poultry faeces. Journal of Food, Agriculture and Environment. 2005;4 (1):41-42.

[15] Nagarajan S, Arjun P, Raaman N, Mohan Das T. Regioselective facile one-pot Friedländer synthesis of sugar-based heterocyclic biomolecules. Carbohydrate Research. 2010;345(14):1988-1997.

[16] Diarra M, Malouin F. Antibiotics in Canadian poultry productions and anticipated alternatives. Frontiers in Microbiology. 2014;5:1-15.

[17] Apata DF. Antibiotic resistance in poultry. International Journal of Poultry Sciences. 2009;8(4):404-8.

[18] Mathan P, Patricia T, Aswin A, Gunaseelan S, Karthick R, Mayur K, Narayanan M. Isolation of pathogenic bacteria from poultry wastages at chennai suburban. IOSR Journal of Environmental Science, Toxicology and Food Technology (IOSR-JESTFT). 2013;6(6):50-54.

[19] Santovito E, Greco D, Marquis V, Raspoet R, D'Ascanio V, Logrieco AF, Avantaggiato G. Antimicrobial activity of yeast cell wall products against Clostridium perfringens. Foodborne Pathogens and Disease. 2019;16(9):638-647.

[20] Krishnamoorthy M, Arjun P. Probiotic and antimicrobial activity of bacteria from fermented toddy of Cocus nucifera. Journal of Academia and Industrial Research. 2012;1(3):127-131.

[21] Kianpour S, Ebrahiminezhad A, Mohkam M, Tamaddon AM, Dehshahri A, Heidari R, Ghasemi Y. Physicochemical and biological characteristics of the nanostructured polysaccharide-iron hydrogel produced by microorganism Klebsiella oxytoca. Journal of basic microbiology. 2017;57(2):132-140.

[22] Microbiology: A Laboratory Manual (9th Edition) By James Cappuccino, Natalie Sherman; 1993.

[23] Muthusamy K, Sajen SH, Sharmila SMI, Murugan P, Ramaiah S. Microbiota from Intestinal Gut of Dawkinsia filamentosa (Valenciennes, 1844) as Indicator of Environmental Condition. Asian Journal of Fisheries and Aquatic Research. 2023; 22(2):15-24.

[24] Bauer AW, Kirby WM, Sherris JC, Turck M. Antibiotic susceptibility testing by a standardized single disk method. American Journal of Clinical Pathology. 1966;45(4): 493-6.

[25] The European committee on antibiotic susceptibility testing - EUCAST. Isolaztion and identification of bacterial isolates from poultry and fish feeds sold in abraka, delta state, nigeria. Journal of Industrial Technology. 2017;2(1):14-20.

[26] Daoud A, Saud A, Arjun P. Somatic embryogenesis and in vitro plant regeneration of Bacopa monnieri (Linn.) Wettst., a potential medicinal water hyssop plant. Saudi Journal of Biological Sciences. 2021;28:353-359.

[27] Islam MR, Martinez-Soto CE, Lin JT, Khursigara CM, Barbut S, Anany H. A systematic review from basics to omics on bacteriophage applications in poultry production and processing. Critical Reviews in Food Science and Nutrition. 2021;1-33.