

### DEGRADATION OF KERATIN SUBSTRATE BY MICROORGANISMS: A REVIEW

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Keratin is an insoluble proteins present in the epidermis and important composition of feather, hair, wool, nail, hoofand horns.Keratin is important composition and provide strength to the keratinous substrate.In the environment, feather, hair, wool, nail, hoof and horns are considered as waste material and are major causes of environmental pollution.Keratin is also found in products of agro-industrial processing, dairy industry, poultry farms and slaughterhouses.These keratinous waste are structurally stable and very hard to degrade.The keratinous waste can be managed by biological degradation.Microbial keratinase present in some species of bacteria and fungi is best for the degradation of keratin waste in natural condition.The present review paper explains keratinolytic microorganisms and their keratinase for the biological degradation of keratin waste.

Keywords: keratinolytic microorganisms, keratinase, biological degradation and waste material.

#### Introduction

Keratin waste are the waste produced by leather industry, poultry industry, slaughter houses and hospitals. Because of their resistance to biodegradation, keratinous wastes in the environment pose a significant threat to the ecosystem. Burning, landfilling, heating under pressure and chemical hydrolysis are common ways for their treatment. These techniques consume a lot of energy are not environmentally friendly. The development biological/green of technologies for the breakdown of keratinous wastes has become critical in light of these issues. Microorganisms degrade keratin and produce biotechnologically important products such as keratinolytic enzymes, peptides, and amino acid-rich keratin hydrolysates. This

is a cost-effective and simple method that reduces the environmental problem posed by these wastes in an environmentally friendly manner.

Microbial keratin degradation has been widely explored in recent years, as various bacteria, fungi, and actinomycetes are found to be good keratin degraders and are easily isolated from soil and keratin-rich wastes materials.

Degradation by Bacteria:

Most of the keratinolytic bacteria are Gram-positive bacteria, but Gram-negative bacteria are also found to degrade keratin by few strains. The common species of Keratinolytic bacteria include *Bacillus*, *Brevibacillus*, *Chryseobacterium*, *Stenotrophomonas*, *Pseudomonas*, *Keratinibaculum*, *Paenibacillus*, *Meiothermus*, *Rhodococcus*, Achromobacter, Exiguobacterium and Aeromonas<sup>1,2,3,4</sup>. Several researchers have indicated that species of Bacillus have high efficiency to degrade keratin namely B. licheniformis, B. pumilus, B. cereus, and B. subtilis <sup>5,6,7</sup>. Some new species of keratinolytic fungi are also isolated including Bacillus safensis LAU 13 was isolated by Lateef et al.  $(2015)^8$  and Bacillus cereus wasisolated by Ahmadpour et al.  $(2017)^9$ .

Degradation by Fungi:

Several fungal species contribute enormously in the recycling of recalcitrant keratinous waste. Most of the research papers for keratin degrading capability have registered mostly fungi as important keratinolytic organisms. Keratin degradation capability among the fungi varies due to some factors such as culture conditions, sources of carbon, nitrogen, and energy source. The morphological feature of filamentous fungi facilitates keratin degradation through the firm attachment of mycelia andpenetration of keratin substrates by fungal hyphae. Keratin degradation capability of fungi has been documented in various literature studies. Various species and strains of fungi have been reported for keratinolytic property including species of Aspergillus, Penicillium. Chrvsosporium. Fusarium. *Microsporum*, Trichophyton and Acremonium, Aphanoascus, Chaetomium, *Penicillium*<sup>10</sup>. Some other species of fungi which were isolated from soil and poultry waste, have also been recorded for keratinolytic activity including species of Scopulariopsis, Myceliophthora, Candida, Cladosporium. Metarrhizium. Neurospora. Cunninghamella and Westerdikella<sup>11,12</sup>. Morophology of fungi are suitable for keratin degradation as their mycelium assist in attachment tokeratin substrate. After attachment fungi perforate keratin by their hyphae. The ideal example to

understand the mechanism of keratin degration by nonpathogenic fungi is, *Onygena corvine*, which is a very effective keratin degrader<sup>13</sup>.

Dermatophytes are а category of keratinolytic fungi that includes certain members of the keratinolytic fungi. These fungi secrete keratinases, which are necessary for their penetration into the body<sup>15</sup>. The addition of starch and maltose to the fermentation medium at the right pH can improve keratinase synthesis and keratin breakdown. Fungi are the most effective at degrading feathers. The degradation of keratin substrate was achieved by producing nutritional medium with C, N, S, and energy sources in a static situation. The degradation of keratin is also providing achieved by optimum environment condition to the keratinolytic fungi such as pH and temperature.

Degradation by Actinomycetes:

Most of the keratin degrading microbes are among bacteria, fungi, and actinomycetes. Actinomycetese are important keratin degraders because they can hydrolyze a wide range of keratin wastes, including feathers, hair, and wool<sup>16</sup>. In some studies, isolation and keratinolytic activity of some actinobacteria like Actinomaduraand, Actinoalloteichus have been reported for hydrolysis of keratin substrates<sup>17,18</sup>. Some members of Streptomyces, viz: S. fradiae and S. gulbargensis have also been reported to show keratinolytic activity<sup>19</sup>. Species of Nocardiopsis have been found effective for keratinolytic activity on feather waste<sup>20</sup>

Keratinase from Actinomyces:

The keratinase obtained from the species of actinomyces and Streptomyces was efficient in degradation of different substrates including keratin azure, human hair, cock feathers, and collage as indicated by experiments.Another important keratinase from actinomycetese was reported by Elhoul et al.,(2016)as a novel thermostable keratinasefrom strain of *Actinomadura viridilutea* and these keratinase displayed effective chicken feather hydrolysis within 96 hours<sup>21</sup>.

Similarly, Ningthoujam et al., (2016) reported the production of keratinase from*Amycolatopsis* sp. isolated from a limestone habitat, showing promising keratinolytic activity on keratin azure and chicken feather substrates<sup>22</sup>.

Mechanism of Keratin Degradation by Microorganisms:

The exact method by which microbes degrade keratin is unknown. According to some studies, keratinolysis is caused by two mechanisms: sulfitolysis, which involves the reduction of disulfide bonds, which involves and proteolysis, the hydrolysis of peptide bonds of keratin.Sulfitolysis requires the presence of disulfide reductases enzyme in the microorganism which work in combination keratinases for the with complete degradation of keratin $^{23}$ .

According to some researches keratinases are more effective keratin degraders when combine with some enzymes produced by microorganismsfor example disulfide reductasesand cysteine dioxygenase <sup>24</sup>. The main composition of keratin structure formed by disulfide bridges and Disulfide reductase and cysteine dioxygenase initiate keratin by breaking the cross-linkages between disulfide bonds which make the availability of peptide bonds for hydrolysis by keratinaseenzyme.

Ramnani et al., (2005) and Rahayu et al., (2012) combined the different enzymes for keratinolvsis of feather waste and keratinolysis concluded that by combinedenzymes were higher than single enzymes<sup>23,25</sup>. Microbial keratin degradation process is also dependent on someinternal factors like rate of agitation and sources of carbon, energy, and nitrogen. It is also suggested that keratin may be added as a sole source of carbon, nitrogen and energy for the efficient keratin decomposition. Availability of keratinase enzyme is also achieved by supplementing the media with keratin source which lead to the production of keratinase enzyme by the microorganisms. During keratin degradation process by microorganisms, exo-polysaccharides can also be synthesized which help in adhesion of microorganisms to the surface of keratin substrate enhancing more degradation of keratin<sup>26</sup>.

Application of mechanical force with enzymatic lysis has also been suggested for efficient keratin degradation<sup>27</sup>. It can be concluded that for efficientkeratin degradation, purified keratinase enzyme can be applied to keratin source.

Keratin and its structure:

It is important to understand the structure of keratin for degradation of keratin waste in both natural and in vitro conditions. Keratin is fibrous protein and found in the outer covering of almost all reptiles, vertebrates and fishes<sup>28</sup>. Keratin exists widely in nature and found as the third most abundant protein after cellulose and chitin<sup>29</sup>. Presence of high cross linking between disulfide bonds, make it a recalcitrant structural protein. Keratin can be divided into soft keratin and hard keratin. In Soft keratin cysteine content is less than 10% and found epidermis of skin and feathers.Hard keratin has cysteine content of10-14% and present in in hair, nails, wool, claws, and hooves<sup>30</sup>.

Keratin is insoluble in water and organic solvent because it forms recalcitrant polymers. Keratin is composed by some forces including disulfide bonds, hydrogen bonds and hydrophobic interactions. This recalcitrant polymer is resistant to degradation by enzymes such as pepsin and trypsin<sup>28</sup>.

On the basis of secondary structure, keratins are classified into  $\alpha$ -keratin and  $\beta$ keratin. The classification is based on the presence of  $\alpha$ -helix and  $\beta$ -sheet in the secondary structure of keratin. β-Keratin is composed of domains, rich in  $\beta$ -pleated sheets and formcrosslinking via disulfide bonds <sup>31</sup>. Beta-keratin is present in reptiles and birds<sup>32</sup>. The molecular weight of a keratin protein usually in the range of 10-14 kDa<sup>33</sup>. Beta ( $\beta$ ) keratin has high cysteine content with disulfide bonds which provide it properties of rigidityand resistance to degradation. Total β-keratin composition of a mature feather is about 80-90% of its content.

 $\alpha$ -Keratin consists of  $\alpha$ -helical-coil and can form intermediate filaments through selfassembling<sup>34</sup>. Alpha keratins are good in toughness, strength elasticity, and flexibility. These properties of Alpha ( $\alpha$ ) keratin is due to presence of some amino methionine, phenylalanine, acid like valine, isoleucine and alanine<sup>35</sup>. On the basis of sulfur content. Alpha keratin is further classified into hard and soft keratins<sup>36</sup>.

The  $\alpha$ -keratin has two keratin polypeptides in its structure, with head to tail structure, a dimeric coiled coil. in Thisselfassembled dimeric structure make head to tail polypeptides. The Dimers again make thier self-assembly to form the tetramers. Four such tetramer units assemble to make an intermediate filament<sup>34</sup>. This type of intermediate filament is found in skin and hair with similar pattern. The uncoiled head is rich in threonine and serine and then make a secondary structure due to phosphorylation and glycosylation activity. N-acetyl glucosamine is found as sugar moiety head domain of the polypeptide. Some molecular techniques can be adopted to increase the stability of keratin such as site specific phosphorylation and Posttranslational modifications including phosphorylation, sumoylation and glycosylation<sup>37</sup>.

Different organs may have varied content of  $\alpha$ -Keratin and  $\beta$ -keratin. For example, wool has  $\alpha$ -Keratinwhile feather is reported to have both keratins<sup>38</sup>. Feathers are found to have 41–67% content of $\alpha$ keratins and 33–38% content of  $\beta$ - keratin. Similarly, some other keratin substrate such as hair, bristle and woolhave 50–60%  $\alpha$ -keratins, some keratin-associated matrix proteins (20–30 %) and  $\beta$ -keratins. Not only organ but also part of organs may also have different content of keratin protein. The feather's outer rachis contains  $\beta$ keratin which make it more stable<sup>39</sup>.

Sources of keratin protein:

Keratin protein is present inliving organism and these living organisms are largest source for the isolation of keratin protein after their death. Various keratin sources are body parts of living organisms such as feathers, nails, wool, horns, hair, hoof, scales and stratum corneum<sup>40</sup>.

Feathers of birds are largest sources of keratin protein andare found mostly as waste products. Feathers are reported to contain 90% of keratin protein<sup>41</sup>. The chicken feathers also have about 90% of keratin making it very stable from water, heat, rain and cold. After slaughtering, feathers are not used and considered as biological waste for the environment<sup>42</sup>.

Hair is a good source of keratin and can be taken from human and animals. Animal hairs have more keratin on their hair. Keratin is responsible for the strength and flexibility of the hair.

The human hair contains approximate 80% of keratin protein in its composition. The accumulation of hair after hair cut and animal death causes unusual accumulation of hair, which are counted in the environmental waste.

Human nail is composed of a highlycrosslinked keratin network, with several disulfide linkages and high sulfur content (3.8%). These structural features make human nails a highly strong keratin substrate. Similar conformation with more sulfur content and crosslinking, is found in animal horns making it very tough organ. amino Some free acids. calcium aluminium, copper, iron, zinc, manganese, zinc and chromium also provide toughness to the animal horn<sup>43</sup>. The hoof of animals has high thermal stability due to presence of both  $\alpha$ -helical conformation and  $\beta$ sheet<sup>44</sup>. The extremal shell of hard keratin is found in beak of birds which make it very stable. So it is clear that different organs are good source of keratin and can be isolated from these body parts. Keratinases:

The term keratinase is used for important proteases which possess keratinolytic activities. These keratinase proteases show proteolytic activity against stable proteinkeratin. Keratinase contain serine or metal protease activity for keratin degradation. Keratinase complete their keratinolytic activity with combined action of some other proteolytic enzymes<sup>16</sup>.

Keratinase are enzymes which are synthesized by microorganisms when keratin substrate is available in the media. The keratinases produced by various types of bacteria and fungi of different sources differ in terms of amino acid sequence, molecular weight, optimum pH and temperature<sup>45,46</sup>. Due to keratinolytic activity of keratinase for degradation offeathers, hair and wool, keratinase are really important for industrial applications<sup>47,48</sup>.

Keratinase have important role in different industrial and agrircultural purposes including fertilizers, leather industries, biomedical fields, detergents, cosmetics and material production of Nitrogen fertilizer, biofertilizer, textile industry, pharmaceutical industry and Nanobiotechnology <sup>49</sup>. Keratinase are also being used in poultry industries for improved meat production in broiler chicken<sup>50</sup>.

Several researchers have isolated and purified keratinase from microorganisms and characterized the purified keratinase. Keratinase enzyme is produced bv supplementing the media with keratin substrates like keratin azure, azokeratin, human hair, cow horn, feather and keratin Microorganisms powder. produce keratinase enzymes when optimum conditions are present including pH, temperature and buffer. Bacteria and fungi produce keratin when suitable temperature range is present ranging from  $28-40^{\circ}$ C and mostly between 28-35<sup>o</sup>C. Similarly, these microorganisms show their higher efficiency of keratinase production at various pH ranging from 5 to  $13^{51}$ . Different enzymes produced by fungi, bacteria and other extremophiles combine with keratinase which are capable of keratindegradation<sup>52</sup>. efficient highly Biochemical and molecular studies of proteases indicate that enzyme engineering can be adopted for the development of disulfide boding between the keratinase for higher stability.

The mode of action of keratinase for keratin degradation is started with keratinase attachment to keratin. Keratin degradation by Keratinases is not site specific and keratinase attack peptide bonds of keratin at numeroussites. Keratinase can also cleave the bond between aromatic and nonpolar amino acids.

The most keratinolytic group belongs to fungi deuteromycetese including the species of Acremonium, Aspergillus, Chrysosporium, spergillus, Alternaria. Trichurus, Ctenomyces, Curvularia, Cladosporium, Geomyces, Fusarium, Geotrichum. Gleomastis, Monodictys, Mvrothecium. Paecilomvces.

Pacecilomyces, Penicillium, Stachybotrys, Urocladium, Scopulariopsis, Sepedonium, Trichurus, Doratomyces. Keratinase produced by these fungi has demonstrated effective cleavage of keratin substrate. More than 200 species have been used for the isolation of fungal keratinase. Fungal keratinase is very promising for the degradation of keratin and utilize it efficiently for degradation<sup>53</sup>.

Dermatophytic species also have been reported to produce keratinase enzyme including *Microsporum canis*, *M. cookie*, *M. persicolor*, *Trichophyton krajdenii*, *T. mentagrophytes*, *T. raubitschekii*, *T. rubrum*, *T. Simii*and*Epidermophyton floccosum*.These reviews indicate that keratinolytic species are good sources of keratinase enzyme.<sup>54</sup>

## **Conclusions:**

Keratinolyticmicroorganisms degrades the various keratinous waste in an efficient way. Degradation of keratinous waste materials by microorganisms reduces environmental problems in an eco friendly way. Biological degradation of keratinous waste is a simple and cheap method and dependent on some specific enzyme like keratinase. Keratinase enzyme along with some proteins, peptides and amino acids degrade keratin substrates efficiently. Keratinase from microorganisms can be used for recycling of poultry waste, animal feed and in leather industry.

# References

- Lateef A, Oloke JK, Gueguim-Kana EB, Sobowale BO, Ajao SO and Bello BY 2010, Keratinolytic activities of a new feather-degrading isolate of *Bacillus cereus* LAU 08 isolated from Nigeriansoil. *Int. Biodeterior. Biodegr.* 64 162–165.
- Gareth LJ, Julian P, Callum T and Erik B 2010, Protease- and keratinaseproducing microbial strainsfor compost

bioaugmentation. *Int. Biodeterior. Biodegr.* **64** 574–580.

- 3. Huang Y, Sun Y, Ma S, Chen L, Zhang H and Deng Y 2013, Isolation and characterization of *Keratinibaculum paraultunense* gen. nov., sp. nov., a novel thermophilic, anaerobic bacterium withkeratinolytic activity. *FEMS Microbiol. Lett.* **345** 56–63.
- 4. Paul T, Das A, Mandal A, Halder SK, DasMohapatra PK, Pati Brand and Mondal KC 2014, Valorization of chicken feather waste for concomitant production of keratinase, oligopeptides and essential amino acids under submerged fermentation by *Paenibacillus woosongensis* TKB2. *Waste Biomass Valoriz.* **5** 575–584.
- Lin X, Inglis GD, Yanke LJ and Cheng KJ 1999, Selection and characterization of feather degrading bacteria from canola meal compost. *J.Ind. Microbiol. Biotechnol.* 23 149– 153.
- Kim JM, Lim WJ and Suh HJ 2001, Feather-degrading *Bacillus* species from poultry waste. *Process Biochem.* 37 287–291.
- Werlang PO and Brandelli A 2005, Characterization of a novel featherdegrading *Bacillus* sp. strain. *Appl. Biochem. Biotechnol.* 120 71–79.
- Lateef A, Adelere IA and Gueguim-Kana EB 2015, *Bacillus safensis* LAU 13: a new source of keratinase and its multi-functional biocatalytic applications. *Biotechnol. Biotechnol. Equip.* 29 54–63.
- Ahmadpour F, Yakhchali B and Musavi MS 2017, Isolation and identification of a keratinolytic *Bacillus cereus* and optimization of keratinase production. J. Appl. Biotechnol. Report. 3 507–512.

- Blyskal B 2009, Fungi utilizing keratinous substrates. *Int. Biodeterior. Biodegr.* 63(6) 631–653.
- 11. Eliades L, Cabello M, Voget C, Galarza B and Saparrat M 2010, Screening for alkaline eratinolytic activity in fungi isolated from soils of the biosphere reserve "Parque Costero del Sur" Argentina. World J. *Microbiol. Biotechnol.* **26** 2105–2111.
- Liang JD, Han YF, Zhang JW, Du W, Liang ZQ and Li ZZ 2011, Optimal culture conditions for keratinase production by a novel thermophilic *Myceliophthora* strain GZUIFR-H4.9-1. J. Appl. Microbiol. 110 871–880.
- Lange L, Busk PK and Huang Y 2014, Use of a microbial composition for the degradation of keratinaceousmaterials. Patent No. WO2014169920.
- Gradisar H, Kern S and Friedrich J 2000, Keratinase of Doratomyces microsporus. Appl. Microbiol. Biotechnol. 53 196–200.
- 15. Gilardi GL 1965, Nutrition of systemic and subcutaneous pathogenic fungi. *Bacteriol. Rev.* **29** 406–424.
- 16. Gupta R and Ramnani P 2006, Microbial keratinases and their prospective applications: an overview. *Appl. Microbiol. Biotechnol.* **70** 21-37.
- 17. Habbeche A, Saoudi B, Jaouadi B, Haberra S, Kerouaz B, Boudelaa M, Badis A and Ladjama A 2013, Purification and biochemical characterization of a detergent-stable keratinase from a newlv thermophilicactinomycete Actinomadura keratinilytica strain Cpt29 isolated from poultry compost. J. Biosci. Bioeng. 117 413-421.
- 18. Manivasagan P, Sivakumar K, Gnanam S, Venkatesan J and Kim SK 2014, Production, iochemicalcharacterization and detergents application of keratinase from the marine

Actinobacterium Actinoalloteichus sp. MA-32. Journal of Surfactants and Detergents. **17** 669–682.

- 19. Syed DG, Lee JC, Li WJ, Kim CJ and Agasar D 2009, Production, characterization and application of keratinase from *Streptomyces gulbargensis*. *Bioresour Technol*. **100** 1868–1871
- 20. Saha S, Dhanasekaran D, Shanmugapriya S and Latha S 2013, *Nocardiopsis* sp. SD5: a potent feather degrading rare actinobacterium isolated from feather waste in TamilNadu, India. J. Basic Microbiol. 53 608–616.
- 21. Elhoul MB, Jaouadi NZ, Rekik H, Benmrad MO, Mechri S, Moujehed E, Kourdali S, El Hattab M, Badis A, Bejar S and Jaouadi В 2016. Biochemical and molecular characterization of new keratinovtic protease from Actinomadura viridilutea DZ50. Int. J. Biol Macromol. 92 299-315.
- 22. Ningthoujam DS, Devi LJ, Devi PJ, Kshetri P and Tamreihao K 2016, Optimization of keratinase productionby *Amycolatopsissp.* Strain MBRL 40 from a limestone habitat. *Journal of Bioprocess Biotech.* **6**(5) 6.
- Ramnani P, Singh R and Gupta R 2005, Keratinolytic potential of *Bacillus licheniformis* RG1: structural and biochemical mechanism of feather degradation. *Can. J. Microbiol.* 51 191–196.
- 24. Yamamura S, MoritaY, Hasan Q, Yokoyama K and Tamiya E 2002, Keratin degradation: a cooperative action of two enzymes from *Stenotrophomonas* sp. *Biochem. Biophys. Res. Comm.* **294** 1138–1143.
- 25. Rahayu S, Syah D and Suhartono MT 2012, Degradation of keratin by keratinase and disulfide reductase from

*Bacillus* sp. MTS of Indonesian origin. *Biocatal. Agric. Biotechno.* **1** 152–158.

- 26. Vasileva-Tonkova E, Gousterova A and Neshev G 2009, Ecologically safe method for improved featherwastes biodegradation. *Int. Biodeterior. Biodegr.* 63 1008–1012.
- 27. Kunert J 2000, Physiology of keratinophilic fungi. *Rev. Iberoamericana de Micologia* 1 77–85.
- Meyers MA, Chen P, Lin AY and Seki Y 2008, Biological materials: structure and mechanical properties. *Prog. Mater. Sci.* 53 1–206.
- 29. Lange L, Huang Y, Busk PK 2016, Microbial decomposition of keratin in nature—a newhypothesis of industrial relevance. *Appl Microbiol Biotechnol* **100** 2083–2096.
- 30. Jin H-S, Park SY, Kim K, Lee Y-J, Nam G-W and Kang NJ 2017, Development of a keratinase activity assay using recombinant chicken feather keratin substrates. *PloS one.* **12**(2) p.e0172712
- 31. Fraser RDB and Parry DA 2011, The structural basis of the filament-matrix texture in the avian/reptilian group of hard  $\beta$ -keratins. *Journal of Structural Biology.* **173** 391–405.
- 32. Greenwold MJ, Sawyer RH 2013, Molecular evolution and expression of archosaurian β-keratins: Diversification and expansion of archosaurian  $\beta$ -keratins and the origin of feather β-keratins.  $J_{\cdot}$ Exp. Zoology.Part *B*: Molecular and Developmental Evolution. 320(6) 393-405.
- Woodin AM 1956, Structure and composition of soluble feather keratin. *Biochemical Journal*. 63(4) 576-581.
- 34. McKittrick J, Chen PY, Bodde SG, YangW, Novitskaya EE and Meyers MA 2012, The structure, functions, and

mechanical properties of keratin. *JOM* **64** 449–468.

- 35. Schowen 1993. Principles of biochemistry 2nd ed. (Lehninger, Albert L.; Nelson, David L.; Cox, Michael M.). Journal of Chemical Education. 70 (8).
- 36. Nickerson WJ 1947, Biology of Pathogenic Fungi. Medical mycology. Annual Reviews in Microbiology, 7(1) 245-272.
- 37. Snider NT and Omary MB 2014, Posttranslational modifications of intermediate filament proteins: mechanisms and functions. *Nat.Rev.Mol.Cell Biol.* 15 163.
- 38. Bodde SG, Meyers MA and McKittrick J 2011, Correlation of the mechanical and structural properties of cortical rachis keratin of rectrices of the Toco Toucan (Ramphastos toco). J. Mech. Behav. Biomed. Mater. 4 723– 732.
- 39. Tesfaye T, Sithole B andRamjugernath D 2017, Valorisation of chicken feathers: a review on recycling and recovery route—current status and future prospects. *Clean Technol. Environ Policy.* 19 2363–2378.
- 40. Kim JD 2007, Purification and characterization of a keratinase from a feather degrading fungus, *Aspergillus flavus* strain K-03. *Mycobiology*. **35** 219–225.
- 41. Nagal S and Jain PC2010, Feather degradation by strains of *Bacillus* isolated from decomposing feathers. *Brazilian J. Microbiol.* **41**(1) 196-200.
- 42. Chaturvedi V and Verma P 2014, Metabolism of chicken feathers and concomitant electricity generationby *Pseudomonas aeruginosa* by employing microbial fuel cell (MFC). *Journal of Waste Management*. 1-9.
- 43. Romanov OE 2005, The comparative analysis of horn properties and their

composition. *Scientific Research of Caucasus* 91-95.

- 44. Kakkar P, Madhan B and Shanmugam G 2014, Extraction and characterization of keratin frombovine hoof: A potential material for biomedical applications. *Springer Plus*. 3(1) 596.
- 45. Brandelli A 2008, Bacterial Keratinases: Useful enzymes for bioprocessing agroindustrial waste and beyond. *Food and Bioprocess Technol.* 1(2) 105-116.
- 46. Kaewsalud Τ, Yakul K, Jantanasakulwong K, Watanabe M. Tapingkae W and Chaiyaso 2021, Biochemical Characterization and Application of Thermostable-Alkaline Keratinase from *Bacillus* halodurans SW-X to Valorize Chicken Feather Wastes. *Waste* **Biomass** Valor 12 3951–3964.
- 47. Yahaya RS, Phang LY, Normi YM, Abdullah JO, Ahmad SA and Sabri S 2022, Feather-Degrading Bacillus cereus HD1: Genomic Analysis and Its Optimization for Keratinase Production and Feather Degradation. *Current Microbiology*. **79**(6) 1-5.
- 48. Peng Z, Mao X, Mu W, Du G, Chen J, and Zhang J 2022, Modifying the Substrate Specificity of Keratinase for Industrial Dehairing to Replace Lime-Sulfide. ACS Sustainable Chemistry & Engineering 10(20) 6863-6870.
- 49. Paul T, Jana A, Mandal, AK, Mandal A, Das Mohpatra PK and Mondal KC 2016, Bacterial keratinolytic protease, imminent starter for NextGen leather and detergent industries. *Sustain. Chem. Pharm.* **3** 8–22.
- 50. Xu KL, Gong GX, Liu M, Yang L, Xu ZJ, Gao S, Xiao MY, Ren T, Zhao BJ, Khalil MM and Zhao L 2022, Keratinase improves the growth performance, meat quality and redox

status of broiler chickens fed a diet containing feather meal. *Poultry Science* **101**(6) 101913.

- Brandelli A, Salab L and Kalil SJ 2015. Microbial enzymes for bioconversion of poultry waste into added-value products. *Food Res. Int.* 7(3) 3-12.
- 52. Kanoksilapatham W and Intagun W 2017, A review: biodegradation and applications of keratin degrading microorganisms and keratinolytic enzymes, focusing on thermophiles and thermostable serine proteases. *Am. J. Appl. Sci.***14** 1016–1023.
- 53. Daroit DJ, Brandelli A 2014, A current assessment on the production of bacterial keratinases. *Crit. Rev. Biotechnol.*34 372–384.
- 54. Khan AM and Bhadauria S 2015, A review on chemical and molecular characterization of keratinophilic fungi. *Int. J. Sci. Res.* **4**(1) 420-423.