

Keratinolytic Activity of *Arthroderma* and *Chaetomium* Species Isolated From Ujjain Soils

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Annotation: Twelve species of keratin decomposers isolated from 923 samples of feathers, nests and pellets of 90 species of free-living birds were clustered in 4 "econs" (numerical ecological groups): (A) *Arthroderma tuberculatum*, *A. ciferrii*, *A. multifidum*, *A. cuniculi* (associated with the birds nesting in hollows); (B) *Arthroderma quadrifidum*, (moderately alkalophilic species associated with the birds having a frequent contact with the soil); (C) *Aphanoascus fulvescens* (D) *Arthroderma curreyi*, (the species common in water habitats and associated frequently with exoanthropic birds) also isolated from Ujjain soils. *Chaetomium* is a genus of fungi in the Chaetomiaceae family. It is a dematiaceous (dark-walled) mold normally found in soil, air, cellulose and plant debris. According to the Dictionary of the Fungi (10th edition, 2008), there are about 95 species in the widespread genus. Members of this genus typically have superficial, ostiolar perithecia, covered in hairs. Asci are often clavate and evanescent, bearing eight spores. Ascospores are usually lemon-shaped, commonly colored olive-brown. Mycelia often grows in conglomerate masses that resemble ropes. *Chaetomium* spp. are also encountered as causative agents of infections in humans. Many cases cause type 1 allergic reactions and infections. A few cases of fatal deep infections due to *Chaetomium atrobrunneum* have been reported in immunocompromised people. Other clinical syndromes include brain abscess, peritonitis, and onychomycosis. these have also been isolated from Ujjain soils.

Keywords: *Arthroderma*, *Chaetomium*, fungi, soils, Ujjain, habitat, keratinolytic, infections, human.

Introduction

The hydrolysis of keratin wastes by *Arthroderma* & *Chaetomium* is considered a biotechnological alternative for recycling and valorization through keratinolytic fungi. Despite their resistant structure, keratin wastes can be efficiently degraded by various microorganisms through the secretion of keratinases, which are promising enzymes for several applications, including detergents, fertilizers, and leather and textile industry. In an attempt to isolate keratinolytic microorganisms that can reach commercial exploitation as keratinase producers, the current work assesses the dynamics of keratin biodegradation by several keratinolytic fungal strains isolated from Ujjain soil.[1,2]

Keratin is an insoluble fibrous protein characterized by high stability due to the high degree of cross-linkages by disulfide and hydrogen bonds. It also contains a variety of amino acids, predominantly cystine, lysine, proline, and serine. Considering the secondary structural conformation, keratins have been classified into α - (α -helix of hair and wool) and β -keratins (β -sheets of feather). In addition, keratins are grouped into hard keratin (hair, feather, nails, wool, etc.) having a high disulfide bond content and soft keratin (skin) with a low disulfide bond content. Keratin-rich wastes are troublesome environmental contaminants and are released in increasing quantities as byproducts from agro-industrial processes in the form of feathers, hair, nails, and horns. Fungi *Arthroderma* & *Chaetomium* play an ecological role in the degradation of keratin substrates through their contribution to recycling the carbon, nitrogen, and sulfur from keratins isolated from Ujjain soils.[3,4] Considering from both economic and environmental point of view, attention was focused on the management of recalcitrant keratinous wastes. The hydrolysis of keratin wastes by microorganisms is considered a biotechnological alternative for recycling and

valorization through keratinolytic microorganisms. Keratinases, considered as proteases with keratinolytic function, act synergistically with other keratinolytic enzymes to degrade the complex supramolecular organization of keratin. After disrupting the disulfide bonds of cysteine, the major amino acid in keratin, the keratin substrate is more easily available to the hydrolytic enzymes secreted by *Arthroderma* & *Chaetomium* fungi. [5, 6]

Despite the resistant structure, keratin wastes can be efficiently degraded by various microorganisms that secrete of keratinolytic enzymes, such as keratinases, which are a group of serine or metalloproteases. These enzymes are predominantly extracellular and are produced by *Arthroderma* & *Chaetomium* isolated from Ujjain soils. It is believed that in the future, fungal keratinases will occupy a special niche among proteases as valuable enzymes for the bioprocessing of the keratinous wastes, which are released into the environment in huge amounts because of human activities. [7,8]

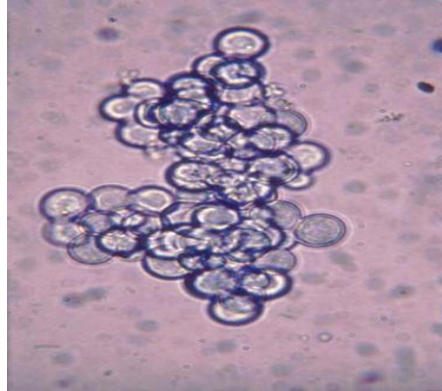
During the biodegradation process, only the S–S bonds were affected by *Arthroderma* & *Chaetomium*, and after long contact, the microbial enzymes secreted disrupt the peptide bonds of the keratin chains. Because keratinolytic fungi and their enzymes keratinases are a subject of scientific and economic interest because of their capability to hydrolyze keratin *Arthroderma* & *Chaetomium* isolated from Ujjain soils has potential in the biotechnological process of biodegrading keratin. [9,10]

Discussion

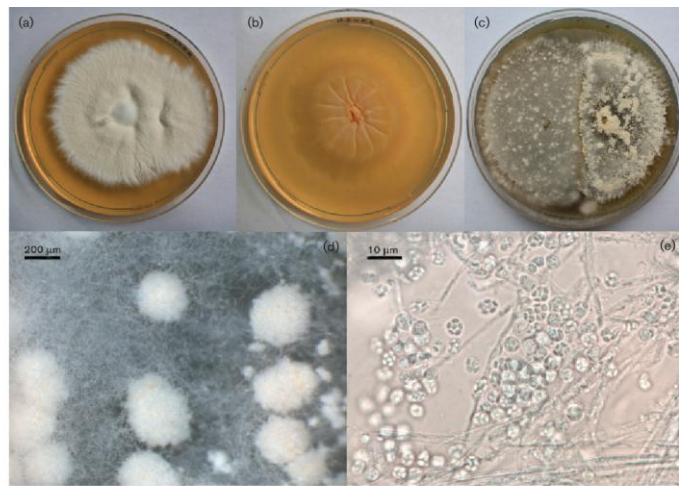
In this descriptive study, 3 soil samples were collected from various sites of 2 different parks in Ujjain were isolated during 2022. The samples were collected from the superficial layer of the soil whose depth did not exceed 5–10 cm by using an iron spatula. In doing so, 300–400 gram of soil was collected in sterile polyethylene bags and brought to laboratory for further processing. PH of each soil sample was measured after preparation of soil suspension (one gram of soil to five mL deionized water) using pH meter. [29] We used Vanbreuseghem's hair bait technique for isolation of keratinophilic fungi. Briefly, each soil sample was thoroughly mixed, and about 70 gram of the soil was packed in a sterile 90 mm Petri dish. Then, several pieces of sterile healthy children hair fragments were dispersed over the surface of the soil samples and moistened with sterile distilled water supplemented with antibiotic solutions, chloramphenicol (0.2 g/mL), and strepto-penicillin (1000 IU/mL). [11,12] All the baited soil Petri dishes were incubated at room temperature (20–25°C) in the dark for 2 months and got moistened if necessary. After observation of colony growth around the hairs, the colonies were subcultured on Sabouraud's dextrose agar (SDA) with and without chloramphenicol (50 mg/L) and cycloheximide (500 mg/L) and purified. The fungi were identified based on the conventional method (colony morphology and macro- and microconidia characteristics) and DNA-based identification techniques. [27,28] Molecular identification of the unknown isolates was achieved by DNA sequence analysis. First, the fungi were grown in flasks containing Sabouraud's dextrose broth and incubated at 25°C for several days using shaker rotator. After colony growth, the culture was filtered. First, the frozen mycelium mass was smashed by mechanical pressure using sterile poulder and liquid nitrogen. The acquired powder was then mixed with lyses buffer and the DNA was extracted. [13,14]

The fungal isolates belonged to 2 genera as follows: *Arthroderma* & *Chaetomium*

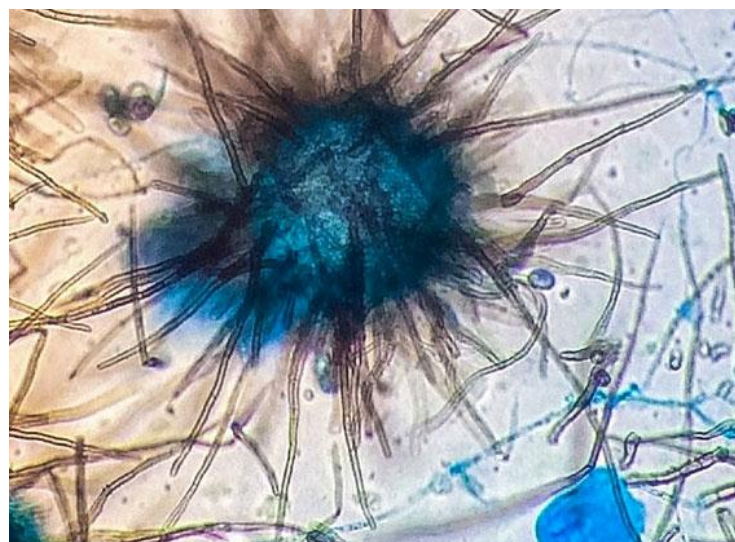
Results



Arthroderma



Arthroderma (Colonies and fungal morphology)



Chaetomium (fruiting body releasing spores)

The keratinolytic activity of keratinophilic fungi is important for ecology [25,26] and has attracted many researchers' attention around the world. Keratinophilic fungi play an important role in the natural degradation of keratinized residues in the soil of Ujjain. Parks are among the popular public places for the people to spend their time and have fun with their family. These parks potentially have a high risk for transmission of fungal infections from *Arthroderma* & *Chaetomium*. [15,16] These keratinophilic fungi responsible for the majority of fungal skin infections in humans. Infections are primarily localized to the hair, nails, and outer layers of the skin, reflecting the affinity of the fungi for keratin and the inability of the fungi to invade deeper tissues or organs in immunocompetent skin. Infections can cause a number of symptoms, including extensive inflammation, itching, and discolored, scaly skin lesions. These infections, referred to as dermatophytoses. [17,18] Dermatophytes express cell-surface associated and secreted proteins that may contribute to virulence.



Sporotrichosis by fungal spores of *Arthroderma* from Ujjain soils

Coincubation of peripheral blood mononuclear cells (PBMCs) with mannan, a glycoprotein found in the fungal cell wall, [23,24] inhibits lymphocyte proliferation in response to mitogens and antigenic stimuli. Additionally, these fungi express a number of proteases, including elastase and keratinases, which likely contribute to the spread of the fungi within the skin by degrading substrates in the extracellular matrix.

Treatment options for dermatophytoses depend on the location and extent of the infection. As such, topical antifungals are used to treat the majority of dermatophyte infections, although oral therapy is preferable.[19,20]

Dermatophyte spores survive more than a year in the environment under optimal conditions of temperature and humidity, and they resist most routinely used hospital disinfectants, which facilitates transmission. Dermatophytosis is an infection of hair shafts and stratum corneum caused by keratinophilic fungi. Skin involvement may be localized, multifocal, or generalized. Pruritus, if present, is usually minimal to mild but occasionally may be intense. Lesions usually include areas of circular, irregular, or diffuse alopecia with variable scaling. Remaining hairs may appear stubbled or broken off.[21,22]



Onychomycosis caused by *Chaetomium* in Ujjain soil

Conclusions

A number of soil fungi are known as potential pathogens for humans and animals isolated from Ujjain soils i.e. *Artroderma* & *Chaetomium*. Keratinophilic fungi are among important groups of fungi living in the soil that colonize in various keratinous substrates, produce keratinases, and decompose them into components with lower molecular weight. [30]

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