

Fungal Biology

Amritesh Chandra Shukla *Editor*

Applied Mycology

Entrepreneurship with Fungi

 Springer

Fungal Biology

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Fungal biology has an integral role to play in the development of the biotechnology and biomedical sectors. It has become a subject of increasing importance as new fungi and their associated biomolecules are identified. The interaction between fungi and their environment is central to many natural processes that occur in the biosphere. The hosts and habitats of these eukaryotic microorganisms are very diverse; fungi are present in every ecosystem on Earth. The fungal kingdom is equally diverse, consisting of seven different known phyla. Yet detailed knowledge is limited to relatively few species. The relationship between fungi and humans has been characterized by the juxtaposed viewpoints of fungi as infectious agents of much dread and their exploitation as highly versatile systems for a range of economically important biotechnological applications. Understanding the biology of different fungi in diverse ecosystems as well as their interactions with living and non-living is essential to underpin effective and innovative technological developments. This series will provide a detailed compendium of methods and information used to investigate different aspects of mycology, including fungal biology and biochemistry, genetics, phylogenetics, genomics, proteomics, molecular enzymology, and biotechnological applications in a manner that reflects the many recent developments of relevance to researchers and scientists investigating the Kingdom Fungi. Rapid screening techniques based on screening specific regions in the DNA of fungi have been used in species comparison and identification, and are now being extended across fungal phyla. The majorities of fungi are multicellular eukaryotic systems and therefore may be excellent model systems by which to answer fundamental biological questions. A greater understanding of the cell biology of these versatile eukaryotes will underpin efforts to engineer certain fungal species to provide novel cell factories for production of proteins for pharmaceutical applications. Renewed interest in all aspects of the biology and biotechnology of fungi may also enable the development of “one pot” microbial cell factories to meet consumer energy needs in the 21st century. To realize this potential and to truly understand the diversity and biology of these eukaryotes, continued development of scientific tools and techniques is essential. As a professional reference, this series will be very helpful to all people who work with fungi and should be useful both to academic institutions and research teams, as well as to teachers, and graduate and postgraduate students with its information on the continuous developments in fungal biology with the publication of each volume.

More information about this series at <http://link.springer.com/series/11224>

Amritesh Chandra Shukla
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Applied Mycology

Entrepreneurship with Fungi

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This volume is dedicated to my daughters, Baby Aaradhya (08 yrs) and Baby Arunnima (05 yrs), who sacrificed their sport time and helped me during compilation of this work.



Foreword

Fungi is a group of organisms that are an important link in the food chain in all ecosystems with immense potential and myriad of useful bioactive compounds. Fungi feature in a wide range of diverse processes and applications in the modern agriculture, food, and pharmaceutical industry. In the food and drink arena as well as in fermented foods, the role of fungi has been important historically in products such as mushrooms, and yeasts for baking and brewing, respectively. These roles are supplemented by the use of fungal food processing enzymes and additives, and more recently the development of protein-based foodstuffs from fungi. Further, certain mycorrhizal fungi (VAM & AM) may be necessary for seed germination and plant health or may be used as biocontrol agents against weeds and invertebrates; formulations of biofertilizer cum biopesticide can be used as biostimulants and bioprotectants of crops. The successful application of biotechnological processes in agriculture and food using fungi may therefore require the integration of number of scientific disciplines and technologies. These may include subjects as diverse as agriculture, biotechnology, biochemistry, molecular biology, and process engineering. The practical use of newer techniques such as genetic recombination and robotics has revolutionized the modern agricultural biotechnology industry and has created an enormous range of possible applications of fungal products in developing entrepreneurship.

Applied Mycology: Entrepreneurship with Fungi, published by Springer Nature, is a momentous volume, and I strongly deem that it will attract readers working in the field of microbiology, especially in mycology. The current volume has 20 chapters contributed by academicians, scientists, and researchers working in the field of mycology throughout the world. I congratulate the editor for bringing out this

volume with amazing commitments from professionals working in the area of fungi and their application for developing entrepreneurship in various sectors including agriculture, food, health, nutraceuticals, pharmaceuticals, and other allied sectors.



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Arvind Deshmukh

Preface

Microorganisms are a part and parcel of our life. They are our friends and can also be our foe. They help in maintaining a balance in our ecosystem. We have tapped their potential for the benefit of society. Hence, the objective of this volume, *Applied Mycology: Entrepreneurship with Fungi*, is to keep the readers informed about the recent developments that took place in the applied mycology and their possible application for developing entrepreneurship.

The volume consists of 20 chapters contributed by authors with vast experience in the field of applied mycology. The main focus of this book is to popularize various methods and protocols related to formulation of different value-added fungal products and their application for developing entrepreneurship. The book chapters cover wide applications of fungi in nutraceutical and baking purposes; fungi in cosmetics; fungi in industrial biotechnology/mycotechnology (such as production of alcohols, organic acids, enzymes, vitamins, polysaccharides, pigments, lipids, glycolipids) and brewing industry; fungi in pharmaceuticals and production of antibiotics; mycometabolites from endophytic fungi having potential in pharmaceutical and agricultural industries; fungi in bioremediation/biodegradation; fungal consortium for organic municipal solid waste composting; fungal biopesticides, fungal biofertilizer, plant growth promoter, mycotoxins, and mycorrhizal symbioses; as well as fungi in micro-based bio-refinery for gold nanoparticles production.

Further, advance researches and their applications, as given in the book, will give more ideas to researchers, scientists, policy makers, and professionals working in the area of applied and industrial mycology. Besides, the information will not only be helpful for developing new startup/fungal entrepreneurship and revenue generation for the welfare of society but also be helpful to reduce un-employability at the global level.

Lucknow, India

Amritesh Chandra Shukla

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I am highly grateful and thankful to the authorities of the University of Lucknow, especially Professor Alok Kumar Rai, Hon'ble Vice-Chancellor, University of Lucknow, for his endeavor and motivations at all stages of the progress. I am also thankful to my colleagues and research scholars in the Department of Botany, University of Lucknow, for providing their valuable help and support during compilation of this book.

Further, I don't have words to thank my parents, Sri Kailash Chandra Shukla and late Smt. Sandhya Shukla, the makers of my destiny who with all difficulties sent me away from home to get this highly specialized knowledge of 'Science'; without their well wishes, this work would have never been completed.

My acknowledgments are incomplete if I fail to remember the constant encouragement and great sacrifice of my wife Dr. Neetu Shukla, and my daughters (baby Aaradhya Shukla and baby Arunnima Shukla); without her understanding forbearance and help, this work would have been impossible.

I also express my deepest gratitude to my family members, friends, and well-wishers, who rendered their support during this work. Finally, with folded hands and bent knees, I bow before the almighty to thank Him for the continuing graces and blessings He showers on me as I tread through the scientific path of service for humanity.

Lucknow, India

Amritesh Chandra Shukla

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Chapter 1

Fungal Cosmetics: Mushrooms in Beauty Care and the New Age of Natural Cosmetics



Sridevi Visvanathan, Ravishankar Krishnamoorthy,
and Gokul Shankar Sabesan

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1.1 Introduction

Fungi are a group of eukaryotic organisms existing in the ecosystem as chemoheterotrophs, predominantly as saprophytes on dead and decaying organic matter. Mushroom is a term usually used to describe the conspicuous ‘umbrella-like’ shaped fruiting bodies or sporophores of certain fungi. Mushrooms typically are members of the order Agaricales and phylum Basidiomycota (produces basidia or cup-shaped fruiting bodies). Mushrooms are also referred to as ‘gill fungi’, because they have a characteristic gill-like appearance on the underside of the cap. In general, mushrooms are popular around the globe due to their edible nature and it forms part of the recipe for a variety of delicious food because of the taste and meat-like texture. But it should be remembered that all mushrooms are not edible and there are certain toxic varieties because of the secondary metabolites produced by them. However, the deadly mushroom groups are limited and the toxicity displayed by these species is an inborn mechanism of protection of the basidiocarp from consumption and premature destruction by animals. Nevertheless, the versatility of the edible mushrooms is much and beyond; they are widely exploited for their medicinal values and cosmeceutical properties. Numerous mushrooms are used as functional ‘do-good’ ingredients in cosmetic formulations. The surging demand for mushroom or fungal cosmetics is on a rise owing to their inherent properties that confer natural beauty benefits in the cosmetics and personal care formulations. The nutrient-rich extracts of various mushrooms can serve as great skin and hair care ingredients in enhancing the quality of beauty care products.

Most of the time, the mushrooms are more easily recognized by their common names than the botanical names. The common names and scientific names of mushrooms are listed in Table 1.1.

The incredible functional benefits of the mushrooms are multifold. Mushrooms are powerhouses of natural bioactive compounds and have been used traditionally

Table 1.1 Popular mushrooms/fungi and their scientific names

| S. no. | Common name | Scientific name |
|--------|--|---|
| 1 | Snow mushroom or white jelly mushroom or silver ear fungus | <i>Tremella fuciformis</i> |
| 2 | Reishi mushroom or Lingzhi | <i>Ganoderma lucidum</i> |
| 3 | Caterpillar fungus | <i>Ophiocordyceps sinensis</i> |
| 4 | Oyster mushroom | <i>Pleurotostreatus</i> |
| 5 | Button mushroom | <i>Agaricus bisporus</i> |
| 6 | Turkey tail mushroom | <i>Trametes versicolor/Coriolus versicolor and Polyporus versicolor</i> |
| 7 | Shiitake mushroom | <i>Lentinula edodes</i> |
| 8 | Chaga mushroom | <i>Inonotus obliquus</i> |
| 9 | Penny bun or Porcini mushroom | <i>Boletus edulis</i> |

used for centuries in alternative medicine and beautifying decoctions for skin and hair care benefits (Taofiq et al., 2016).

1.2 Application in Skincare

1.2.1 Anti-microbial Properties

The popular saying goes as ‘beauty is skin deep’, but the beauty of the skin can be spoiled by a variety of skin infections that are caused by bacteria and fungi. The extracts of mushrooms contain bioactive components that exhibit inhibitory action of several bacteria and fungi that cause skin conditions and therefore find its place in the cosmetic industry.

1.2.1.1 Anti-dermatophytic and Skin Protective

Dermatophytes are a group of keratinophilic fungi which cause superficial mycoses on human beings and animals. The disease caused by dermatophytes is called dermatophytosis (pl dermatophytoses), which affects the superficial keratinized layers of the body, namely, the skin, hair and the nail. Dermatophytes are pathogenic fungi with the ability to colonize the skin and degrade the keratinous layers by virtue of producing proteolytic enzymes such as keratinases and collagenases. Dermatophyte infections are most common among the skin infections and can be easily acquired indirectly by fomite transmission from soil, by sharing of bed linen, clothing and footwears and directly from infected pets or domestic animals. Hot and humid climate, coupled with a habit of tight-fitted clothes like jeans, jeggings, leggings, etc., can further facilitate ward of dermatophyte infection. The exorbitant cost of the conventional antifungal drugs coupled with the rampant development of anti-fungal resistance by the skin fungi justifies the need to look for alternative sources of remedies. The use of mushrooms extracts that have anti-dermatophytic activity in skin-care formulations would indirectly help to prevent such infections. Extracts of *Lenzitesquercina*, *Ganoderma lucidum* and *Rigidoporusulmarius* exhibited a wide range of anti-dermatophytic activity against dermatophytes similar to the commonly used antifungal agents (Ogidi & Oyetayo, 2016). This activity may be attributed to the presence of several compounds such as alkaloids, phenols, steroids, flavonoids, tannins, cardiac glycosides and terpenoids. Further, Wong et al. (2010) had substantiated the anti-fungal activity of certain mushrooms to the activity of glucanase. Glucanase is an enzyme exhibiting anti-fungal and anti-dermatophytic activities. It acts on glycan in the fungal cell wall and thereby weakens the wall by hydrolysing the glycan of the pathogenic fungi.

Further, the fungal extracts of these mushrooms are documented to have a medicinal effect on certain skin conditions such as eczema, rosacea and psoriasis. The

extract of chaga mushroom has betulinic acid which has a beneficial role in several skin conditions. Few commercially available medicinal products are also known to contain mushroom extracts with soothing effects on these skin conditions.

1.2.1.2 Anti-acne and Anti-inflammatory Properties

Acne vulgaris is a most common chronic skin disease and a major cosmetic concern for teenagers. The skin condition results because of blockage and/or inflammation of the pilosebaceous units of the skin. The pilosebaceous unit is composed of the hair follicles and their accompanying sebaceous gland. The skin bacterium *Propionibacterium acnes* (currently *Cutibacterium acnes*) plays an important role in acne vulgaris (pimples). Acne can manifest as both inflammatory lesions (pustules and papules) and non-inflammatory lesions (blackheads and whiteheads). The common area affected is mostly the face but can also affect the back and chest. The causative organism produces lipases that break down oily sebum to fatty acids and glycerol which causes inflammation and the sebum starts to accumulate when the gland is blocked, resulting in acne.

Polycephalomycesphaothaiensis, an insect-infecting fungus (entomopathogenic), is known for its anti-acne activity (Sonyot et al., 2020). The anti-bacterial and anti-inflammatory property exhibited by the extracts of this fungus is a promising new avenue for acne treatment. Several mushrooms also exhibit anti-inflammatory and anti-microbial properties which may have a beneficial effect on acne. In addition, mushrooms are also a rich source of vitamin D that, when applied topically, enables and speeds up the healing process.

1.2.2 Emollient Properties

Emollient property is the quality of softening or soothing the skin. Several studies have established that mushrooms such as *Tremella fuciformis* extracts contain certain polysaccharides that are similar to the functional benefits offered by hyaluronic acid. Hyaluronic acid, which is also referred to as hyaluronan, is an important glycosaminoglycan that is widely distributed throughout the epithelial and connective tissues of the body. Since the mushroom extract mimics the function of hyaluronic acid, it serves as an external moisturizer to the body. The application of snow mushroom extracts can provide adequate moisturizing and hydration to our skin and makes it soft and supple, thereby providing a plumping effect making the skin look youthful.

1.2.3 Exfoliation Properties

Exfoliation is a natural process by which the dead skin cells from the outer layer of the skin are removed. It happens at a frequency of around 30 days, but during this nature's process, not all the dead skin cells are shed completely and therefore a part of the dead cells get accumulated on the surface of the skin, making it prone to infections by bacteria. Chemical exfoliators are available, but it can be sometimes harsh on sensitive skin. The extracts of shiitake mushrooms have bioactive components that can work as natural exfoliators.

1.2.4 Skin Lightening and Brightening Properties

Skin colour is of major cosmetic significance and it is perceived that fairness is an indication of beauty for long. In the quest to achieve fairness, the consumers use skin-lightening cosmetics. Most of these cosmetics work by inhibiting melanin, the pigment that is present in the skin and determines the skin colour of man. In humans, melanin pigments are derivatives of the amino acid tyrosine (precursor). The pathway of conversion of tyrosine to melanin involves a key enzyme called tyrosinase, which catalyzes two steps. Most of the skin-lightening agents work by inhibiting the tyrosinase enzyme, thereby preventing the conversion of melanin. Other ingredients like kojic acid are even more special. It can make melanocytes non-dendritic and thereby decreases the melanin content. Shiitake mushroom is a key source of kojic acid and therefore a natural remedy for skin lightening as well as brightening. Shiitake extracts can reduce both melanin production and their transfer; thus, it can limit the melanin content in the newly formed cells and make them appear relatively lighter. It is also helpful to remove blemishes and scars caused by acne.

1.2.5 Anti-aging

The reishi mushroom is a notable skincare component, since it moisturizes and detoxifies the skin while also preventing premature ageing, redness and fine wrinkles. The high content of beta-glucan aids soften and tone skin complexion; furthermore, the free radical-scavenging ability of this mushroom renders age-related problems in skin and protects skin from damages due to pollution and UV rays. Enriched with antioxidants, mushroom extracts can help to reduce oxidative stress caused by free radicals in our body. This free radical-scavenging activity makes it a potent natural do-good ingredient in anti-aging formulations. The benefit would be double-fold if mushrooms were a part of the daily diet besides being applied externally to the skin surface. Mushroom extracts help to ward off signs of aging like fine

lines, wrinkles, sunspots, etc. The extract can also keep the skin smooth and firm, toned and young looking.

1.3 Application in Hair Care

1.3.1 *Anti-hair Fall*

Pleurotusostreatus extract helps to strengthen the root hairs and reduces hair fall. Mushrooms that are rich in iron such as oyster variety when taken orally also benefit by preventing hair fall. Iron is an essential component that boosts circulation of oxygen to the hair follicles; it enriches the hair roots, thereby enabling the hair to grow healthier, longer and faster. Iron deficiency in our body can lead to anaemia. Therefore, iron deficiency indirectly results in loss of vitality in hair, making the hair prone to breakage. Mushrooms are natural iron supplements which help to invigorate the hair roots.

1.3.2 *Anti-dandruff*

Dandruff is a harmless, chronic condition that usually occurs when the scalp becomes dry or greasy and produces white flakes of dead skin. It is neither contagious nor serious but can be a source of embarrassment with people constantly trying to brush it off their collar or shoulders. Even though dandruff is common, there is no complete cure to this condition till date, although it has been effectively managed. Though dandruff can be related to several predisposing host conditions, it is widely associated with the hyper-proliferation of a commensal lipophilic fungus belonging to the genus, *Malassezia*. *M. globose*, *M. furfur* and other *Malassezia* species have different types of lipases, phospholipases and acid sphingomyelinases, which help in the breakdown of the sebaceous secretions (oils) in the scalp, thereby providing suitable ecological niche for their rapid multiplication. Therefore, most anti-dandruff (AD) products are designed to exert fungicidal action on *Malassezia* species, the primary target in management of dandruff. However, being a commensal flora, the recurrence of growth of these fungi is inevitable. Hence, the challenges in management should focus on the use of synergistic combination of anti-fungals to minimize resistance development and recurrence while also making the AD product skin friendly.

1.3.3 *Hair Moisturizing and Conditioning*

Mushroom extracts, besides preventing hair fall, also help to retain hydration and hereby keeps the hair conditioned and shiny.

1.4 Popular Mushrooms with Cosmetic Benefits

1.4.1 *Tremella fuciformis*

Tremella fuciformis is commonly referred to as a snow mushroom or a white jelly mushroom because of its appearance. The Chinese were the first to discover that this fungus may be used in cosmetics and skincare as well as cuisine. In Korea, Japan and China, the use of mushrooms in skincare has been practised for more than thousands of years. *Tremella fuciformis* has the ability to disseminate and infuse the skin, hydrating it deeply. It also reduces the formation of melanin in the skin, resulting in a brighter complexion. This mushroom is high in vitamin D content that aids in the fading of old acne scars. Oxidative stress is caused by the formation of free radicals in the body, which affects skin structure and promotes premature skin ageing. However, on the other hand, this mushroom has the capacity to combat free radical damage to the skin and may prevent age-related problems. Furthermore, due to its anti-oxidant activity, it can be utilized as an ingredient in eye gel masks to brighten dark under-eye circles and prevent hair loss and is therefore included in shampoos.

1.4.2 *Ganoderma lucidum*

It is popularly known as Reishi mushroom or Lingzhi, which has been celebrated as a food delicacy for centuries and mostly reserved for royal feasts. In addition to the medicinal value, this mushroom had played a pivotal role in skin and hair care routine. *G. lucidum* has a great potential as a skincare ingredient, especially for 'skin-whitening' products. So far, only a few publications have been published that elucidate the chemical makeup of this fungi and its melanogenesis inhibiting actions. Ganodermanondiol, a bioactive molecule, has been discovered to suppress the activity and expression of cellular tyrosinase, as well as the expression of tyrosinase-related protein-1 (TRP-1), TRP-2, and microphthalmia-associated transcription factor (MITF), lowering melanin formation. Furthermore, ganodermanondiol had an effect on the melanogenesis signalling pathway. As a result of these findings, this mushroom is an excellent choice for use as a skin-lightening agent. (Kim et al., 2016). *G. lucidum* polysaccharide (GLP) is a natural antioxidant with no toxic side effects, which can antagonize UVB-induced fibroblast photo aging (Hu et al., 2019). Based on the recent research, *Ganoderma* extracts that contain *Ganoderma* polysaccharides have been used in promoting skin wound healing, mitigating post-burn infection, and preventing skin flap ischemia-reperfusion injury (Yin et al., 2019).

1.4.3 *Ophiocordyceps sinensis*

Ophiocordyceps sinensis, a genus of ascomycete fungi, parasitize mainly on insects. The presence of [polysaccharides](#), sterols, nucleosides, steroids and several other [bioactive compounds](#) in *O. sinensis* makes it useful in the field of alternative medicine for the treatment of several disease conditions. Several research studies have revealed that this mushroom has anti-tyrosinase, anti-elastase and anti-collagenase activity and a photo-protective effect with a sun protection factor up to 25. Therefore, *O. sinensis* can be the best source of cosmetic ingredient for skincare applications. (Cheng et al., [2018](#))

1.4.4 *Agaricus bisporus*

The button mushroom, *Agaricus bisporus*, has recently acquired attention as a source of active substances such as glucans and proteins, as well as for its antioxidant properties and as a functional ingredient in cream as a replacement for natural antioxidants. Because it is a natural material, this mushroom extract is a suitable substitute for EDTA and BHT in all-natural goods. The levels needed to provide the same effect as EDTA have been assessed, and future use of such extracts in natural cosmetics is encouraged (Tables [1.2](#), [1.3](#), and [1.4](#)).

1.5 Formulations

The real challenge in the development of the cosmetic formulation is the right choice of ingredients and the choice of the mushroom/ fungal extract to be used. The choice of the mushroom extract is based on the functional benefits and the planned claims on the product.

Further, in the formulations, the choice of using an oil-soluble or water-soluble extract depends on the type of product. If it is a hair oil formulation, an oil-soluble mushroom extract can be used, whereas for a hair cream, and shampoo formulation, you can use water-soluble extract of mushroom.

In a normal formulation like an anti-dandruff (AD) shampoo or anti-acne cream, one of the major concerns is the possibility of the development of that resistance in commensal flora: For example, (1) resistance in *Malassezia* spp to anti-fungals like zinc pyrithione (ZPTO), octopirox, etc. and (2) resistance in *Cutibacterium acnes* to anti-bacterial ingredients in anti-acne products. This is used in AD products/ anti-acne products due to either cross-resistance or co-resistance mechanisms. Hence, there is a need for the development of eco-friendly and skin-friendly cosmetics and personal care products by lowering the levels of synthetic anti-fungals/anti-bacterials. The use of herbal anti-microbials (fungal/mushroom extracts) in

Table 1.2 Functional/key ingredient in mushrooms with cosmetic applications

| Name of the mushroom | Ingredient/active principle | Properties | Applications | Functional benefits | References |
|--|-----------------------------|--|--|---|---|
| <i>Rhizopus</i> species | Lactic acid | Exfoliating agent, Skin lightening and reduce acne eruption | Skin-care cream | Skin moisturizer, impart smoothness and suppleness | Wilkinson and Goldberg (2018) |
| <i>Candida albicans</i> , <i>Agaricus bisporus</i> , <i>Armillaria tabescens</i> <i>Saccharomyces cerevisiae</i> | Ceramides | Skin-hydrating agents and anti-aging | Skin-care cream, Shampoo, deodorant, restorative facial mask | Prevent moisture loss | Gao et al. (2004) and Anon (2017) |
| <i>Portbellas</i> and <i>Criminis</i> species | L-ergothioneine | Anti-oxidant | Anti-aging creams and lotions | Guard skin from oxidative and DNA damage | Dubost et al. (2006) and Bazela et al. (2014) |
| <i>Lentinula edodes</i> , <i>Grifolafondosa</i> , <i>Pholiota nameko</i> and <i>Auricularia auricula-judae</i> | Trehalose | Free radical scavenging | Moisturizing cream, hydrating cleansing gel | Water retention | Hyde et al. (2010) |
| <i>Inonotus Obliquus</i> | Betulinic acid | Accelerates the regeneration of tissues and skin. Anti-septic helps to heal wounds. Useful in treatment of eczema, psoriasis and dermatitis. | Regenerative creams, anti-septic cream | Prevent skin infections | Megan (2020) |
| <i>Ganoderma lingzhi</i> | Beta-glucans | Anti-inflammatory, anti-redness, antioxidant, anti-wrinkle benefits. | Anti-aging creams, anti-aging cream and moisturizing cream | Keep the skin youthful, healthy and hydrated Attracts water and helps in synthesis of ceramides, which are the building blocks in the protective layer of the skin | Megan (2020) |

(continued)

Table 1.2 (continued)

| Name of the mushroom | Ingredient/active principle | Properties | Applications | Functional benefits | References |
|--|---|--|--------------------------------------|---|---------------------|
| <i>Lentinula edodes</i> | Kojic acid | Skin brightening | Fairness cream, anti-mark cream | Lighten areas of hyperpigmentation, fade dark spots and scars, improve overall complexion | Megan (2020) |
| <i>Lentinula edodes</i> | Selenium, vitamin D, and complex B vitamins | Anti-oxidant | Anti-aging cream | Help combat the effects of inflammation | Megan (2020) |
| <i>Lentinula edodes</i> | L-ergothioneine | Anti-oxidant | Anti-aging cream | Cellular breakdown and encourages a faster cellular renewal process, resulting in healthy collagen and elastin production | Megan (2020) |
| <i>Tremella fuciformis</i> | Similar to hyaluronic acid | Hydration, Antioxidant | Anti-aging cream | Can retain up to 500 times its weight in water, making it a high-efficacy humectant | Megan (2020) |
| <i>Trametes Versicolor</i> | Polysaccharopeptides | Anti-tyrosinase and anti-inflammatory | Skin-lightening cream | To help minimize dark spots, as well as can help reduce redness and irritation | Jhan et al. (2016) |
| <i>Ophiocordyceps sinensis</i> | Ethyl acetate extracts | Anti-elastase activity, Anti-collagenase activity and photo-protective | Anti-aging cream | Can help hydrate dry skin, boost collagen and elastin production, and has even shown some promise as a topical treatment for inflammatory conditions like eczema It helps reduce one of the most important sources of skin aging | Cheng et al. (2018) |
| <i>Boletus edulis</i> , <i>B. pinicola</i> , <i>B. aereus</i> , <i>B. aestivalis</i> , <i>B. erythropus</i> , and <i>B. appendiculatus</i> , can also accumulate considerable amounts of selenium. <i>Agaricus bisporus</i> , <i>A. bitorquis</i> , <i>A. campestris</i> , <i>A. cesarea</i> , <i>A. campestris</i> , <i>A. edulis</i> , <i>A. macrosporus</i> , and <i>A. silvaticus</i> . | Selenium | Anti-dandruff | Anti-dandruff shampoo, cream and oil | Amazing source of selenium, anti-oxidant and anti-dandruff activity | Falandysz (2008) |

Table 1.3 Popular skin -care products in the global market with mushroom extract as functional do-good ingredient

| Product name | Company name | Name of fungi | Properties | Application | References/ website |
|--|--|---|---|-------------|---|
| SS Smile | Shenzhen Youdi Cosmetics Co., Ltd. and Beijing Tongrentang Skincare Research Institute | <i>Ophiotocordyceps sinensis</i> (caterpillar mushroom) | Analeptic and beautifying effects, anti-tyrosinase and anti-elastase activity, anti-collagenase and photo-protective effect | Skincare | Cheng et al. (2018) and Ching Li Tor (2020) |
| Snow mushroom moisturize | The Inkey list | <i>Tremella fuciformis</i> | Enhances skin elasticity | Skincare | Devash (2018) |
| Raw job's tears pore brightening essence | Naruko | <i>Tremella fuciformis</i> | Anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |
| Virgin skin serum | Code of harmony | <i>Tremella fuciformis</i> | For soothe stressed and breakout prone skin, anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |
| Virgin skin tightening and brightening serum | Crave skincare | <i>Tremella fuciformis</i> | Anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |
| Nu-shroom hydrafill serum | Cyberderm | <i>Tremella fuciformis</i> | Anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |
| Hydrating leave on gel primer | Nuface | <i>Tremella fuciformis</i> | Anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |
| Hyaluronic concentrate | The chemistry brand | <i>Tremella fuciformis</i> | Anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |
| Intensive skin boosting toner | Luoki | <i>Tremella fuciformis</i> | Exfoliant, anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |
| Hydro boost 100% hydrogel mask | Neutrogena | <i>Tremella fuciformis</i> | Anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |
| Hydro boost the super hydrator hydrogel mask | Neutrogena | <i>Tremella fuciformis</i> | Anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |
| Hydro boost hydrogel recovery mask hyaluronic acid | Neutrogena | <i>Tremella fuciformis</i> | Anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |

(continued)

Table 1.3 (continued)

| Product name | Company name | Name of fungi | Properties | Application | References/ website |
|--|------------------------------|---|---|----------------------|---|
| 20% vitamin C serum | Purad'or | <i>Tremella fuciformis</i> | Anti-aging, anti-oxidant, moisturizer, humectant | Skincare | https://incidecoder.com/ |
| Enbioment cleanser | DMK | <i>Tremella fuciformis</i> | Skin feels fresh, clean and vibrant | Skincare | https://incidecoder.com/ |
| Sensitive eye smoother | Bioelements | <i>Tremella fuciformis</i> | Shields against irritants and reduce puffiness | Skincare Eye care | https://incidecoder.com/ |
| Soothing reset mist | Bioelements | <i>Tremella fuciformis</i> | Forms a barrier-protective hydrating film to keep skin nourished | Skincare | https://incidecoder.com/ |
| Ufo H ₂ O overdose | Foreo | <i>Tremella fuciformis</i> | Microfiber facemask instantly drenches dry, thirsty skin in soothing moisture | Skincare | https://incidecoder.com/ |
| Euphoria face mask | Joanna vargas | <i>Tremella fuciformis</i> | Instantly soothes, calms and nourishes the skin | Skincare | https://incidecoder.com/ |
| Ultra whitening perfect ampoules | Miguhara | <i>Tremella fuciformis</i> | Anti-oxidant | Skincare | https://incidecoder.com/ |
| Volition's Snow mushroom water serum | Sephora | <i>Tremella fuciformis</i> | Hydrates and locks in moisture for long-term retention | Skincare | www.sephora.com |
| Moon juice plump jelly Hydrating serum | Revolve | <i>Tremella fuciformis</i> | Instantly quenches the skin, gives it some bounce | Skincare | Devash (2018) |
| tata harper clarifying moisturizer | Sephora | <i>Tremella fuciformis</i> | Moisturizer | Skincare | Devash (2018) |
| Kypris glow philtrea | Credo | <i>Tremella fuciformis</i> | Exfoliation and prepare for makeup application | Skincare | Devash (2018) |
| Mega mushroom relief and Resilience soothing face serum, mask and lotion | Dr. Andrew weil for origins™ | <i>Tremella fuciformis</i> and <i>Ganoderma lingzhi</i> | Soothing | Skincare | www.origins.my |

| | | | | | |
|---|------------------------------|--|--|------------------|--|
| Mega-mushroom skin relief advanced face serum | Dr. Andrew weil for origins™ | <i>Inonotus obliquus</i> | Revitalizing | Skincare | www.origins.my |
| Mega-mushroom relief & resilience fortifying emulsion | Dr. Andrew weil for origins™ | <i>Inonotus obliquus</i> | Revitalizing | Skincare | www.origins.my |
| Mega-mushroom relief & resilience soothing face mask | Dr. Andrew weil for origins™ | <i>Inonotus obliquus</i> | Soothing | Skincare | www.origins.my |
| Mega-mushroom relief & resilience soothing cream | Dr. Andrew weil for origins™ | <i>Inonotus obliquus</i> | Soothing | Skincare | www.origins.my |
| Mega-mushroom hydra burst gel lotion | Dr. Andrew weil for origins™ | <i>Inonotus obliquus</i> | Hydration | Skincare | www.origins.my |
| Mega-mushroom skin relief face cleanser | Dr. Andrew weil for origins™ | <i>Inonotus Obliquus</i> | Cleansing | Skincare | www.origins.my |
| Mega-mushroom skin relief micellar cleanser | Dr. Andrew weil for origins™ | <i>Inonotus obliquus</i> | Cleansing | Skincare | www.origins.my |
| Make Prem Chaga concentrate essence | Dr. Andrew weil for origins™ | <i>Inonotus obliquus</i> | Anti-oxidants, soothing protect the skin from free radical damage | Skincare | Driscoll (2019) |
| Embellir night cream | Menard | <i>Ganoderma lingzhi</i> and <i>G. sinense</i> | Eliminate toxins that slow cellular renewal, stimulate cell production and help repair damage | Skincare | www.menard-cosmetic.com |
| Gempyuri self-aesthetic skin care | DXN | <i>Ganoderma lucidum</i> | Multi-massage cleaner Tightening fresh toner Volume energy serum Lifting impact ampoule oil Derma perfection rejuvenation cream DXN Gempyuri mask | Skincare | https://www.dxn2u.com/ |
| Ganzhi E UV defense day cream | DXN | <i>Ganoderma lucidum</i> | Protect against UV rays | Photo-protective | https://www.dxn2u.com/ |

(continued)

Table 1.3 (continued)

| Product name | Company name | Name of fungi | Properties | Application | References/ website |
|---|---------------------|----------------------------|---|----------------------|---|
| Ganzhi Soap | DXN | <i>Ganoderma lucidum</i> | Revitalize the skin and delays the aging process | Cleansing | https://www.dxn2u.com/ |
| Ganzhi liquid cleanser | DXN | <i>Ganoderma lucidum</i> | Cleanses the skin deep into the pores and leaving it clean and refreshed | Cleansing | https://www.dxn2u.com/ |
| Ganzhimoisturising microemulsion | DXN | <i>Ganoderma lucidum</i> | Leaving your skin smooth and radiant all day long | Moisturizer | https://www.dxn2u.com/ |
| Ganzhi toner | DXN | <i>Ganoderma lucidum</i> | Minimize skin pores while leaving skin soft and hydrated | Tones skin | https://www.dxn2u.com/ |
| GanzhiHydrasoft toner | DXN | <i>Ganoderma lucidum</i> | Cleanses and minimizes pores, penetrates and tones skin from the deepest levels | Tones skin | https://www.dxn2u.com/ |
| Ganzhi nourishing night cream | DXN | <i>Ganoderma lucidum</i> | Intensely nourish when the skin naturally renews itself at night | Revitalizes the skin | https://www.dxn2u.com/ |
| Ganzhi™ Lipstick | DXN | <i>Ganoderma lucidum</i> | Hydration | Lip care | https://www.dxn2u.com/ |
| AluminiEye | Alumier | <i>Tremella fuciformis</i> | Improve elasticity and firmness, while diminishing the appearance of fine lines, dark circles and puffiness | Eye cream | Sapphi (2020) |
| Anti-aging liftaway mud face sheet mask | Starskin | <i>Tremella fuciformis</i> | Brightening and rejuvenation | Skincare | Sapphi (2020) |
| Full of grace-naked facial oil | Lush | <i>Agaricus bisporus</i> | Protective and conditioning for skin | Skincare | Sapphi (2020) |
| Adaptogen deep moisture cream | Youth to the people | <i>Reishi mushroom</i> | Hydrate, detoxify and combat premature signs of aging | Skincare | Sapphi (2020) |
| Time stop collagen ampoule | The plant base | <i>Hericiumerinaceus</i> | Anti-oxidants activity, to brighten, soothe and strengthen skin | Skincare | Sapphi (2020) |
| Evercalm- ultra comforting rescue mask | REN clean skincare | <i>Albatrellusovinus</i> | Smooth skin, with even tone and a luminous, healthy complexion | Skincare | Sapphi (2020) |

Table 1.4 Popular hair care products in the market with mushroom extract as functional do-good ingredient

| Product name | Company name | Name of fungi | Properties | References/website |
|---|---------------------------------|---|---|---|
| Ganozhi Plus Shampoo | DXX | Ganoderma | Makes hair silky soft and glossy | https://www.dxn2u.com/ |
| Lingzhi hair strengthening shampoo and conditioner | tianDe | Lingzhi | Powerful anti-oxidant, activating hair growth, protection from UV radiation | https://www.naturalniproducti.com/ |
| Color me brilliant conditioner | Briogeo | <i>Trametes versicolor</i> | Active and natural colour-lock complex that coats colour treated hair, locking in colour and protecting against fade caused by frequent washing | https://briogehair.com/ |
| Anti-hairloss tonifying shampoo | Anuva cosmetics haircare | Reishi mushroom | Prevents hair loss, acts as anti-oxidant | https://www.amazon.in/Anti-Hair-Loss-Tonifying-Mushroom |
| Derma solution shampoo and conditioner | Dr. Groot | <i>Pleurotusostreatitus</i> | strengthens hair elasticity from the roots | https://shopee.com.my/Dr. Groot-Anti-Hair-Loss-Shampoo |
| Reishi mushroom & Wasabi radiant color protection shampoo | Natural hygiene company limited | Reishi mushroom | Hair care and UV protection | https://www2.thaitrade.com |
| Ryo Chouibang premium multi-nutri scalp shampoo | Ryo | White tree ear mushroom, Reishi mushroom, Flower mushroom | Scalp care shampoo | Ryo-Chouibang-Premium-Multi-Nutri-Scalp-Shampoo |
| Organic hair oil-reishi mushroom based | Shazos | Reishi mushroom | Hair oil | https://shazos.com/ |
| Natural nourish snow mushroom deep conditioning mask | AG haircare | <i>Tremella fuciformis</i> | Deep conditioning mask | https://www.ulta.com/p/natural-nourish-snow-mushroom-deep-conditioning-mask-pimprod2023923 |
| The key to volume and growth chanterelle mushroom | Madara | Chanterelle mushroom | 3 min growth boost scalp treatment | https://www.madaracosmetics.com/en/blog/let-your-hair-grow-like-mushrooms |

synergistic combination with synthetic anti-microbial agents with novel delivery methods seems to be a better alternative.

Based on the experimental studies, the authors have shown that the addition of one or more mushroom extracts not only helps to reduce the level of synthetic anti-fungals like zinc pyrithione, climbazole, octopirox in the AD formulations viz shampoo, cream and oil but also make them more efficacious than those popular brands that use stand-alone synthetic anti-fungal ingredients. Similar benefits are obtained in documented skincare products as well.

The complete ready-to-use data of eight cosmetic formulations are provided for the benefit of entrepreneurs. The data provided includes a complete ingredient list of eight formulations, their lab-scale manufacturing process and test specifications. Each of these formulations is standardized based on intensive research and development. The choice of ingredients is based on established literature and extensive R&D and therefore provides the desired functional benefits. Minor modifications and tweaking to suit the personal likings and preferences on colours, perfumes, etc., can be done and standardized.

Formulation 1

1.1 Product Name: Amla and Shiitake Hair Oil

Variant: Coconut Oil

| Phase | Ingredients | Qty in % |
|-------|--|----------|
| A | Mineral oil (Light Liquid Paraffin) | 63.5911 |
| | Cyclopentasiloxane (SF 1202) | 1.0000 |
| B | Mineral oil (Light Liquid Paraffin) | 10.0000 |
| | Paraffin (Paraffin Wax) | 0.1000 |
| | Butylated Hydroxy Toluene (BHT) | 0.0500 |
| C | Mineral oil (Light Liquid Paraffin) | 2.0000 |
| D | <i>Phyllanthus emblica</i> oil-soluble extract | 1.2000 |
| | <i>Lentinula edodes</i> extract | 1.0000 |
| E | E1 <i>Cocos nucifera</i> (Coconut Oil) | 10.0000 |
| | Perfume Amla | 1.0000 |
| | Propylparaben | 0.0500 |
| | D & C Red No 17, C.I.No 26100 | 0.0002 |
| | D&C Green No 6, C.I. No 61565 | 0.0017 |
| | D&C Yellow Np 11, C.I. No 47000 | 0.0070 |
| F | Mineral oil (Light Liquid Paraffin) | 3.0000 |
| | Total | 100.0000 |

1.2 Process

Step 1:

Weigh Mineral oil of Phase A in the main kettle, start stirring. Add SF 1202 into main kettle under mixing.

Step 2:

Weigh mineral Oil of Phase B in the side jacketed kettle, add Paraffin wax and BHT one after another under stirring. Heat to 60 °C under mixing. Stir well to clarity.

Step 3:

Add Phase B into Phase A main kettle under mixing. Rinse side kettle with Phase C mineral oil and add to Phase AB under constant stirring. Stir till phase clear.

Step 4:

Add Amla oil and *Lentinula edodes* soluble extract of Phase D in Phase ABC under constant stirring. Continue stirring for 20 min.

Step 5:

Take Phase E1 coconut oil in side kettle, start stirring.

Take Perfume Phase E2 in separate vessel; add propylparaben and colours one after another under slow mixing. Mix well until Clear. Add E2 into E1 under slow mixing until clear solution.

Rinse this vessel with coconut Oil Phase F and add the rinses to Phase E1 E2. Continue stirring for 20 min.

Step 6:

Add Phase EF into Phase ABCD under continue stirring. Continue stirring for 20 min.

Step 7:

Adjust colour if required.

Step 8:

Sample out for QC Approval.

Step 9:

Discharge bulk through filer after QC Clearance.

1.3 Analytical Parameters

| S. no. | Test | Specification | Method |
|--------|------------------------------|---------------------------------------|----------------|
| 01 | Appearance | Clear, oily liquid | Visual |
| 02 | Colour | yellowish green: To match with master | Visual |
| 03 | Perfume | To match with Master | Olfactory |
| 04 | Specific gravity at 30 °C | 0.84–0.86 | Pycnometer |
| 05 | Refractive index at 40 °C | 1.445–1.455 | Refracto meter |
| 06 | Acid value max | 1.0 | Volumetric |
| 07 | Peroxide value max | 7.5 | Volumetric |
| 08 | Microbial limit ^a | | Micro Manual |
| | a. Total viable count fu/g | Max.100 | |
| | b. Pathogens | Absent | |

^aMeans of the average counts in Nutrient agar and Sabouraud media

Formulation 2

2.1 Product Name: Antidandruff Hair Oil

Variant: Hair nourishing and antidandruff

| Phase | Ingredients | Qty in % |
|-------|--|----------|
| A | Mineral oil (Light Liquid Paraffin) | 91.2361 |
| | Cyclopentasiloxane (SF 1202) | 2.0000 |
| | Tocopheryl Acetate (Vitamin E Acetate) | 0.0050 |
| B | Perfume amla | 0.8000 |
| | Butylated Hydroxy Toluene (BHT) | 0.0500 |
| | Piroctone Olamine (Octopirox) | 0.0250 |
| | <i>Boletus edulis</i> extract | 0.0250 |
| | Propylparaben | 0.0500 |
| C | D & C Red No 17, C.I.No 26100 | 0.0002 |
| | D&C Green No 6, C.I. No 61565 | 0.0017 |
| | D&C Yellow Np 11, C.I. No 47000 | 0.0070 |
| D | Sorbitan Oleate (Arlacel 80) | 1.6000 |
| E | <i>Phyllanthus Emblica</i> Oil Soluble Extract | 2.2000 |
| F | Mineral oil (Light Liquid Paraffin) | 2.0000 |
| | Total | 100.0000 |

2.2 Process

Step 1:

Weigh Mineral oil of Phase A in the main kettle, start stirring. Add SF 1202 and vitamin E Acetate one after another into main kettle under mixing.

Step 2:

Weigh perfume of Phase B in the side vessel with an overhead stirrer. Start stirring, add BHT, Octopirox, *Boletus edulis* extract and Propylparaben. Stir well to clarity.

Then add colours Phase C one after another into Phase B mix under stirring. Continue the stirring under clear solution.

Take Phase D Arlacel 80 into Phase BC under constant stirring. Stir for 30 min.

Then add Amla Extract of Phase E in Phase BCD under constant stirring. Continue stirring for 20 min.

Step 3:

Transfer Phase BCDE mixture under slow stirring in Phase A in the main kettle, continue constant stirring.

Step 4:

Use Phase F for rinsing the vessel of BCDE and add the rinsing to the main kettle of Phase A. Continue stirring Phase APCDEF for 45 min till clarity.

Step 5:

Sample out for QC Approval.

Step 6:

Discharge bulk through filter after QC Clearance.

2.3 Analytical Parameter

| S. no | Test | Specification | Method |
|-------|------------------------------|--|----------------|
| 01 | Appearance | Clear, Oily Liquid | Visual |
| 02 | Colour | Light yellowish green: To match with master | Visual |
| 03 | Perfume | To match with Master | Olfactory |
| 04 | Specific Gravity at 30 °C | 0.824–0.826 | Pycnometer |
| 05 | Refractive Index at 40 °C | 1.445–1.455 | Refracto meter |
| 06 | Acid Value Max | 1.0 | Volumetric |
| 07 | Peroxide Value Max | 7.5 | Volumetric |
| 08 | Microbial Limit ^a | | Micro Manual |
| | a. Total Viable Count fu/g | Max.100 | |
| | b. Pathogens | Absent | |

^aMeans of the average counts in Nutrient agar and Sabouraud media

Formulation 3**3.1 Product: Hair Cream with Oyster Mushroom****Variant: Hair Strengthening**

| Phase | Ingredient | Qty in % |
|-------|---|----------|
| A | A1 Sterile DM water | 67.100 |
| | Disodium EDTA | 0.050 |
| | Methylparaben | 0.230 |
| | Carbomer (Carbopol 940) [To adjust viscosity] | 0.500 |
| | A2 Polysorbate | 1.000 |
| B | Mineral oil-LLP | 13.000 |
| | Stearic acid | 2.000 |
| | Emulsifying wax | 1.000 |

(continued)

| Phase | Ingredient | Qty in % |
|-------|---|--------------|
| | Cetearyl alcohol | 3.200 |
| | Dimethicone | 0.200 |
| | Propylparaben | 0.120 |
| C | Sterile DM water | 3.000 |
| | Triethanolamine 97% | 1.200 |
| D | Sterile DM water | 3.000 |
| | Glycerin | 1.000 |
| | <i>Aloe barbadensis</i> Leaf Juice (Aloe Juice-MJAL 41) | 0.500 |
| | <i>Pleurotusostreatus</i> Extract | 0.500 |
| E | Perfume Ref.PoloGreen-Paria 174069 | 0.400 |
| F | Sterile DM water | 2.00 |
| | Total | 100 |

3.2 Process:

Step 1:

Weigh in sterile DM water from phase A1 in main kettle, add Disodium EDTA and Methylparaben.

Start Stirring. Disperse Carbomer by steady sifting into the vortex. Continue Stirring (with high speed if necessary) for 20 min. Add this to main kettle. Continue stirring. Add Polysorbate of phase A2Heat to 80 °C under constant stirring.

Step 2:

Weigh in all the materials of phase B into a side jacketed vessel. Heat 80 °C under constant gentle stirring. Mix to homogeneity.

Step 3:

Apply vacuum in main kettle. Suck in through muslin cloth of phase B at 90 °C into the main kettle to phase A at 80 °C under full vacuum with slow speed mixing. Mix well at high speed for 5 min and at low speed for 10 min. Cut off steam. Start cooling with cold water.

Step 4:

Suck in TEA solution from [phase C into the kettle to phase AB] at 70 °C. Mix well for 10 min under vacuum. Continue mixing.

Step 5:

Suck in Phase D into the kettle to phase ABC at 55 °C.

Step 6:

At 40 °C, stop mixing and cut off vacuum. Open the kettle, add perfume of phase E into the bulk ABCD. Close the kettle, apply vacuum and mix for 10 min.

Step 7:

Use Sterile DM water of phase F for rinsing of vessels under different steps wherever necessary and suck into the main kettle. Continue mixing for 20 min. Cool to RT.

Step 8:

Stop mixing. Remove vacuum. Sample out the bulk for QC.

Step 9:

Discharge after QC clearance.

Note:

- (i) Hot mixing under open vacuum leads to evaporation losses of water. This has to be compensated by extra water of about 1–2% which may be added to the main kettle along with phase A. This is specific to the type of kettle, its capacity, rpm of stirrer during mixing and cycle time. This process requires standardization to ensure that the moisture content of the finished product is not affected.
- (ii) Ideally, the mixing has to be carried out under closed vacuum as far as possible to keep the moisture loss to the minimum.
- (iii) Cycle time for the batch has to be arrived at on the shop floor over the first 4–5 batches.

3.3 Analytical parameters

| S. no | Test | Specification | Method |
|-------|---|--------------------------|---------------------|
| 01 | Appearance | Opaque, homogenous cream | Visual |
| 02 | Colour | White: to match master | Visual |
| 03 | Perfume | To match master | Olfactory |
| 04 | Viscosity, cps, Brookfield T#6, 0.3rpm, at 30C | 1.2–3.2 Lakhs | Viscometer |
| 05 | pH | 6.50–7.50 | pH meter |
| 06 | Bulk density at 30C, g/cc | 0.970 ± 0.010 | Density Cup |
| 07 | Total fatty matter %B by weight | 19.0 ± 1.0 | Extraction |
| 08 | Loss on drying at 105 C % by weight | 77.5 ± 1.0 | Oven Method |
| 09 | Rancidity | Absent | Extraction & Visual |
| 10 | Microbial Test ^a 1. Total count, cfu/gm 2. Pathogens | <100 Absent | Micro manual |

^aMean of two average numbers on Nutrient agar and Sabouraud Agar

Formulation 4

4.1 Product: Reishi Mushroom Hair Cream

Variants: Hair Fall Control

| Hair cream–hair fall control | | |
|------------------------------|--|----------|
| Phase | Raw materials | %Qty |
| A | A1 Sterile DM water | 36.100 |
| | Disodium | 0.050 |
| | Methyl Paraben | 0.250 |
| | A2 Sterile DM water | 26.000 |
| | Carbomer (Carbopol 940) | 0.400 |
| | A3 Sterile DM water | 2.000 |
| | Polyvinylpyrrolidone (PVP K30) | 0.300 |
| | A4 Polysorbate(Tween 20) | 1.500 |
| B | Mineral oil (Light Liquid Paraffin) | 12.500 |
| | Glycerylmonostearate-SE | 3.000 |
| | Emulsifying wax (Monecol-EW) | 1.000 |
| | White bees wax | 1.000 |
| | Olive oil | 1.000 |
| | Fatty alcohol ethoxylate (Noigen 375) | 0.800 |
| | Cetyl alcohol | 0.600 |
| | Sorbitan Mono Oleate(Arlacel80) | 0.500 |
| C | Phenoxyethanol | 0.400 |
| | Octylmethoxycinnamate | 0.200 |
| | Propylparaben | 0.150 |
| D | Sterile DM water | 2.000 |
| | Triethanolamine 98% | 0.550 |
| E | Cyclopentasiloxane(SF1202) | 1.000 |
| | Cyclopentasiloxane & Dimethiconol(DC 1502) | 1.000 |
| | Amino silicone fluid (DC 2 8566) | 0.300 |
| F | Sterile DM water | 2.000 |
| | Glycerine | 2.000 |
| | Aloe juice | 0.500 |
| | Ganoderma lingzhi extract | 0.200 |
| G | Perfume | 0.600 |
| | Wheat germ oil | 0.100 |
| Total | Sterile DM water | 2.000 |
| | Total | 1000.000 |

4.2 Process

Step 1:

Weigh in sterile DM water from phase A1 in main kettle, add Disodium EDTA and Methylparaben.

Start Stirring.

Disperse Carbomer by steady sifting into the vortex. Continue Stirring (with high speed if necessary) for 20 min. Add this to main kettle. Continue stirring.

Prepare Phase A2 and A3 and add one after another into Main Kettle Phase A1 under slow mixing.

Add Polysorbate of phase A4 to main kettle. Heat to 80 °C under constant stirring.

Step 2:

Weigh in all the materials of phase B into a side jacketed vessel. Heat 80 °C under constant gentle stirring. Mix to homogeneity

Step 3:

Apply vacuum in main kettle. Suck in through muslin cloth of phase B at 90 °C into the main kettle to phase A at 80 °C under full vacuum with slow speed mixing. Mix well at high speed for 5 min and at low speed for 10 min. Cut off steam. Start cooling with cold water

Step 4:

Suck in TEA solution from phase C into the kettle to phase AB at 70 °C. Mix well for 10 min under vacuum. Continue mixing.

Step 5:

Prepare Phase D, Suck in Phase D into the kettle to phase ABC at 55 °C. Continue mixing.

Step 6:

Prepare Phase E, Suck in Phase E into the kettle to phase ABCD at 50 °C. Continue mixing.

Step 7:

At 40 °C, stop mixing and cut off vacuum. Open the kettle; add perfume of phase F into the bulk ABCDE. Close the kettle, apply vacuum and mix for 10 min.

Step 8:

Use Sterile DM water of phase F for rinsing of vessels under different steps wherever necessary and suck into the main kettle. Continue mixing for 20 min. Cool to RT.

Note:

- (i) Hot mixing under open vacuum leads to evaporation losses of water. This has to be compensated by extra water of about 1–2%, which may be added to the main kettle along with phase A. This is specific to the type of kettle, its capacity, rpm of stirrer during mixing and cycle time. This process requires standardization to ensure that the moisture content of the finished product is not affected.
- (ii) Ideally the mixing has to be carried out under closed vacuum as far as possible to keep the moisture loss to the minimum
- (iii) Cycle time for the batch has to be arrived at on the shop floor over the first 4–5 batches.

4.3 Analytical parameters

| S. no | Test | Specification | Method |
|-------|---|--------------------------|-----------------------|
| 01 | Appearance | Opaque, homogenous cream | Visual |
| 02 | colour | White: to match master | Visual |
| 03 | Perfume | To match master | Olfactory |
| 04 | Viscosity, cps, Brookfield T#6,0.3rpm,at 30C | 2–4 Lakhs | Viscometer |
| 05 | pH | 6.50–7.50 | pH meter |
| 06 | Bulk density at 30C,g/cc | 0.970 ± 0.010 | Density Cup |
| 07 | Total fatty matter %B by weight | 23.0 ± 1.0 | Extraction |
| 08 | Loss on drying at 105 C % by weight | 75.0 ± 1.0 | Oven Method |
| 09 | Rancidity | Absent | Extraction and Visual |
| 10 | Microbial Test ^a 1. Total count, cfu/gm 2. Pathogens | <100 Absent | Micro manual |

^aMean of two average numbers on Nutrient agar and Sabouraud Agar

Formulation 5

5.1 Product: Shampoo with Snow Mushroom

Variant: Herbal Conditioner

| Phase | Raw materials | % |
|-------|-------------------|-------|
| A | DM water | 35.72 |
| | Polyquaternium 10 | 0.30 |
| B | SLES 28% | 27.00 |
| | DLSL | 4.00 |

(continued)

| Phase | Raw materials | % |
|-------|------------------------------------|--------|
| | Dimethiconal | 2.00 |
| | Flocare 7 | 1.00 |
| | Glycerine | 1.00 |
| | Kathon CG | 0.05 |
| | Amla extract | 0.25 |
| | Henna extract | 0.25 |
| | Hibiscus extract | 0.25 |
| | <i>Tremella fuciformis</i> extract | 0.25 |
| C | DM Water | 5.00 |
| | EDTA DS | 0.05 |
| | SLES | 9.00 |
| | CMEA | 1.50 |
| | EGDS | 0.05 |
| D | CAPB | 3.00 |
| | CDEA | 1.50 |
| E | DM Water | 2.00 |
| | D panthanol | 0.10 |
| F | Perfume | 0.50 |
| G | Citric acid 25% | 0.20 |
| H | Sodium Chloride 25% | 1.00 |
| I | Tartrazine 1% | 0.50 |
| | Caramnel 10% | 1.40 |
| | B. Blue 0.1% | 0.13 |
| J | DM Water | 2.00 |
| | | 100.00 |

5.2 Process

Check the kettle and clean previous residue before starting the batch. Confirm the cleaning of Kettle.

Water should be heated at 90 °C and cooled.

Step 1

Transfer DM Water from Phase A into main kettle and start stirring and disperse Polyquaternium 10 into water under slow stirring. Avoid lump formulation during dispersion of Polyquaternium.

Step 2

Transfer slowly all ingredients from Phase B into main kettle Phase A one after another by slow stirring. Continue stirring (avoid foaming during mixing).

Step 3

Add Phase C one after another inside jacketed oil kettle and heat to 70 °C under slow mixing. Once bulk reaches temperature 70 °C, stop heating. Cool room

temperature. Transfer Phase C mixture in main kettle bulk Phase ABC under slow mixing. Continue the mixing for 15 min.

Step 4

Add Phase D materials one after another into main kettle Phase ABC by slow mixing. Mix continuously at slow speed-mixing for 15 min.

Step 5

Prepare Phase E & F and add one phase after another phase into bulk main kettle Phase ABCD r slow stirring. Continue the mixing.

Step 6

Adjust pH of the bulk with Citric acid 25% solution. Add Phase G to phase ABCDEF (adjust pH to 6.0 – 7.0). Continue mixing for 5 min.

Step 7

Add sodium chloride 25% solution of phase H into Phase ABCDEFG in small instalments under constant mixing to adjust the viscosity 3000–4000 cps. Continue mixing for 10 min.

Step 8

Add colour Phase I into bulk Phase ABCDEFGH under slow constant mixing.

Step 9

Use Phase J DM water for rinsing of vessels under different steps wherever necessary and add leftover into the bulk.

Step 10

Sample out for QC approval

Step 11

Adjust pH, adjust viscosity if required

Step 12

Discharge after QC clearance

Note:

- (i) Adjusted for SLES active content according to the active content in the supply. The short fall/excess quantity has to be adjusted with DM water of Phase j.
- (ii) Thoroughly mix well the Zinc pyrithione 48% solution drum/pack before weighing for proper homogeneity of the material.
- (iii) Adjust the shortfall/excess quantity of citric acid solution of phase G with DM water of Phase J.
- (iv) Adjust the shortfall/excess quantity of Sodium chloride solution of phase H with DM water of Phase J.

- (v) Stirring has to be slow and uniform so that development of foam during process would be very normal.
- (vi) Cycle time for the batch has to be arrived at on the shop floor over the 4–5 batches.

5.3 Analytical Parameters

| S.no | Test | Specification | Method |
|------|---|---|--------------------------|
| 01 | Appearance | Pearly, light red coloured viscose liquid | Visual |
| 02 | Colour | Match with master | Visual |
| 03 | Perfume | Match with master | Olfactory |
| 04 | pH of product | 6.0–7.0 | pH Meter |
| 05 | Viscosity in cps LV#3, 12 rpm | 3000–4000 | Brookfield Viscometer |
| 06 | Active content as SLES (Mol Wt 384), % by weight | 12–14 | Volumetric |
| 07 | Sodium chloride content % by weight max | 2.0 | Volumetric |
| 08 | Microbial test ^a | | Micro Manual |
| | 1. Total count cfu/g | <100 | |
| | 2. Pathogens | Absent | |

Note:

- (i) Goods should be transferred to dispatch only after QC and microbial clearance.

Formulation 6

6.1 Product: Shampoo with Button Mushroom

Variant: Anti-dandruff

| Phase | Raw materials | % |
|-------|----------------------------------|-------|
| A | DM water | 29.92 |
| | ETD 2020 | 0.20 |
| | TEA | 0.30 |
| B | SLES 28% | 36.00 |
| | DSLS | 4.00 |
| | ZPTO (Zinc Pyrithione 48%) | 1.00 |
| | Agaricus bisporus extract | 1.00 |
| | Flocare 7 | 1.00 |
| | Propylene glycol | 1.00 |
| | Kathon CG | 0.05 |
| C | DM water | 5.00 |
| | Zinc chloride | 0.05 |
| | SLES 28% | 9.00 |
| | CMEA | 1.00 |

(continued)

| Phase | Raw materials | % |
|-------|---------------------|--------|
| | EGDS | 0.05 |
| D | CAPB | 3.00 |
| | CDEA | 1.00 |
| E | DM water | 2.00 |
| | D panthanol | 0.10 |
| F | Perfume | 0.80 |
| | Menthol | 0.20 |
| G | Citric acid 25% | 0.20 |
| H | Sodium chloride 25% | 1.00 |
| I | B. Blue 0.1% | 0.13 |
| J | DM water | 2.00 |
| | | 100.00 |

6.2 Process

Check the kettle and clean previous residue before starting the batch. Confirm the cleaning of Kettle.

Water should be heated to 90 °C and cooled.

Step 1

Transfer DM Water from Phase A into main kettle and start stirring and disperse carbomer into water under slow stirring. Avoid lump formulation during dispersion of carbomer. Neutralize carbomer using TEA from Phase A under missing.

Step 2

Transfer slowly all ingredients from Phase B into main kettle Phase A one after another by slow stirring. Continue stirring (Avoid foaming during mixing).

Step 3

Add Phase C one after another inside jacketed oil kettle and heat to 70 °C under slow mixing. Once bulk reaches temperature 70 °C, stop heating. Cool room temperature.

Step3

Add Phase D materials one after another into main kettle Phase AB by slow mixing. Mix continuously at slow speed mixing for 15 min.

Step 4

Prepare Phases E and F. Add one phase after another into the bulk main kettle Phase ABCD r slow stirring. Continue the mixing.

Step 5

Adjust pH of the bulk with citric acid 25% solution of Phase G to phase ABCDEF to 6.0–7.0. mix 5 min.

Step 6

Add sodium chloride 25% solution of phase H into Phase ABCDEFG in small instalments under constant mixing to adjust the viscosity 3000–4000 cps. Continue mixing for 10 min.

Step 7

Add colour Phase I into bulk Phase ABCDEFGH under slow constant mixing.

Step 8

Use Phase J DM water for rinsing of vessels under different steps wherever necessary and add leftover into the bulk.

Step 9

Sample out for QC approval

Step 10

Adjust pH, adjust viscosity if required

Step 11

Discharge after QC Clearance

Note:

- (i) Adjusted for SLES active content according to the active content in the supply. The short fall/excess quantity has to be adjusted with DM water of Phase j.
- (ii) Thoroughly mix well the Zinc pyrithione 48% solution drum/pack before weighing for proper homogeneity of the material.
- (iii) Adjust the shortfall/excess quantity of citric acid solution of phase G with DM water of Phase J.
- (iv) Adjust the shortfall/excess quantity of Sodium chloride solution of phase H with DM water of Phase J.
- (v) Stirring has to be slow and uniform so that development of foam during process would be very normal.
- (vi) Cycle time for the batch has to be arrived at on the shop floor over the 4–5 batches.

6.3 Analytical Parameters

| S. no | Test | Specification | Method |
|-------|----------------------------------|--|--------------------------|
| 01 | Appearance | Pearly, light blue coloured viscous liquid | Visual |
| 02 | Colour | Match with master | Visual |
| 03 | Perfume | Match with master | Olfactory |
| 04 | pH of product | 6.0–7.0 | pH Meter |
| 05 | Viscosity in cps LV#3, 12 rpm | 3000–4000 | Brookfield Viscometer |

(continued)

| S. no | Test | Specification | Method |
|-------|--|---------------|--------------|
| 06 | Active content as SLES (Mol Wt 384), % by weight | 12–14 | Volumetric |
| 07 | Sodium chloride content % by weight max | 2.0 | Volumetric |
| 08 | Microbial Test ^a | | Micro Manual |
| | 1. Total count cfu/g | <100 | |
| | 2. Pathogens | Absent | |

Note:

(i) Goods should be transferred to dispatch only after QC and microbial clearance.

Formulation 7

7.1 *Product* Face Scrub Cream with Snow Mushroom

Variant Walnut and Snow Mushroom

| Phase | Ingredients | % |
|-------|------------------------------------|---------|
| A | DM water | 63.000 |
| | EDTA DS | 0.050 |
| | SLS powder | 0.200 |
| | Allantoin | 0.100 |
| B | Stearic acid | 16.000 |
| | LLP | 2.000 |
| | IPM | 2.000 |
| | EGDS | 1.000 |
| | Cetyl alcohol | 1.000 |
| | Phenoxy ethanol | 0.400 |
| | Silicone DC 200 | 0.500 |
| C | DM water | 3.000 |
| | KOH | 0.500 |
| D | Walnut scrub | 2.000 |
| E | Glycerine | 1.000 |
| | Cucumber powder | 1.000 |
| | <i>Tremella fuciformis</i> extract | 1.000 |
| F | DM Water | 2.000 |
| | Niacinamide | 0.500 |
| | Kathon CG | 0.050 |
| G | Vitamin E acetate | 0.100 |
| | Vitamin A palmitate | 0.050 |
| | Perfume | 0.500 |
| H | DM water | 2.150 |
| | | 100.000 |

7.2 Process

Check the Kettle and clean previous residue before starting the batch. Confirm the cleaning of Kettle.

Water should heat 90 °C and cool.

Step 1

Transfer DM Water from Phase A into main kettle and start heat at 80 °C. Once water reaches 80 °C, add one after another of remaining phase A material in main kettle under slow stirring.

Step 2

Transfer all ingredients from Phase B into side jacketed oil kettle and heat to 80 °C under slow mixing. Once bulk reaches temperature 80 °C, stop heating.

Step 3

Apply vacuum in main kettle. Suck In, through muslin cloth of Phase B oil Phase at 80 °C into the main kettle to Phase A at 80 °C under full vacuum with under slowing speed mixing. Mix continuously at high-speed mixing for 5 min and at low speed for 10 min at 80 °C. Cut off stream and start cooling.

Step 4

Suck in KOH solution from Phase C into the kettle to phase AB at 70 °C. Mix well for 10 min under vacuum. Continue mixing.

Step 5

Stop stirring and remove the vacuum, start slow stirring and add slowly phase D into Phase AB at 60 °C under mixing. Continue mixing for 10 min and confirm even spreading of scrub in bulk. Continue mixing,

Step 6

Prepare Phase E & F and suck Phase E & F one after another into bulk main kettle Phase ABCD at 55 °C under slow stirring. Continue the mixing under vacuum.

Step 7

Prepare Phase G, and suck Phase G into bulk main kettle Phase ABCDEF at 40 °C under slow mixing. Continue mixing under vacuum 10 min.

Phase 8

Use DM water Phase H for rinsing of vessels under different steps wherever necessary and suck into main kettle. Continue mixing for 20 min. Cool to RT.

Step 9

Stop mixing, remove vacuum. Sample out the bulk for QC approval.

Step 10

Discharge after QC Clearance.

Note:

- (i) Hot mixing under vacuum leads to evaporation losses of water. This has to be compensated by extra water of about 1–2% which may be added to the main kettle along with phase A. This is specific to the type of kettle, its capacity, and rpm of stirrer during mixing and cycle time. The requires to standardize vis-à-vis the moisture content of the finished product over the first 4–5 batches.
- (ii) Ideally, the mixing has to be carried out under closed vacuum as far as possible to keep the moisture loss to the minimum.
- (iii) Cycle time for the batch has to be arrived at on the shop floor over the 4–5 batches.

7.3 Analytical Parameters

| S. no | Test | Specification | Method |
|-------|---|---|-------------------|
| 01 | Appearance | Opaque | Visual |
| 02 | Colour | Light brownish white Match with Master | Visual |
| 03 | Perfumes | Match with master sample | olfactory |
| 04 | Viscosity cps, Brookfield T#4, 0.3 rpm, at 30 °C | 2–4 lakhs | viscometer |
| 05 | pH | 6.50–7.50 | pH Meter |
| 06 | Bulk Density at 30 °C | 0.960–0.980 | Density Cup |
| 07 | Total Fatty Mater % by weight | 14–18 | Extraction method |
| 08 | Loss on drying at 105 °C by weight | 23–26 | oven |
| 09 | Rancidity | absent | Olfactory |
| 10 | Microbial Test ^a | | Micro Manual |
| | 1. Total Count cfu/g | <100 | |
| | 2. Pathogens | Absent | |

^aMeans of the two average numbers on Nutrient and Sabouraud Agar

Formulation 8

8.1 Product: Hand and Body Lotion with snow mushroom

Variant: Moisturizer

| Phase | Ingredients | % |
|-------|-----------------|---------|
| A | DM Water | 74.0500 |
| | EDTA | 0.0500 |
| | Carbomer 940 | 0.2500 |
| | Allantoin | 0.1000 |
| | Methylparaben | 0.2500 |
| | Polsysorbate 20 | 0.5000 |
| B | LLP | 10.0000 |

(continued)

| Phase | Ingredients | % |
|-------|------------------------------------|----------|
| | Stearic Acid | 2.0000 |
| | Cetostearyl alcohol | 2.0000 |
| | Emulsifying Wax | 1.0000 |
| | Isopropyl Myristate | 2.0000 |
| | DC 200 | 0.5000 |
| | Almond Oil | 0.2000 |
| | Propyl paraben | 0.1500 |
| | phenoxy ethanol | 0.4000 |
| C | DM Water | 1.0000 |
| | TEA | 0.7000 |
| D | Glycerin | 1.0000 |
| | Tremella fuciformis extract | 1.0000 |
| | Sodium Lactate 60% | 0.2000 |
| E | Perfume | 0.5000 |
| | Vitamin E Acetate | 0.1000 |
| | Vitamin A Palmitate | 0.0500 |
| F | DM Water | 2.0000 |
| | Total | 100.0000 |

8.2 Process

Check the Kettle cleaning and pervious residue before starting the batch. Confirm the cleaning of Kettle.

Water should heat 90 °C and cool.

Step 1

Transfer DM Water form Phase A into main kettle and start stirring, dissolve EDTA, and disperse slowly carbomer in the water, start to heat at 80 °C. Once bulk reaches 80 °C, add one after another of remaining phase A material in main kettle under slow stirring.

Step 2

Transfer all ingredients from Phase B into side jacketed oil kettle and heat to 80 °C under slow mixing. Once bulk reaches temperature 80 °C, stop heating.

Step 3

Apply vacuum in main kettle. Suck in through the muslin cloth. Add Phase B oil at 80 °C into the main kettle to Phase A at 80 °C under full vacuum and slowing speed mixing. Mix continues high speed mixing for 5 min and at low speed for 10 min at 80 °C. cut off stream and start cooling.

Step 4

Suck in TEA solution from Phase C into the kettle to phase AB at 70 °C. Mix well for 10 min under vacuum. Continue mixing.

Step 5

Prepare Phase D and add into the main kettle Phase ABC at 60 °C under mixing. Continue mixing for 10 min. Continue mixing.

Step 6

Prepare Phase E and suck Phase E into bulk main kettle Phase ABCD at 40 °C under slow stirring. Continue the mixing under vacuum.

Step 7

Use DM water Phase F for rinsing of vessels under different steps wherever necessary and suck into main kettle. Continue mixing for 20 min. Cool to RT.

Step 8

Stop mixing, remove vacuum. Sample out the bulk for QC approval.

Step 9

Discharge after QC Clearance.

Note:

- (i) Hot mixing under vacuum leads to evaporation losses of water. This has to be compensated by extra water of about 1–2% which may be added to the main kettle along with phase A. This is to specific to the type of kettle, it is capacity, rpm of stirrer during mixing and cycle time
 - (a) The requires to standardize vis-à-vis the moisture content of the finished product over the first 4–5 batches.
- (ii) Ideally, the mixing has to be carried out under closed vacuum as far as possible to keep the moisture loss to the minimum.
- (iii) Cycle time for the batch has to be arrived at on the shop floor over the 4–5 batches.

8.3 Analytical Parameters

| S.No | Test | Specification | Method |
|------|------------|---|-----------|
| 01 | Appearance | Opaque lotion | Visual |
| 02 | Colour | Light brownish white Match with Master | Visual |
| 03 | Perfumes | Match with master sample | olfactory |

(continued)

| S.No | Test | Specification | Method |
|------|--|-------------------|-------------------|
| 04 | Viscosity cps, Brookfield 63 0.3 rpm, at 30 °C | 10,000–20,000 cps | viscometer |
| 05 | pH | 5.50–7.50 | pH Meter |
| 06 | Bulk density at 30 °C | 0.960–0.980 | Density Cup |
| 07 | Total fatty matter % by weight | 2–4 | Extraction method |
| 08 | Loss on drying at 105 °C by weight | 23–26 | oven |
| 09 | Rancidity | absent | Olfactory |
| 10 | Microbial test | | Micro Manual |
| | 3. Total Count cfu/g | <100 | |
| | 4. Pathogens | Absent | |

1.6 Conclusion

Mushrooms as a rich source of natural bioactive compounds are finding growing applications in nutraceutical, cosmeceutical and nutritional industries. Several mushrooms and their extracts had already found their places as functional do-good ingredients in various skin-care, lip care and hair care products. Besides cosmetics, mushrooms are now being exploited for their role in personal care products. Several global companies have patented the use of mushrooms/fungi in their formulations with functional claims such as anti-aging, anti-oxidant, anti-wrinkle, anti-aging, skin lightening, moisturizing, anti-acne and anti-dandruff effects. The scope for entrepreneurs to exploit the global opportunity to utilize the appropriate mushroom in their commercial personal care products is expanding. With the immense hidden potential and the resource of a variety of bio-active compounds, there are still several species of fungi/ mushrooms which remain untapped in the industry. In future, more fungi and mushrooms would be studied for their efficacy, safety and utility in cosmetic products. A new world of unexplored fungi/ wild mushrooms would also be discovered, identified, taxonomically classified, verified and cultivated in the future which would boost the cosmetic industry. The future research should be strengthened with cutting-edge technology coupled with the growing omics-driven knowledge on mushrooms. The use of mushroom extracts with multi-functions in the skin and hair care industry would be a trend of the future glamour-driven cosmetic industry.

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Chapter 2

Fungal Endophytes: A Potential Source of Low-Cost Entrepreneurship



Richa Sharma, Amrithesh Chandra Shukla, and Sumpam Tangjang

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2.1 Introduction

De Barry first time coined the term “endophyte”. (De Barry, 1866) and to detect fungi, which live intracellular as well as intracellular in plant tissues without giving any symptom to the plant (Jeewon et al., 2017). While the means of endophytes is “in the plant” (in Greek word *endon*= within, *phyton* = plant) and endophytes emerged mainly from the rhizosphere along with phyllosphere (Ryan et al., 2008). Further, the definition of fungal endophyte as fungal endophytes are colonizers of the alive internal tissues of their host plant (Rollinger & Langenheim, 2018). Endophytic fungi were reported in almost all the plant species viz. algae, mosses, ferns, mainly in angiosperms, as well as gymnosperms from various areas of the world (Doilom et al., 2017).

Several current studies showed the ubiquity of these microbes, and approximately 1 million species of fungal endophytes have been reported from plants (Dreyfuss & Chapela, 1994) and even from lichens (Li et al., 2007a). This prediction is based on a rational assumption that every individual plant species can be occupied by one or more fungal species (Huang et al., 2007).

2.1.1 Classification of Fungal Endophytes

Fungal endophytes generally belong to Ascomycetes, Basidiomycetes, Oomycetes, or Zygomycetes and anamorphic fungi (Arnold, 2007; Sieber, 2007). In addition, there are also various ways of assemblage of fungal endophytes. Transmission way in specific and between the fungal endophytes of the Clavicipitaceae known as clavicipitaceous endophytes while rest are known as non-clavicipitaceous endophytes (White, 1988). Numerous studies have recognized that different fungal endophytes interactions align in the environmental conditions and depend on the host conditions (Faeth et al., 2006).

2.1.2 Reproduction Way of Fungal Endophytes

Reproductive ways of fungal endophytes are normally used synonymously to move to the host along with the population of host plants. Fungal endophytes may produce spores and promote horizontal transmission. Otherwise, hyphae of fungal endophyte may grow into host seeds and transmit to offspring of infested plants, which is generally termed as vertical transmission. However, it is essential to identify those endophytic fungi that may produce mitotic asexual or meiotic sexual spores. While sexual reproduction needs to the production of sexual spores and is therefore always horizontal. Further, asexual reproduction of endophytic fungi is probably through vertical transmission through the host seeds and horizontal

transmission via spores or maybe hyphae (Tadych et al., 2014). Literature reveals that endophytic fungi associated with grasses have concentrated with two associated genera such as *Neotyphodium* and *Epichloë*. Both of them arise systemically and are transmitted vertical way from maternal plants to progeny. However, *Epichloe* fungal endophytes are capable to transmit sexually via spores (Schard et al., 2004). But *Neotyphodium* fungal endophytes are assumed to be harshly transmitted vertically and, hence, considered cached in the host plant (Eaton et al., 2011). On other hand, *Neotyphodium* produces asexual conidia on living plants and on growth culture media (di Menna et al., 2012). Moreover, in woody plants, fungal endophytes have also been observed in seeds and acorns (Petrini et al., 1992), but vertical transmission fungal endophytes in woody plant are probably unusual (Saikkonen et al., 2004). Current facts indicate the horizontal transmission in usually common in grass (Tadych et al., 2014).

2.1.3 Fungal Endophytes of Tissue, Organ, and Host Specificity

Fungal endophytes have been screened from roughly entire aerial tissues; there are very few studies on separately from tissue, organ, and host specificity. Literature reveals that fungal endophytes show remarkable host specificity, comparable to another group, such as smut, rust, and mycorrhizal fungi (Fisher & Petrini, 1990). Organ specificity of fungal endophytes have been reported from wheat plant, Piceaabies white fir Norway and spruce (Sieber, 1985, 1988, 1989). The organ specificity of fungal endophytes occurred due to adaptation to specific microecological and physiological situations (microcosm) found in a particular organ (Petrini et al., 1992). Moreover, previous studies also established that certain level of tissue specificity in fungal endophytes (Wang & Guo, 2007). The host specificity of fungal endophytes from *Fagus sylvatica* and *Pinus sylvestris* and both were growing at the same place, as reported by Petrini and Fisher (Petrini & Fisher, 1988).

2.1.4 Interactions Between Fungal Endophytes and Host Plants

Endophytic fungi can develop a range of forms of interactions with their host plants covering from beneficent mutualism to hidden and dormant pathogens transient through symbiotic microorganisms (Schulz & Boyle, 2005). The asymptomatic colonization of fungal endophytes inside the host-plant tissues might be transient throughout the host plants. The steadiness of the position of communication or the exchange from a type of interaction to a different depends on factors: for example, their nutritional status, developmental stages, genetic character of the two partners,

and different environmental factors (Schulz & Boyle, 2006). However, the development phases of the pathogens of their occurrence within the plants are not observable till the symptoms of the disease come out. This is common to separate identified pathogens as fungal endophytes (Bacon & Hinton, 1996). Asymptomatic colonization may correspond to a prolonged incubation period or a recess infection, which is a style of tolerance of the host plant to meticulous micro-organisms. Different species of *Colletotrichum*, *Alternaria alternata*, *Phomopsis* spp., and *Fusarium* sp., regularly reported in hidden fungal endophytes infections of various host plants (Sinclair & Cerkauskas, 1996). Latent-infecting endophytic fungi may be accountable for the significant deprivation in pre- and postharvest of economically essential crops (Hartman et al., 1986). Fungal endophytes represent thus a continuum of different interactions (Schulz & Boyle, 2005). Like pathogens, fungal endophytes possess necessary tools such as phytotoxic metabolites and exoenzymes to cross plant barriers and to grow within plants (Castro & Fontes, 2005). However, in variation of pathogens, the phenotypic plasticity of fungal endophytes permits them to set up asymptotically within host-plant tissue, intercellularly or intracellularly with additional options such as hidden pathogens, mutualism, and saprophytism, local or also systemic colonization (Schulz & Boyle, 2005).

2.1.5 Plant Protection by Fungal Endophytes

Endophytic fungi are known to protect the host plant in numerous ways, such as producing toxic alkaloids in grasses to prevent herbivores, increasing the tolerance in hot springs to grow, and providing defense against pests in plants (Zhang et al., 2006). Endophytic fungi give to everything with existing pathogenic microorganisms in the host plant. Increasing evidences suggested that fungal endophytes interconnect with the pathogenic microorganisms in diverse ways in different hosts' plants and, resultantly, changes in nutrient balance in support of endophytic fungi or support the defense mechanism of plants, may suppress the growth of the pathogens may be due to altered physiology conditions (Busby et al., 2016). Many fungal endophytes species generate antifungal, antibacterial, antiviral, and antibiotics novel compounds (Istifadah & Mcgee, 2006) and also provide protection against pathogenic microorganisms with compact harshness (Zabalgoeazcoa, 2008). Colonization of fungal endophytes inside the plants enhance the protection against plant nematodes. Fungal endophytes control the functioning of consequently plant's survival, pathogen system, diversity, and conservation (Busby et al., 2016).

Fungal endophytes also protect different herbivores and pests. Roughly thousands of insect pathogenic fungi belong to the class Basidiomycetes and Chytridiomycetes are identified as fungal endophytes; they are closely related with grass fungal endophytes: for example, *Epichloë* and *Claviceps* (Moonjely et al., 2016). Many different mechanisms of fungal endophytes are capable to suggest for this struggle and inducing resistance to diseases. The mechanisms of fungal endophyte encourage decreasing the fitness of plants by increasing their tolerance to

abiotic and biotic stress, resistance is also connected with the nutritional status of the host plants (Aguilar & Barea, 1996). Endophytic fungi *Cryptosporiopsis cf. quercina* and *Colletotrichum* sp. are found capable against phytopathogens such as *Gaeumannomyces graminis*, *Pyricularia oryzae*, *Rhizoctonia cerealis*, and *Phytophthora capsici* (Lu et al., 2000).

Current study suggests that plants infected with fungal endophytes have distinguishable benefits against biotic and abiotic stresses, other nonendophytic counterparts (Bae et al., 2008). Many valuable features have regularly been reported improved competitiveness (Hill et al., 1991), improved tolerance to stressful factors such as microbial infections, heavy metal presence (Reiter et al., 2002), low pH and high salinity (Waller et al., 2005), drought acclimation (Cheplick et al., 2000), enhanced resistance to insect and pests, and herbivores (Akello et al., 2008).

2.2 Fungal Endophytes Producer of Secondary Metabolites

Endophytic fungi are among the generally major groups of eukaryotic microorganisms that are well known for producing numerous new metabolites, which are directly used as drugs or function as lead structures for synthetic modifications (Chin et al., 2006; Mitchell et al., 2008). Endophytes guard their hosts from infectious agents and adverse conditions by secreting bioactive secondary metabolites (Nongalleima et al., 2013). Endophytic fungi benefit the infected plants, for example, it increases the resistance to herbivore grazing through the production of different alkaloids (Owen & Hundley, 2004), improves the growth and competitive capability. They also influence the nutrients uptake and endurance of these plants (Jia et al., 2016), temperature and drought tolerance, plant phenotypic traits, leaf chemistry, propensity for vegetative reproduction and tolerance of heavy metals in soils (Redman et al., 2002). In addition, endophytic fungi have the ability to protect host plants from diseases caused by pathogen microorganisms either through direct effects, for example, by producing antibiotics and lytic enzymes, or indirect effects, for example, by enhancing the plant defense (Ye et al., 2017). Endophytic fungi are able of producing bioactive compounds, which are alike to their host plants, and are able in the conservation of world's deteriorating biodiversity (Mane et al., 2017).

Endophytic fungi producing toxic substances to protect the host plant from pets, insects, and herbivores. Host beneficial endophytes such as *Neotyphodium* and *Epichloë*, which provide the drought tolerance, antiherbivore protection, as well as improved nutrient uptake to the host plants (Schard et al., 2004). Some species of *Stagonospora*, *Piriformosporoides*, and *Acremonium strictum* show parallel functions of defense and growth promotion (Hol et al., 2007). Moreover, endophytic fungi have developed mechanisms to live inside the host plant by shielding themselves beside every chemical and physical weapon of the plants: for example, *Fusarium solani* customized its poisons-binding site by means of alterations in amino acids to escape from damaging effects of camptothecin (Hestekamp, 2017). Endophytic fungi are gaining importance due to their role in plant growth

stimulation, defense against biotic and abiotic stresses, plant growth hormones, higher seed yield, and pests via modulation of growth hormone signaling (Vandenkoornhuysen et al., 2015). Wild relatives of *Aegilops sharonensis* and *Triticum dicoccoides* (wheat) have several valuable endophytic fungi of diverse taxonomic groups, which are not present in cultivated modern-day *T. Aestivum* (Ofek-Lalzar et al., 2016).

Studying endophytic fungal metabolites found that a correlation of habitat and entire biological activity of fungal metabolites (Schulz et al., 2002). Several significant medicines are derived from plants such as taxol, camptothecin, quinine, vincristine, and vinblastine (Ramawat et al., 2009), whereas more than eight thousand five hundred bioactive compounds or secondary metabolites are identified by fungal endophyte origin (Goyal et al., 2017). Fungal endophytes may produce different chemicals as overstated by standard example, like gibberellin production by *Fusarium oxysporum* causing agent of foolish seedling disease in rice. Fungal endophytes produce a plethora of bioactive metabolites of distinctive structure, including alkaloids, phenolic acids, benzopyranones, quinones, flavonoids, steroids, xanthenes, terpenoids, and tetralones (Tan & Zou, 2001). At the present, it has been found that *Cladosporium cladosporioides* and *Metarhizium anisopliae* fungal endophyte can be produced taxol (El-Maali et al., 2018). Accumulation of a fungal endophyte *Taxomyces andreanae* from *Taxus baccata* to produce taxol biosynthesis fueled discover for fungal endophyte associated with promise bioactive compounds and their derivatives (Nicoletti & Fiorentino, 2015). Fungal endophyte strains have been found to be rich sources of numerous novel and valuable bioactive compounds like antioxidants, pigments, anticancer compound, antiviral, antibacterial, antifungal, anti-inflammatory, antitumor, immuno-suppressants, antimalarial, insecticidal, immunomodulatory, antitubercular activities and antimicrobials like azadirachtin A, B, cytochalasin N, citrinal B, diosgenin, germacrane-type ginkgolide-B, gliotoxin, huperzine A, penicillide derivatives, a-pyrone analogues, piperine, taxolcamptothecin, podophyllotoxin, and biocontrol agents (Nassimi & Taheri, 2017). Many researchers have proven that fungal endophytes from medicinal plants are a possible source of novel natural products, especially valuable resources for pharmaceutically and industrially important compounds that can be used in the treatment of diverse life-threatening diseases, as well as different industrial applications (Kaul et al., 2013; Bengtsson-Palme, 2018). Table 2.1 shows the diversity of fungal endophytes and a large quantity of different secondary metabolite or bioactive compounds or production by various fungal endophytes reported from different host plants universally.

2.3 Extracellular Enzyme Activities from Fungal Endophytes

Generally, fungal endophytes have the ability to utilize diverse organic compounds (carbon), which enables them in degradation of structural components such as proteins, lipids, cellulose, hemicelluloses, lignin, glucose, keratin, pectin, and

Table 2.1 Production of secondary metabolites by fungal endophytes

| Fungal endophytes | Secondary metabolites | References |
|--|---|----------------------------|
| <i>Phoma</i> sp. | (3S,4S)-3,8-dihydroxy-6-methoxy-3,4,5-trimethylisochroman-1-one | Kim et al. (2019) |
| <i>Aspergillus niger</i> | Trypacidin A, Methylsulochrin. | Isabelle et al. (2019) |
| <i>Chaetomium</i> , <i>Aspergillus</i> , <i>Alternaria</i> , <i>Penicillium</i> and <i>Charobacter</i> | favonoids, terpenoids | Yuan et al. (2019) |
| <i>Alternaria</i> sp. <i>Fusarium</i> sp. and mycelia sterilia | bioactive secondary metabolites | Prabha et al. (2018) |
| <i>Aspergillus flocculus</i> | Ergosterol peroxide, ergosterol and campesterol | Tawfike et al. (2019) |
| <i>Trichoderma atroviride</i> | Atrichodermone A, B, C | Zhou et al. (2017) |
| <i>Fusarium oxysporum</i> | 4-hydroxybenzoic acid | Bogner et al. (2017) |
| <i>Alternaria</i> sp. Samif01 | Altenuisol | Tian et al. (2017) |
| <i>Chaetomium globosum</i> | Chaetoglobosin A | Dissanayake et al. (2016) |
| <i>Pestalotiopsis neglecta</i> | Phenols, flavonoids | Sharma et al. (2016) |
| <i>Hypocrea virens</i> | Gliotoxin | Ratnaweera et al. (2015) |
| <i>Diaporthe phaseolorum</i> , | Baccatin III | Li et al. (2015) |
| <i>Trichoderma stromaticum</i> and <i>Rhizoctonia solani</i> | Ethanolic extract and Solanionic acid | Ratnaweera et al. (2015) |
| <i>Xylaria</i> sp. | Helvolic acid | Ratnaweera et al. (2014) |
| <i>Stemphylium</i> sp. | Infectopyrones A and B | Zhou et al. (2014) |
| <i>Taxomyces andreanae</i> | Taxol (paclitaxel) | Kusari et al. (2014) |
| <i>Eupenicillium</i> sp. | Eupenicinicol | Li et al. (2014) |
| <i>Aspergillus versicolor</i> | 8-O-methylversicolorin | Dou et al. (2014) |
| <i>Myxotrichum</i> sp. | Myxotrichin A, Myxotrichin D | Yuan et al. (2013) |
| <i>Chaetomium elatum</i> | Xanthoquinodin | Chen et al. (2013) |
| <i>Fusarium oxysporum</i> | Vincristine | Kumar and Kaushik (2013) |
| <i>Ulocladium</i> sp. | Ophiobolin P | Wang et al. (2013) |
| <i>Phomopsis longicolla</i> | Dicerandrol C (24) | Erbert et al. (2012) |
| <i>Penicillium vinaceum</i> | Quinazoline alkaloid | Erbert et al. (2012) |
| <i>Phaeosphaeria</i> sp. | Phaeosphaerin A | Li et al. (2012) |
| <i>Acremonium persicinium</i> | Cordyheptapeptides C–E | Chen et al. (2012) |
| <i>Eupenicillium parvum</i> | Azadirachtin, Azadirachtin A | Kusari and Spittler (2012) |
| <i>Preussia africana</i> | Preussochrome C | Zhang et al. (2012) |
| <i>Phoma</i> sp. | b-sitosterol | Wang et al. (2012) |

(continued)

Table 2.1 (continued)

| Fungal endophytes | Secondary metabolites | References |
|----------------------------------|--------------------------|-----------------------------|
| <i>Muscodorsutura</i> | Isocaryophyllene | Kudalkar et al. (2012) |
| <i>Chaetomium globosum</i> | Gliotoxin | Li et al. (2011) |
| <i>Phomopsis</i> sp. | Cytochalasin N | Fu et al. (2011) |
| <i>Colletotrichum</i> sp. | Methanol | Arivudainambi et al. (2011) |
| <i>Coniochaeta</i> sp. | Coniothiepinol A | Wang et al. (2012) |
| <i>Coniochaeta</i> sp. | Conioxepinol B | Wang et al. (2010) |
| <i>Alternaria</i> sp. | Altenusin | Kjer et al. (2009) |
| <i>Nigrospora</i> sp. YB-141 | Solanapyrone C | Wu et al. (2009) |
| <i>Phomapidodella</i> | Phomodione (43) | Hoffman et al. (2008) |
| <i>Penicillium</i> sp. | Penicilllenols A1 and B1 | Lin et al. (2008) |
| <i>Ampelomyces</i> sp. | Altersolanol A | Alves et al. (2008) |
| <i>Pestalotiopsis</i> sp. | Ambuic acid derivative | Ding et al. (2008) |
| <i>Geotrichum</i> sp. AL4 | 1,3-oxazinane | Li et al. (2007a) |
| <i>Cytonaema</i> sp. | Cytonic acid B | Li, Zhou, et al. (2007b) |
| <i>Aspergillus niger</i> | Naptha-y-pyrone | Zhang and Qi-Yong (2007) |
| <i>Alternaria</i> sp. | Podophyllotoxin | Eyberger et al. (2006) |
| <i>Phialocephala fortinii</i> | Podophyllotoxin | Eyberger et al. (2006) |
| <i>Penicillium janthinellum</i> | Polyketide citrinin | Marinho et al. (2005) |
| <i>Entrophospora</i> | Campotheicin | Puri et al. (2005) |
| <i>Dothiorella</i> sp. | Cytosporone B | Xu et al. (2005) |
| <i>Phomopsis</i> sp. | Phomol | Weber et al. (2004) |
| <i>Fusarium subglutinans</i> | Periconicins | Kim et al. (2004) |
| <i>Pestalotiopsis microspora</i> | Pestacin and isopestacin | Harper et al. (2003) |
| <i>P. microspora</i> | Isopestacin and pestacin | Strobel et al. (2002) |
| <i>Xylaria</i> sp. | Cytochalasins | Wagenaar et al. (2000) |
| <i>Cytonaema</i> sp. | Cytonic acid A | Guo et al. (2000) |

oligosaccharides (Kudanga & Mwenje, 2005). Microorganisms signify a viable substitute source of enzymes, as they may be cultured in large quantities within short time frames by fermentation, they are biochemically diverse, and are agreeable to genetic manipulation (Anbu et al., 2017). Literature reveals that fungal endophytes produce numerous extracellular enzymes, such as amylases, cellulases,

protease, lipases, and laccases. The association of such fungal endophytes with the host plant initiates the release of extracellular hydrolysis (Leo et al., 2016). They hydrolysis the food substances and protect against pathogens (Desire et al., 2014). Moreover, endophytic fungal enzymes are used in the beverage, food, confectionary, pulp and paper, textile and leather industries to shorten the processing of unprocessed materials and pharmaceutical uses (Raju et al., 2015). Further, endophytic fungi were isolated from several medicinal plants, traditionally used in the local communities of the Western Ghats, India, the analysis exposed cellulase, amylase, and pectinase enzymes (D'Souza and Hiremath 2015). Similarly, the qualitative and quantitative analyses of the fungal endophyte have also been reported from some medicinal plants, for the production of extracellular enzymes, including amylase, cellulase, lipase, and protease (Patil et al. 2015a). Fungal endophytes are recognized for accumulating diverse extracellular enzymes such as cellulases, amylase, esterase, protease, lipases, pectinases, laccase, and xylanases which play a major role in defending themselves from the protection reply of the host plants or in up taking the food from the soil (Bezerra et al., 2012; Suto et al., 2002). The several industries that devour microbial enzymes together with biomaterials, cellulose, paper, detergents, leather, cosmetics, food, pharmaceuticals, agriculture, and textiles industries (Yadav, 2015). Table 2.2 shows the production of extracellular enzymes by different endophytic fungi reported from various host plants worldwide.

Several fungal endophytes have also been investigated in the production of enzymes. *Penicillium*, *Aspergillus*, *Trichoderma*, and *Humicola* fungal endophytes were identified to produce cellulase enzyme (Sukumaran et al., 2005). *Sclerocystis*, *Nigrospora*, *Microsphaeropsis*, *Cephalosporium*, and *Phomopsis* fungal endophytes were recognized to produce cellulase enzyme (Peng & Chen, 2007). *Pestalotiopsis aurantiogriseum*, *P. guelpini*, *P. glandicola*, *Fusarium lateritium*, *Xylaria* sp., *Acremonium terricola*, *A. japonicas*, *N. sphaerica*, and *Cladosporium cladosporioides* fungal endophytes were identified to produce pectinases, cellulases, proteases, and xylanase enzyme (Bezerra et al., 2012). *Fusarium oxysporum* fungal endophytes have been identified to produce cellulases enzyme (Onofre et al., 2013). *Chaetomium* sp., *Bisporus* sp., *Cladosporium* sp., *Colletotrichum* sp., *Fusarium* sp., *Rhizoctonia* sp., *Aspergillus* sp., and *Curvularia* sp., fungal endophytes, have been identified to produce proteases, cellulases, amylases, lipases, and activity (Patil et al., 2015a). *Pringsheimia smilacis*, *Ulocladium* sp., *Hormonema* sp., *Neofusicoccum austral*, and *Neofusicoccum luteum* fungal endophytes have been identified to produce cellulases, Glucosidase, and phosphatases enzyme (Khan et al., 2016). *Neofusicoccum austral*, *Ulocladium* sp., *Neofusicoccum luteum*, *Pringsheimia smilacis*, and *Hormonema* sp. fungal endophytes have been identified to produce Laccase enzyme (Fillat et al., 2016). *Colletotrichum* sp., *Nigrospora sphaerica*, *Penicillium funiucias*, *Fusarium solani*, *Macrophomina phaseolina* and *Trichoderma viride*, *Pochonia chlamydosporia*, and *Aspergillus* sp. fungal endophytes have been identified to produce Amylase, cellulose, and lipase enzyme (Wagenaar et al., 2000; Escudero et al., 2016). *Alternaria tenuissima*, *P. rubens*, *Curvularia kusanoi*, *Humicolasp*, *P. pinophilum*, *Phoma* sp., and *Fusarium* sp. fungal endophytes have been identified to produce amylase, cellulase, cellulase

Table 2.2 Extracellular enzyme production by fungal endophytes

| Name of fungal endophytes | Enzymes produced | References |
|---|---|-------------------------------|
| <i>Fusarium</i> sp., <i>Humicolasp</i> , <i>Phoma</i> sp. | Cellulase | Liu et al. (2019) |
| <i>Alternaria tenuissima</i> , <i>Penicillium pinophilum</i> and <i>P. rubens</i> | Cellulase tyrosinase and amylase | Yasser et al. (2019) |
| <i>Curvularia</i> sp. | Lipase | Liu et al. (2019) |
| <i>Thielavia arenaria</i> , <i>Aureobasidium pullulans</i> , <i>Phoma</i> sp., <i>Preussia</i> sp., <i>Sordariomycetes</i> sp., <i>Fusarium proliferatum</i> , <i>Preussia</i> sp., <i>Chaetomium</i> sp., <i>Penicillium citrinum</i> , <i>Phomamedicaginis</i> , <i>Aureobasidium</i> sp., <i>A. pullulans</i> , <i>Dothideomycetes</i> sp., <i>Penicillium citrinum</i> , | Glucosidase, phosphatases and cellulases | Khan et al. (2016) |
| <i>Neofusicoccum luteum</i> , <i>Ulocladium</i> sp., <i>Pringsheimiasmilacis</i> , <i>Neofusicoccum austral</i> , <i>Hormonema</i> sp., | Laccase | Fillat et al. (2016) |
| <i>Nigrosporasphaerica</i> , <i>Macrophominaphaseolina</i> , <i>Colletotrichum</i> sp., and <i>Fusarium solani</i> | Cellulase, amylase, and protease | Ayob and Simarani (2016) |
| <i>Trichoderma viride</i> and <i>Penicillium funiucias</i> | Amylase, lipase, cellulose, protease, | Chaturdevi and Gowrie (2016) |
| <i>Pochoniachlamydosporia</i> | Protease | Escudero et al. (2016) |
| <i>Aspergillus</i> sp. | Amylase | Jurynezz et al. (2016) |
| <i>Aureobasidiaceae</i> , <i>Eurotiales</i> , <i>Chaelomiaceae</i> , <i>Sporomiaceae</i> <i>Incertaesadis</i> , <i>Nectriaceae</i> , | Glucosidases, celluloses, and phosphatases | Khan et al. (2016) |
| <i>Aspergillus terreus</i> | l-asparaginase | Kalyanasundaram et al. (2015) |
| <i>Fusarium</i> , <i>Phoma</i> , <i>Colletotrichum</i> , <i>Penicillium</i> | l-Asparaginase | Chow and Ting (2015) |
| <i>Trichoderma piluliferum</i> , <i>Nodulisporium</i> , <i>Cochliobolus lunatus</i> , <i>C. lunatus</i> , <i>C. australiensis</i> , <i>Gibberellabaccata</i> , <i>G. fujikuroi</i> , <i>Myrmecridiums schulzeri</i> , <i>Phomapataminum</i> , <i>Penicillium commune</i> , <i>Acremonium curvulum</i> , <i>Aspergillus niger</i> , <i>A. Ochraceus</i> , <i>A. ochraceus</i> , <i>A. chartarum</i> , <i>Myrothecium verrucaria</i> , <i>Pithomyces atro-olivaceus</i> and <i>P. glabrum</i> | Xylanase, lipase, cellulase, and protease | Bezerra et al. (2015) |
| <i>Alternaria alternate</i> , <i>Penicillium chrysogenum</i> , <i>Sterile hyphae</i> | Gelatinase, cellulase, amylase, pectinase, xylanase, and tyrosinase | Fouda et al. (2015) |

(continued)

Table 2.2 (continued)

| Name of fungal endophytes | Enzymes produced | References |
|---|--|-----------------------------|
| <i>Colletotrichum</i> sp., <i>Fuzarium</i> sp., <i>Rhizoctonia</i> sp., <i>Cladosporium</i> sp., <i>Chaetomium</i> sp., <i>Curvularia</i> sp., <i>Aspergillus</i> sp., <i>Cladosporium</i> sp., <i>Biosporus</i> sp., | Cellulase, lipase, amylase, and protease | Patil et al. (2015b) |
| <i>Colletotrichum gloeosporioides</i> | Protease, chitinase, and amylase | Rabha et al. (2014) |
| <i>Aspegillusniger</i> , <i>A. Fumigatus</i> , <i>A. flavus</i> , <i>Chaetomium</i> sp., <i>Nigrosporasphaerica</i> , <i>Fusarium</i> sp., <i>F. Chlamydosporum</i> , <i>Phoma</i> sp., <i>Discosia</i> sp., <i>Colletotrichum</i> sp., <i>C. falcatum</i> , , <i>Pestalotiopsis disseminate</i> , <i>Talaromycesmersonii</i> , <i>Alternaria</i> sp., <i>Basidiomycetes</i> sp., <i>Cylindrocephalum</i> sp., <i>Mycelia streilia</i> sp., <i>Mycelia sterilia</i> sp., <i>Myrothecium</i> sp., <i>Coniothyrium</i> sp., <i>Fusicoccum</i> sp., <i>Xylaria</i> sp., <i>Phyllosticta</i> sp., <i>Aspergillus</i> sp., <i>Paecilomycesvariotii</i> , <i>Fusarium oxysporum</i> , <i>F.chlamydosporum</i> , <i>F. solani</i> , <i>Myceliasterilia</i> sp., <i>Penicillium</i> sp., <i>P. longicolla</i> , <i>Myrothecium</i> sp. , <i>Acremonium implicatum</i> , | Amylase, cellulase, laccase lipase protease, and pectinase | Sunitha et al. (2013) |
| <i>Nigrosporasphaerica</i> , <i>Drechsleraawaiensis</i> , <i>Lasiodiplodiatheobromae</i> , <i>Cladosporiumcladosporioides</i> , <i>Curvulariabrachyspira</i> , <i>C. verruciformis</i> , <i>Colletotrichum carssipes</i> , <i>C. gloeosporioides</i> , <i>C. falcatum</i> , <i>Phyllosticta</i> sp., | Amylase, protease cellulase, laccase, and lipase | Amirita et al. (2012) |
| <i>Discosia</i> sp. | Amylase | Hegde et al. (2011) |
| <i>Preussia minima</i> and <i>Alternaria</i> sp. | Amylase | Zhang et al. (2010) |
| <i>Monotospora</i> sp. | Laccase | Weihua and Hongzhang (2008) |
| <i>Melanconiumapiocarpum</i> | Laccase, amylase, and cellulase | Guo et al. (2008) |
| <i>Alternaria</i> sp, <i>Pestalotiopsis</i> sp., <i>Acremonium</i> sp. And <i>Fusarium</i> sp. | Amylase, cellulase, protease, and lipase | Maria et al. (2005) |
| <i>Phomopsis</i> sp. and <i>Colletotrichum</i> sp. | Cellulase, Protease, xylanase, and mannanase | Moy et al. (2002) |

tyrosinase, and laccase enzymes (Vázquez et al., 2018; Yasser et al., 2019). *Acremonium implicatum*, *Phyllosticta* sp., *Cylindrocephalum* sp., *C. falcatum*, *C. gloeosporoides*, *C. truncatum*, *C. brachyspora*, *C. vermiformis*, *Alternaria* sp., *A. calidouustus*, *A. niger*, *A. fumigates*, *Cercospora kikuchii*, *Chaetomium* sp., *Fuzarium* sp., *Drechslera hawaiiensis*, *Drechslera* sp., *Mycelia streilia* sp., *F. oxysporum*, *Isaria* sp., *Pestalotiopsis*, *Penicillium* sp., *P. marneffeii*, sp., *Phoma* sp., *P. viridicatum*, *Phomopsis longicolla*, *Catharanthus roseusto*, *Xylaria* sp.,

Calophyllum phyllum, *Rhizoctonia* sp., *Chaetomium* sp., *Trichophyton tonsurans*, *Cladosporium* sp., *Biosporus* sp., *Curvularia* sp., and *Microsporium gypseum* fungal endophytes have been identified to produce lipase enzyme (Sunitha et al., 2013; Fareed et al., 2017). *Fusarium* sp., *F. Proliferatum*, *Xylaria* sp., *Phomopsisliquidambari*, *Chaetomium* sp., *C. globosum*, *C. gloeosporioides*, *Podosporaanserine*, *Neofusicoccumaustrale*, *N. luteum*, *Botryosphaeria* sp., *T. harzianum*, *Botryosphaeria rhodina*, *B. obtuse*, *B. ribis*, *B. dothidea*, *Monotospora* sp., and *Hormonema* sp. Fungal endophytes have been identified to produce lipase enzyme (Sara et al., 2016; El-Ghonemy et al., 2017).

2.4 How to Isolate Fungal Endophytes

The samples thus collected were washed gently in running tap water to remove the soil and debris. Further, isolation of fungal endophytes was investigated, using the method of Suryanarayanan (Suryanarayan et al., 1998). A total of 1350 plant bits of leaves, stems, and roots cut into small pieces of 0.5 cm²; stems and roots samples were cut into 0.5–1.0 cm. The pieces were then surface-sterilized by dipping them serially in 70% ethanol for 5 seconds and in 4% NaOCl for 90 seconds respectively and finally rinsed in sterile distilled water for 10 seconds. Now, one hundred and fifty segments of the samples were randomly selected and plated on Potato Dextrose Agar (PDA) medium (supplemented with 150 mg/L chloramphenicol) contained in Petri dishes (7.5 cm diam). Petri dishes were incubated (12 h dark: 12 h light cycle) for 25 days at 28 °C, to observe the growth of endophytes (Suryanarayanan, 1992). The fungal hyphal tips, which come out from the sample segments, were isolated and then subcultured on PDA plates without using antibiotics. The pure cultures were maintained on PDA slants and identified morphologically as well as used as stock culture for further experimental studies (Fig. 2.1).

2.4.1 Morphological Observations of the Isolated Fungal Endophyte

The various sporulating structures of fungal endophytes, such as conidia, conidiogenous cells, pycnidia (Coelomocytes), ascospores, asci, and ascocarp (Ascomycetes), conidia, and conidiophores (Hyphomycetes), were prepared semipermanent slides of various stages of fungal spores and studied beneath the microscope. The photographs of the mounted specimens were recorded properly. The fungal endophytes thus characterized were positioned in suitable genera, species, and strains of fungi, with standard monographs and taxonomic keys including the morphology of the fungal colony or hyphae, characteristics of the spores, and reproductive structures (Bhat, 2010).

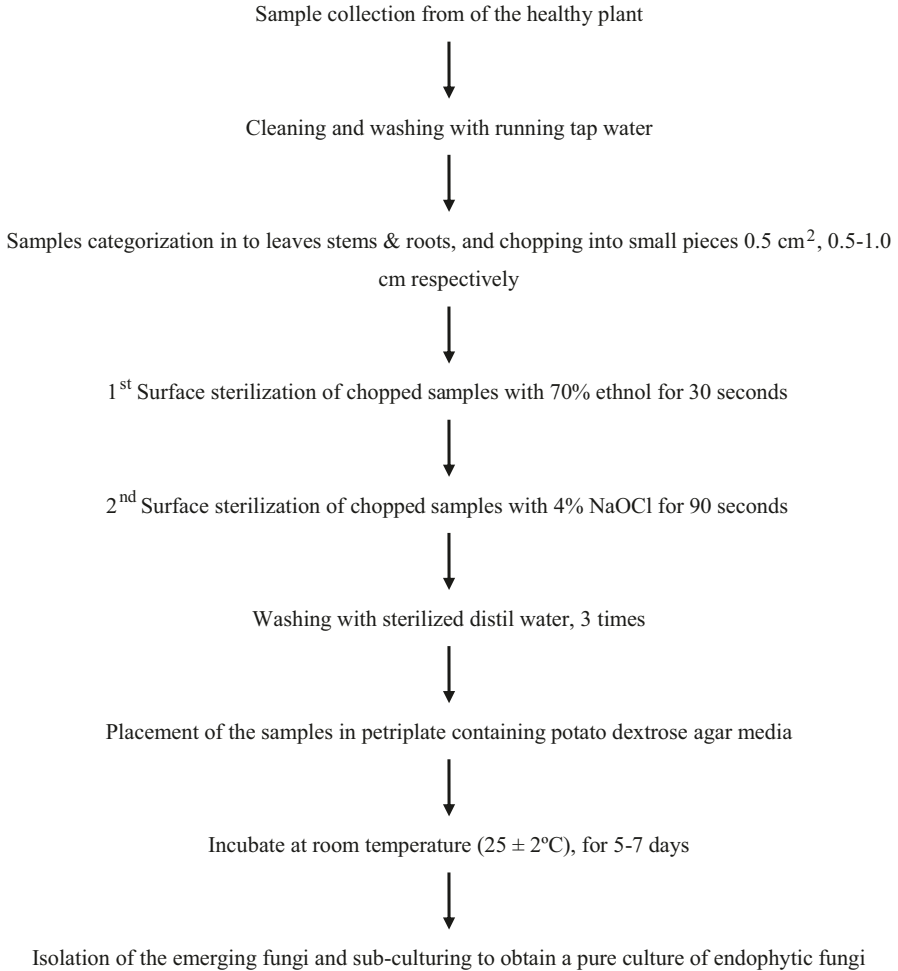


Fig. 2.1 Flow chart procedural steps adopted for isolation of fungal endophytes

2.5 Extracellular Enzyme Analyses from Endophytic Fungi

The production of enzymes by the fungal endophyte was qualitatively screened by growing the fungal mycelium on the solid media having dissolved substrates. All the enzyme activities were carried out at pH 6.0 with slight modifications (Fig. 2.2).

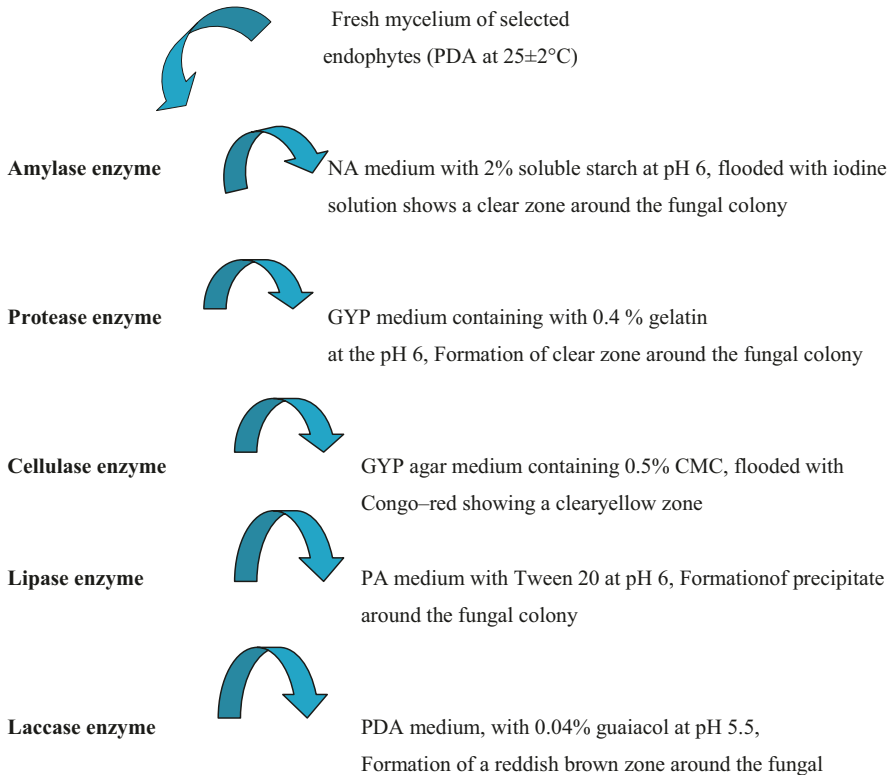


Fig. 2.2 Ray diagram of various extracellular enzyme screening from fungal endophytes

2.5.1 Amylolytic Activity

Amylolytic activity was assessed by growing the isolated fungal endophyte on nutrient agar media with 2% soluble starch at pH 6. After the incubation period of 5–6 days, the plates were applied with 1% iodine along with 2% potassium iodide. The appearance of clear zone surrounding the colony was considered positive for the amylolytic activity (Senthilmurugan et al., 2013).

2.5.2 Proteolytic Activity

Proteolytic activity was screened by growing the fungal isolates on Glucose Yeast Extract Peptone Agar medium (GYP) containing 0.4 % gelatine (pH 6). After 5–6 days of incubation, plates were flooded with saturated aqueous ammonium sulfate, which resulted in a precipitate. This makes the agar hazy and clear zone over all

fungal colony indicating proteolytic activity (Prabavathy & Nachiyar, 2012; Sunitha et al., 2013).

2.5.3 Cellulolytic Activity

The cellulolytic activity of the isolated fungal endophyte was determined by growing it on Glucose yeast extract peptone agar (GYP agar) medium having 0.5% carboxy-methylcellulose (CMC). After incubation (5–6 days) periods, plates were filled with 0.2% aqueous congo red solution and doomed with 1M NaCl for 30 minutes. Which formed yellow areas over all fungal colony indicating cellulose activity (Sunitha et al., 2013; Prabavathy & Nachiyar, 2013).

2.5.4 Lipase Activity

Lipase activity of the isolated fungal endophyte was determined by growing it on Peptone Agar Medium along with sterilized Tween 20 (2 ml). After the incubation period, a clear zone around overall the fungal colony specifies lipase activity (Senthilmurugan et al., 2013; Prabavathy & Nachiyar, 2013).

2.5.5 Laccase Activity

In case of laccase activity, fungal endophyte was cultured in sterilized potato dextrose agar (PDA) medium, supplemented with 0.04% guaiacol (Hi-Media), and adjusted the pH 5.5. These Petri plates were incubated at 28–30 °C for 72 h and then screened for the formation of reddish-brown zones around the fungal colonies indicating the laccase activity (Kalra et al., 2013).

2.6 Fungal Endophytes as a New Source of Entrepreneur

An endophytic fungus is the main source of extracellular enzymes. The association of such fungal endophytes with the host plant initiates the release of extracellular hydrolysis (Leo et al., 2016). Moreover, enzymes are biocatalysts that are involved in catalysis reactions without needing acute conditions, such as high pressures or corrosive environments, temperatures, all of which are often required in chemical processes. Hence, enzymes are used to catalyze reactions in production processes of numerous sectors including environmental bioremediation, industrial bioconversion (biocatalyst), agricultural sectors, pharmaceuticals, and also biotransformations of

several compounds such as flavonoids, alkaloids, and steroids (Wohlgemuth, 2010; Choi et al., 2015). Moreover, there are numerous sources of enzymes including plants, animals, bacteria, fungi, and protists. Microbial enzymes have usually been used because of their easier isolation in high amounts, stability at various extreme conditions, their cocompounds, which are also more controllable low-cost production and less injurious. Microbial enzymes concealed into the media are highly dependable for industrial processes and applications. Microbes isolated from diverse sources, still among species and strains of the same genus, may produce altering levels of enzymes of various properties.

Endophytic fungal enzymes have paying attention for numerous applications, because endophytic fungi secrete large amounts of enzymes into the culture medium and also grow on low-cost materials (Anitha & Palanivelu, 2013). Several fungal enzymes are available commercially including lipases, amylases, cellulases, xylanases, phytases, and proteases (Srilakshmi et al., 2015). The encouraging environmental impact of the manufacture processes is of universally interest and the use of enzymatic reactions as an alternative to organic solvents or chemical reactions is highly valued. Figure 2.3 and Table 2.3 show examples of significant endophytic fungal enzymes and the enzyme sources that are used in several applications; however, only a few fungal strains meet the criteria for industrial production. Generally, applications of enzymes in the food industry have focused on hydrolytic reactions

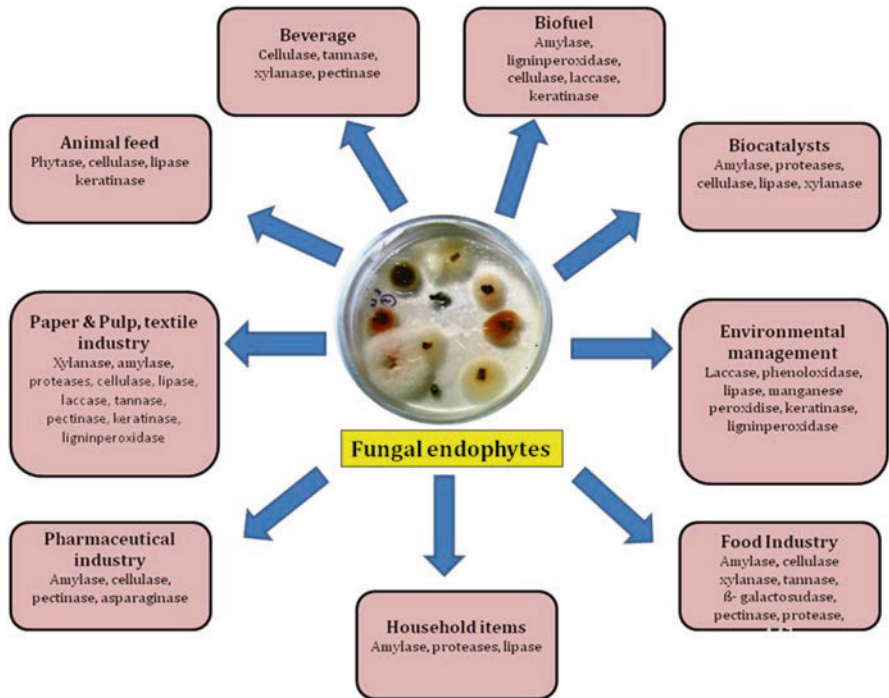


Fig. 2.3 Applications of fungal endophyte origin enzymes

Table 2.3 Examples of enzyme sources and applications of fungi

| Enzymes | Applications | Fungal sources | References |
|-------------------|--|--|---|
| Tannases | Food processing; removal of phenolic substances, juices and wines clarification and stabilize the beverage, reduction of hydrolysable tannin in poultry feeds, production of gallic acid from gallotannins | <i>Aspergillus aculeatus</i> , <i>A. Awamori</i> , <i>A. Caespitosus</i> , <i>A. Niger</i> , <i>A. versicolor</i> , <i>Penicillium charlesii</i> , <i>P. crustosum</i> , <i>P. variable</i> , <i>P. restrictum</i> | Ahmed and Rahman (2014), Bagga et al. (2015), Batra and Saxena (2005) and Srivastava and Kar (2009) |
| Amylases | Hydrolysis of starch in starch processing industry; food and dairy industry; textile industry; pulp and paper industry; detergent industry; pharmaceutical industry; animal feed industry | <i>Aspergillus fumigatus</i> , <i>A. Niger</i> , <i>Cylindrocephalum</i> sp., <i>Lentinula edodes</i> , <i>Penicillium citrinum</i> , <i>P. fellutanum</i> , <i>Rhizopus stolonifer</i> | Ko et al. (2005), Sahoo et al. (2014) and Saleem and Ebrahim (2014) |
| Keratinases | Biomass conversion into biofuels; hydrolysis of keratinous wastes such as feather, hair, and horn; eliminating horny epithelial cells adhering to textile fibers; reducing the environmental pollution | <i>Aspergillus oryzae</i> , <i>A. Parasiticus</i> , <i>Doratomyces microspores</i> , <i>Paecilomycesmarquandii</i> | Anitha and Palanivelu (2013), Farag and Hassan (2004), Friedrich et al. (2005), Gradisar et al. (2000) and Veselá and Friedrich (2009). |
| Cellulases | Animal feed industry; pulp and paper industry; detergent industry; food processing; juices and wines clarification; textile industry; biomass conversion into biofuels | <i>Aspergillus niger</i> , <i>Lentinula edodes</i> , <i>Trichoderma Longibrachiatum</i> , <i>Volvarielladiplasia</i> | Pachauri et al. (2017), Wang and Hsu (2006) and Puntambekar (1995) |
| Lignin peroxidase | Lignin degradation; biomass conversion into biofuels | <i>Aspergillus sclerotiorum</i> , <i>Cladosporium Cladosporioides</i> , <i>Mucor racemosus</i> , <i>Phanerochaete Chrysosporium</i> , <i>Sparassis latifolia</i> | Chandrasekaran et al. (2014), Brugger et al. (2004) and Wen et al. (2009) |
| Laccase | Biopulping biobleaching deinking in pulp and paper industry; lignin degradation; pharmaceutical industry; removal of phenolic substances and stabilize the beverage; biomass conversion into biofuels | <i>Agaricus subrufescens</i> (as "blazei"), <i>Coniophoraputeana</i> , <i>Ganoderma</i> sp., <i>Omphalotusolearius</i> , <i>Phanerochaetefloridensis</i> , <i>Pleurotusostreatus</i> | Arora et al. (2002), Lee et al. (2004), Ullrich et al. (2005) and Songulashvili et al. (2007) |

Table 2.3 (continued)

| Enzymes | Applications | Fungal sources | References |
|----------------------|--|---|--|
| Xylanases | Pulp and paper industry; animal feed industry; bread-making; juice and wine industries; xylitol production; prebiotics production; food processing; textile industry; juices and wines clarification; biomass conversion into biofuels | <i>Aspergillus foetidus</i> , <i>A. Niger</i> , <i>Talaromyces amestolkiae</i> | Chapla et al. (2012), de Alencar et al. (2013) and Nieto-Domínguez et al. (2017) |
| Lipases | Degradation of fat in wastewater treatment; animal feed industry; pulp and paper industry; detergent industry; food processing; leather processing; textile industry; pharmaceutical industry | <i>Aspergillus</i> sp., <i>Curvularia</i> sp., <i>Fusarium solani</i> , <i>F. Verticillioides</i> , <i>Penicillium</i> sp., <i>P. restrictum</i> , <i>P.wortmanii</i> , <i>Rhizopus oligosporus</i> , <i>Trichoderma</i> sp., <i>T. atroviride</i> , <i>T. harzianum</i> , <i>Mucor</i> sp. | Facchini et al. (2015), El-Ghonemy (2017), Marques et al. (2014), Nwuche and Ogbonna (2011), Ul-Haq et al. (2002) and Maia et al. (2001) |
| Manganese peroxidase | Lignin degradation; biomass conversion into biofuels | <i>Lentinula edodes</i> , <i>Phlebiaradiate</i> , <i>Omphalotusolearius</i> , <i>Phellinus robustus</i> | Arora et al. (2002) and Songulashvili et al. (2007) |
| Pectinases | Juices and wines clarification; textile industry <i>Aspergillus japonicus</i> | <i>Aspergillus oryzae</i> , <i>Penicillium viridicatum</i> , <i>P.chrysogenum</i> , <i>Thermoascusaurantiacus</i> | Songulashvili et al. (2007), Banu et al. (2010), Biz et al. (2016) and Martins et al. (2002) |
| Phytases | Feed supplement in diets in animal feed industry; improvement of soil fertilization and nutrient uptake by plants; reduction in the excretion of phosphorus in manure; reducing phosphate pollution in soil and water | <i>Aspergillus</i> sp., <i>A.niger</i> , <i>Mucor</i> sp., <i>Rhizopus oligosporus</i> , <i>Rhizomucor pusillus</i> | Bei et al. (2009), Casey and Walsh (2004), Chadha et al. (2004) and Saxena et al. (2005) |
| Proteases | Detergent industry; food processing; pharmaceutical industry; leather processing; textile industry | <i>Aspergillus</i> sp., <i>Fomitellafraxinea</i> , <i>Humicola</i> sp., <i>Mucor</i> sp., <i>Penicillium</i> sp., <i>Pleurotuscitrinopileatus</i> , <i>Rhizopus</i> sp., <i>Thermoascus</i> sp., <i>Thermomyces</i> sp. | Cui et al. (2007) and Lee et al. (2006) |

(Choi et al., 2015). Glycoside β -galactosidase and hydrolases are involved in the production of a dietary material composed of nonstarch polysaccharides, prebiotics, and oligosaccharides, such as galacto-oligosaccharides, inulin fructo-oligosaccharides, breast milk oligosaccharides, and lactulose that selectively encourage the growth of beneficial intestinal microorganisms in humans (Torres et al., 2010). Pectinases, xylanases, and cellulase are extensively used for the clarification of juices and wines.

Amylases are used to hydrolyze starch in the starch liquefaction process and convert starch into glucose syrups (Souza & Magalhaes, 2010). Further, starch is used as the strengthening agent in the desizing process to prevent the breaking of the warp thread used within the weaving process in the textile industry. Amylases are used for desizing starch in the textile industry, because they provide products that do not degrade the fibers. Moreover, α -amylases are used in the pulp and paper industry instead of starch for coated paper and textiles. The quality of textiles and paper coated with desizing starch is secluded against mechanical harm during processing and finishing. The recycling of waste paper is an eco-friendly trend in the paper industry. Further, amylases, proteases, and lipases enzymes are used for dishwashing detergents. However, amylases are the second other types of enzymes used in the formulation of enzymatic detergent, and 90% of all liquid detergents contain amylases (Souza & Magalhaes, 2010).

Tannases are used to decrease tannin levels in food products such as tea, fruit juices, wines, and beer (Yao et al., 2014). Moreover, these enzymes are used to hydrolyze gallic acid esters and produce gallic acid, which are used as a substrate for the synthesis of food preservatives (Yao et al., 2014; Dhiman et al., 2017). Tannase enzymes are also used for high-grade leather tannin preparation in the leather industry (Dhiman et al., 2017) and this enzymatic process is eco-friendly and also helps to manage waste resulting from production processes.

However, the enzymatic reactions of pectinases, cellulases, xylanases, and lipases aid in the elimination of contaminated ink (Bhat, 2000). Cellulases and proteases are used in the polishing step for clear dyeing to get better color and surface brightness, and oppose textile wrinkling. Lignin peroxidases, xylanases, manganese peroxidases, and laccases are all used to improve the quality of the pulp by removing hemicelluloses and lignin, which are typical impurities in the pulp and paper industries (Choi et al., 2015).

Lipases are hydrolytic enzymes capable of cleaving the ester bond of triacylglycerol as well as catalyze ester synthesis *in vitro* by shifting the equilibrium of the reaction (Mohamed et al., 2011). These enzymes have a broad range of industrial applications in the production and processing of detergents, oils, fats, dairy products, therapeutic agents, and biodiesel (Gog et al., 2012). Ligninolytic enzyme has a variety of isoforms, of which manganese-dependent lignin peroxidase, peroxidase, and laccase are the three main classes. These enzymes are directly concerned with the degradation of diverse xenobiotic compounds (polycyclic aromatic hydrocarbons) and lignin in their normal lignocellulosic substrates (Pointing, 2001). White-rot fungi are capable of degrading the lignin and regarded as the most capable

producers, which secrete ligninolytic enzymes in nature. Laccases (oxygen oxidoreductases and benzenediol) are glycosylated polyphenol oxidases, which degrade lignin powerfully (Kunamneni et al., 2008). These enzymes have significant applications in detoxification of phenolic pollutants, animal biotechnology, pulp and paper industry, and biotransformation (Kunamneni et al., 2008). Protease enzymes are generally used in leather processing for the dehairing and debating of skins and hides instead of chemical processes that have negative implications for the environment. While these proteases enzymes (aspartic protease, protease, cysteine, serine, metalloprotease, and protease) are a significant class of enzymes that catalyze hydrolysis of protein (Oliveira et al., 2014). These enzymes have important applications in food processing industries, leather industries, detergent industry, pharmaceuticals, and biotechnology (Joo et al., 2003).

Xylanases, β -glucanases, and Cellulases have used been in cereal-based feed for monogastric animals for example poultry and pigs, which are not capable to fully degrade and utilize plant-based feeds containing huge amounts of hemicellulose and cellulose (Kirk et al., 2002). However, adding phytases to the feed improves utilization of normal phosphorus bound in phytic acid in cereal-based feed for monogastrics. Usually, 80–90% of whole plant phosphorus is bound to phytic acid, which makes it complicated to monogastrics to utilize. Thus, accessible inorganic phosphates have to be supplemented to the feed in order to reach the necessary concentration. Phytases can reduce the need for supplementation of inorganic phosphorus in the feed (Kirk et al., 2002). However, cellulases are usually used along with the supplement of pectinases, hemicellulases, ligninases, and associated enzymes (Adav & Sze, 2014).

Generally, those fungal endophytes belong to the basidiomycetes, which can enzymatically attack the polymers in the complex-structured lignocellulose. The white rot fungi involve the nutrient recycling of wood required for new plant growth as well as evolutionary impact on diversification of plants (Kües, 2015). However, fungal endophytes are documented as one of the excellent lipase sources among microorganisms (Facchini et al., 2015). The most significant step to recovering the biological degradation of fatty wastewater is pretreatment procedures in order to hydrolyze and dissolve fats, which are able to accelerate the process by declining the fat adsorption to the surface of the anaerobic sludge. Whereas not limiting the transport of the soluble substrate to the biomass (Facchini et al., 2015; Valladao et al., 2009). Moreover, phenoloxidases, laccases, peroxidases, and dioxygenases enzymes are good examples for wastewater treatment (Duran & Esposito, 2000). However, effluents from slaughterhouses have high concentrations of biodegradable organic matter, mainly of which consists of proteins and lipids with low degradability. Subsequently, before effluents are free into the environment, it needs to reduce fat oil and protein from these wastewaters.

Probable applications of enzyme treatments are reduction of the organic matter, which contributes to a cleaner effluent (Valladao et al., 2011). Further, white rot fungi make lignin-degrading enzymes such as, lignin peroxidases, manganese peroxidases and laccases used in biotechnology for degradation of broad-spectrum

intractable organic pollutants, bioremediation of polycyclic aromatic hydrocarbons (PAHs) as well as chlorinated hydrocarbons in the environment (Gao et al., 2010).

Biotechnological applications for practical use have restraining factors, as they have need of large amounts of enzymes. Hence, the productions of low-cost and readily available enzymes possessing satisfactory operating characteristics are demanding. After all, each industrial application can need specific properties of the biocatalysts. Further, researchers still face challenges in searching new enzymes that could result in new applications through better isolation, study of enzyme constancy at severe conditions, and purification procedures. The selection of enzymes for the more stable and stringent conditions is significant for industrial processes and applications. Hence, the search for new microorganisms that have the required properties is a continuous process. Besides, the production and expression of recombinant enzymes through protein engineering technology, it is also required to find enzymes of required characteristics in specific host cell Industrial. Enzymes are usually produced in bioreactors which contain up to over two lakhs liters and are consequently purified in the industrial downstream processes.

Thus, the fungal endophytes that are able to use for such processes have to be speedy growing but not be pathogens or mycotoxins producers. The use of heterologous appearance, recombinant DNA technology, gene cloning to get better production yield of enzymes and their activity can avoid the difficulties related to the production of large quantities.

2.7 International Companies Manufacturing Enzyme from Fungal Endophytes

Fungal endophyte origin enzymes are progressively replacing conformist chemical catalysts in numerous industrial processes. Enzymes have many advantages over chemical catalysts, as well as the capability to function under comparatively gentle conditions of pH, pressure, and temperature. Amylase enzyme was first produced via Solid State Fermentation (SSF) by *Aspergillus oryzae* on wheat bran or moist rice. Jokichi Takamine initially developed this process and got patent in USA in 1884. Moreover, the major production of fungal enzymes was approved out by inundated fermentation technology in the 1940s. Current days, there are numerous international companies having wager in producing industrial enzymes from endophytic fungi. In India, Biocon India Ltd. is a major mass enzyme producer company, but it is not a major at the international level. Table 2.4 is showing the top fourteen companies at the international level.

Table 2.4 Major bulk enzyme-producing companies

| S. no. | International companies |
|--------|---------------------------------|
| 1 | Amano Pharmaceutical Co., Japan |
| 2 | Meito Sankyo Co., Japan |
| 3 | Enzyme Development Corp., USA |
| 4 | Nagase Biochemicals Ltd., Japan |
| 5 | Solavy Enzymes gmbh, Germany |
| 6 | Biocatalysis Ltd., Wales |
| 7 | Rhone-Poulenc, England |
| 8 | Novo Nordisk, Denmark |
| 9 | Rohm gmbh, Germany |
| 10 | Danisco Cultar, Finland |
| 11 | Sankyo Co., Japan |
| 12 | Yakult Biochemical Co., Japan |
| 13 | Shin-Nihon Chemical Co., Japan |
| 14 | DSM-GIST, Netherlands |

2.8 Biotechnological Potential of Fungal Endophytes

Approximately four thousand secondary bioactive compounds or metabolites of fungal origin were reported as biologically active (Dreyfuss & Chapela, 1994). Endophytic fungi are now established as an unused resource for producing significant natural products contributing potential for pharmaceuticals, agriculture, food, and other industrial exploitation (Suryanarayanan et al., 2009). Many novel natural bioactive compounds or secondary metabolic are now known to have immunosuppressive properties, antidiabetic, antioxidants antiviral, anticancer, and insecticidal, etc. (Strobel & Daisy, 2003).

Nowdays broad variety of bioactivities of unstable organic compounds produced by endophytic fungi, *Muscodor albus* Worapong, Strobel & W.M. Hess and *M. roseus*-Worapong, Strobel & W.M. Hess was established in controlling the plant pathogenic fungi (Stinson et al., 2003) and nematodes (Riga et al., 2008). Moreover, *M. albus* against various plants and human-pathogenic fungi and bacteria as an antibiotic (Strobel et al., 2001). In current times *Coryne* sp. (Strain-NRRL 50072) elected as anamorphic form of an *Ascocoryne* sp, was established to produce hydrocarbon derivative (Strobel et al., 2010). Further, *M. albus* strain GBA was newly isolated from *Ginkgo biloba* collected in Newport, RI, USA, which inhibited and killed sure test microbes by production of a mixture of volatile compounds (Banerjee et al., 2010).

However, several endophytic fungi were reported as taxol producers, viz. *Stegoleriumkukenani*, *Alternaria* sp., *Pestalotia* sp., *Pithomyces* sp., *Fusarium* sp., *Tubercularia* sp., *Monochaetia* sp., Bat. & Peres, *Pestalotiopsis guelpini* (Desm.), *Pestalotiopsis microspora* (Speg.) Steyaert, *Sporormia minima* Auersw., *Periconia* sp., *Trichothecium* sp. and *Seimatoantleriumnepalense* Bashyal et al. which were reported from yew as well as other plants (Strobel et al. 20001; Wang et al., 2000). In India, attempts were finished to isolate taxol among potent cytotoxic action from

two fungal endophytes, viz. *Bartaliniarobillardoides* Tassi and *Colletotrichum gloeosporioides* isolated from *Aegle marmelos* and *Justicia gendarussa* Burm F respectively (Gangadevi & Muthumary, 2009).

Fungal endophytes belong to the genus *Pestalotiopsis* have been accepted as the ordinary endophytic fungi from rainforests (Nag Raj, 1993). *Pestalotiopsis* species were produced numerous bioactive compounds and secondary metabolites (Li et al., 2001). In the similar way, *Pestalotiopsis terminaliae* G.P. Agarwal & Hasija from *Terminalia arjuna* plant and *P. Pauciseta* (Sacc.) Y. X. Chen from *Cardiospermum helicacabum* Linn. fungal endophytes were produced taxol (Gangadevi & Muthumary, 2009).

Camptothecin was isolated from different plants such as *Nothapodytesfoetida*, *Camptotheca acuminata* Decais, and *Ervatomiayahyneaana Ophiorrhizamungos* (Rehman et al., 2009). Further, Camptothecin screened from an unidentified fungal endophyte from the inner bark of plant *Nothapodytesfoetida* and *N. nimmoniana* in the Western coast of India (Chapla et al., 2012). However, Camptothecin screened from *Entrophosporainfrequens* (I.R. Hall) R.N. Ames & R.W. Schneid fungal endophyte isolated from the inner bark of *N. foetida* in Jammu and Mahabaleshwar regions of India (Amna et al., 2006).

2.9 Conclusion

Fungal endophytes are the group of microorganisms, which dwell in the inner tissues of the plant parts (viz., roots, stems, leaves, bark, blossoms, and so forth) in symbiotic manner without bringing on any harmful impacts on the plants. Endophytes assume a significant part in assisting plants with battling against biotic and abiotic stresses and improve plant development. Fungal endophytes, obtained from medicinally important plants, are considered as a choice to deliver the bioactive constituents that could be used for the management of various human diseases. Taxol is an anticancer bioactive metabolites that has been extracted from various genera of fungal endophytes. Similarly, fungal endophytes show their efficacy to fight against several pathogens which contain drug resistance activity.

Thus, its wider applicability ranges from agricultural to pharmaceutical sectors could also be explored as potential agent for developing entrepreneurship.

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Chapter 3

The Development of White-Rot Fungi as a Mycoremediation Product



A. A. Ngadin, E. Taghavi, and T. Eaton

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3.1 Introduction

3.1.1 *Habitat's Contamination*

A variety of environmental problems have affected economies and policies around the world (Tsui, 2020). There are many environmental problems related to community behavior, such as producing habitat pollution, health risks and problems, government organizations, the private sector, and industry.

Habitat contamination is one of the environmental issues that all human beings are responsible to solve immediately. Water pollution will lead to the disappearance of biodiversity and aquatic ecosystems. In other words, the aquatic organisms will lose their home. Many reported articles include from Global Water Pollution and Human Health, stating pollution of freshwater resources, such as lakes, rivers, and groundwater, has emphasized chemical pollution. This includes both organic and inorganic pollutants affecting the level of water pollution

A major challenge in determining environmental risks for heavy metals and metalloids is associated with their contrast activity under different redox conditions. Oxidation or reduction reactions, complexes, absorption, and precipitation are key processes that determine their transport and bioavailability. If there is oxygen and in a state of reduction, the solubility of most metal elements is very different. Redox-sensitive metals, iron, and manganese being the most abundant, forming finely dispersed oxide particles that highly absorb heavy metals and metals under toxic conditions.

Interestingly, when oxygen has been depleted, these oxide particles undergo a reductive decomposition solution and release a toxic load. While microorganisms also function to control the precipitation and breakdown of reactive particles in the atmosphere. Most aquatic organisms including microbes are very sensitive to any changes in their environment. They respond to and adapt to this new environment, such as polluted conditions, in many ways. Fewer responses may include reduced reproductive capacity and also the removal of certain enzymatic systems necessary for conventional metabolism. Potential fungi were determined to contribute to the trophic dynamics of freshwater environments. These organisms not only modulate aquatic productivity through an intermediary level of occupying in the food chain, but also indicate the environment status in a definite period. Furthermore, their range has grown in importance in recent years as a result of their specific species' ability to detect any decline in water quality as a result of contamination and eutrophication (Bassem, 2020).

Soil pollution also may affect habitat contamination when the presence of toxic chemicals like herbicides, pesticides, and fungicides in soil, which has caused a high risk to the ecosystem. Various compounds often enter the soil from the atmosphere, such as precipitation water, as well as from surface water sources and shallow groundwater flowing through the soil, as well as from wind movement or other forms of soil disturbances. When the amounts of soil contaminants exceed natural levels, pollution is generated (Osama et al., 2020).

The agricultural industry is very important for food production, but inefficient agricultural practices will continue to contribute to severe soil pollution. For example, copper has been widely used as a fungicide and its concentration can increase soil pollution to a higher level based on how this chemical is used by farmers. These chemicals are also used in animal feed and released into the atmosphere as fertilizer products that are spread through grasslands as well as other agricultural land. In addition, Cadmium is used as a fertilizer and a highly toxic metal in agricultural land. All these pollutants if not handled properly will then produce a lot of metals and natural pollutants from these organic materials such as sewage sludge, dirt, compost, and biowaste (European Environment Agency, 2020).

3.1.2 Health Risks and Issues

Polluted environmental problems can result in serious health risks for humans. The factors of clean air, unpolluted drinking water, nutritious food, and safe living all depend on its environment being well managed by the community. The increase in human population often causes unhealthy environmental problems and affects the health of the population as well as increased medical costs. This will result in a short human life span if the quality of the environment is affected.

Resnik and Portier (2015) stated in the article Environment, Ethics, and Human Health that although the environment can sustain human life, it can also be a source of disease if the environment is mismanaged. Human mortality rates can be high if basic medical needs are inadequate. The risk of cancer, heart disease, asthma, and various other diseases are all due to polluted environmental factors. Solid waste, toxic materials, and food contaminants, or hazardous are factors that can affect the quality of human health such as used vehicle batteries will be dumped on the ground and hydrogen gas can leak from batteries; potassium hydroxide, a caustic agent that can cause respiratory, eye, and skin irritations. If battery acid makes contact with your skin, it can create a skin reaction. Chemical burns can be the result. Unlike thermal burns caused by fire or heat, burns caused by batteries can quickly dissolve your skin. In addition, various infectious illnesses can be caused by contaminated drinking water, inadequate sanitation, and poor hygiene.

Exposure to agricultural chemicals is one of the human challenges in maintaining health and the environment. For example, pesticides play a critical role in improving agricultural yields while also posing health and environmental hazards. Alternatives to the use of pesticides result in the deterioration of human health. Avoiding the use of all pesticides will have a major impact on agricultural production, resulting in food shortages and higher food prices. To increase food production as well as minimize the use of harmful chemicals, public health authorities have chosen to control the use of pesticides and encourage innovation projects based on the idea of integrity in science and technology. These efforts can produce products and reduce human health risks. One of the serious health risks is cancer and this is

due to the agricultural sector such as chemicals, combustion smoke, contaminated drinking water, chemicals found in food, and poor air quality.

3.1.3 Industry Sectors

Private industry needs to have higher standards to prevent any kind of contamination. Most of industrial pollution can be directly linked worldwide. There are a number of forms of industrial pollution that can impact air quality, and it can enter the soil, causing widespread environmental problems. An article on environmental impacts of industrialization highlights four primary impact points: air, water, soil, and habitat (Fig. 3.1).

The most serious problem of the industrial sector is air pollution, caused by smoke or burning of fossil fuels but water pollution is also a serious issue in factory areas especially those built near natural water sources. These contaminants may be solid, liquid, or gaseous, and all have the potential to contaminate local water sources. Industrial sites can also result in soil pollution. Although lead contamination is the most common, heavy metals and other toxic chemicals can also get into the soil and contaminate agricultural crops. In addition, contaminated soil is also not suitable for use if it contains contaminated substances that are harmful to the health of organisms. The destruction of these habitats can disrupt local ecosystems and can result in the extinction of plants and animals if the species is unable to migrate or adapt to its new environment (Folk, 2018).

The harmful impacts of pharmaceutical manufacturing on the environment are well-known. However, because it is not fully managed, its toxic effects on animals

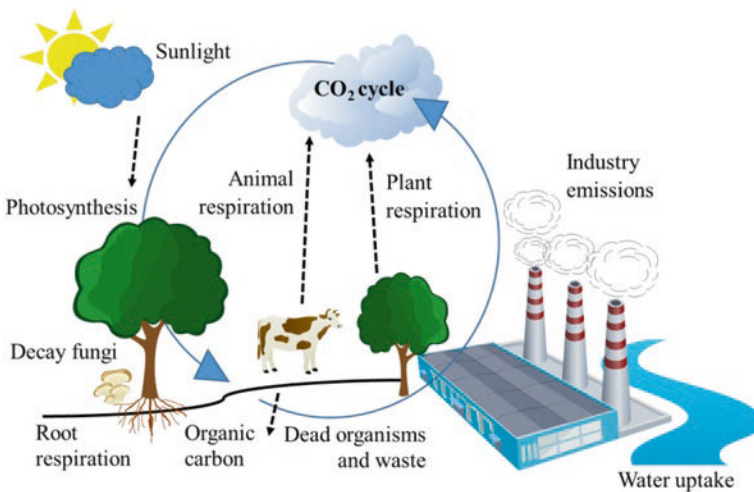


Fig. 3.1 The carbon cycles in natural ecosystem

and humans continue to linger until it is difficult to control. Pharmaceutical wastes are frequently released into the atmosphere. Throwing antibiotics into the environment poses a high risk of the level of resistance of various natural pathogens and pushes their strength for resistance to be treated if infected by humans or animals. It is clear that this global problem is often discussed in terms of pharmaceutical contamination. This problem will have a more significant impact especially on those living near water-processing plants and food supplies that have been contaminated with pharmaceutical products.

Animals, particularly fish that dwell in contaminated water, are negatively affected by pharmaceutical product contamination in the environment. Many rivers in Europe, for example, are home to intersex male fish that show feminine sexual traits, including female reproductive anatomy, according to a research published in the scientific journal *Nature* in 2009. Males can also create vitellogenin, a protein present in eggs that can be stimulated by hormones (Nawrat, 2018).

3.1.4 Communities

Environmental problems are the result of human decisions and activities that harm the environment. Individuals who act irresponsibly will make choices and participate in actions that affect the climate, such as dumping garbage, clearing forest areas for agriculture, roads, and factory development. Many companies, government agencies, and other organizations make decisions and participate in activities that have a significant negative impact on the environment. Individuals and organizations may be fully aware that their actions have damaged the environment, or they may be insensitive without thinking about the possible environmental consequences of their actions.

Humans can influence environmental problems such as the release of chemicals in soil and rivers. These include drain cleaners, laundry powder, paints, and pesticides, all things to avoid at any cost. It has been reported that arts and crafts materials as well as lawn care items can harm our environment and pollute the environment (Lindsey, 2017). The health risks from applying pesticides on crops or animals, drinking dirty water, inhaling polluted air, and being exposed to animal hormones and genetically modified organisms are all present in the lives of every family and farmworker today. Farmers are often affected by risks arising from disturbances to the environment, such as chemical pollution of water and air, sediments, and harmful odors. However, it gave rise to the idea to help them through innovative products that can reduce the low level of health of farmers such as the use of products that can absorb these chemicals. Environmental challenges to land production are more long-term and indirect, but they are still real threats to the viability of farms and farming (Ikerd, 1999).

3.2 The Role of White-Rot Fungi

White-rot fungi are a heterogeneous group that have an extraordinary function in breakdown components of plant cell wall and play an important role in global carbon cycling in forest ecosystems. Interestingly about 90% of all wood-degrading basidiomycetes species are white-rot type (Eriksson et al., 1990; Hatakka & Hammel, 2010). This group of WRF is the most efficient lignin-degrading organisms to convert it into carbon dioxide and water (Fig. 3.2). The WRF are often found on wood trees of angiosperm species and can cause nonselective or simultaneous damage or decay. Some basidiomycetes species can degrade lignin and leave the lighter-colored cellulose, or some of them degrade both lignin and cellulose in tree wood tree cells and this is called selective WRF (Eriksson et al., 1990).

Several previous studies have reported that the WRF are promising bioremediation agents because they have a high tolerance to toxic environment and the ability to transform a complex variety of aromatic pollutants and reduce their concentration in soil or water. In addition, they can also adapt to a wide range of temperatures, pH, humidity, aeration, and biotic factors (Nur et al., 2019). As a result, WRF have the ability to generate enzymes such as lignin peroxidase, manganese peroxidase, and laccase in different abiotic and biotic factors in order to be used in a wide range of mycoremediation or bioenergy applications (Cohen et al., 2002). Fig. 3.3

White-rot fungi potential is an economic resource, since its value may be exploited as a product in commercial biotechnology applications (Table 3.1). Mycoremediation technology, which has the ability to reduce various pollutants in



Fig. 3.2 The wild oyster mushroom (white-rot fungus), *Pleurotus ostreatus*, is grows on wood, and biodegradation processes can be seen based on discoloration and mechanical changes in their natural habitat of forest

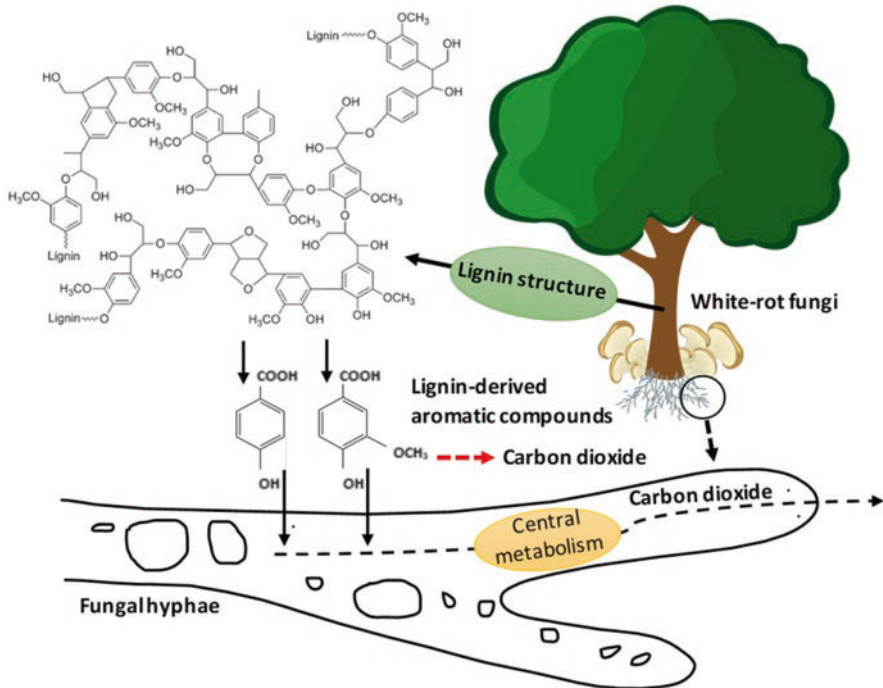


Fig. 3.3 The complex process of lignin-degrading fungi of white-rot fungi and convert it into carbon dioxide and water. (Modified Image from Davinia Salvachúa Rodríguez, National Renewable Energy Laboratory)

soils and sediments, including the dominant conventional explosives, DDT (dichlorodiphenyltrichloroethane), polynuclear aromatic hydrocarbons (PAH), and polychlorine biphenyl PCBs, can be applied to the entrepreneurial aspect of the WRF production process as a product.

3.2.1 Biodegradation Process by White-Rot Fungi

In general, WRF are responsible for the biodegradation of complex biomass, including lignin and a wide variety of environmental pollutants. There are several methods for reducing environmental pollutants and obviously, the WRF technology is different than other well-known methods of bioremediation such as bacteria. The advantage that WRF than bacterial systems are that they do not require preconditioning against certain pollutants. On the other hand, bacteria must be pre-exposed to pollutants to allow the enzymes to be induced and degrade the pollutants. The concentration of pollutants must also be present in significant amounts; otherwise, induction of enzyme synthesis will not occur by bacteria. In addition, the limited nutrients will

Table 3.1 Role of fungi in biotechnology processes of pollutants

| Organism (fungus) | Chemicals/pollutants | References |
|---|---|--|
| <i>Agaricus bisporus</i> , <i>Lactarius piperatus</i> | Cadmium (II) ions | Nagy et al. (2014) |
| <i>Allescheriella</i> sp., <i>Stachybotrys</i> sp., <i>Phlebia</i> sp. | Soil contamination | Mancera-López et al. (2008) |
| <i>Coriolus versicolor</i> | Polycyclic aromatic hydrocarbons | Jang et al. (2009) |
| <i>Fomes fasciatus</i> | Copper (II) | Sutherland and Venkobachar (2013) |
| <i>Flammulina velutipes</i> | Copper | Luo et al. (2013) |
| <i>Ganoderma</i> sp. | Textile dyes Yellow, Blue and Red Procion | Barreto et al. (2011) |
| <i>Jely</i> sp., <i>Schizophyllum commune</i> and <i>Polyporus</i> sp. | Malachite green | (Tsujiyama et al., 2013) |
| <i>Lentinula edodes</i> | 2,4-dichlorophenol Eucalyptus waste Barley straw, Vineyard pruning and Wheat straw | Brienzo et al. (2007), Gaitán-Hernández et al. (2006), and Lechner and Papinutti (2006) |
| <i>Lentinula tigrinus</i> | Wheat straw | Yogita et al. (2011) |
| <i>Pleurotus citrinopileatus</i> | Handmade paper and cardboard industrial waste | Kulshreshtha et al. (2013) |
| <i>Pleurotus eous</i> and <i>Lentinus connotus</i> | Paddy straw, sorghum stalk, and banana pseudostem | Rani et al. (2008) |
| <i>Pleurotus florida</i> | Handmade paper and cardboard industrial waste | Kulshreshtha et al. (2010) |
| <i>Pleurotus ostreatus</i> | Oxo-Biodegradable plastic Cadmium Extract from the sawdust | da Luz et al. (2013), Tay et al. (2011), and Akinyele et al. (2012) |
| <i>Pleurotus pulmonarius</i> | Radioactive cellulosic-based waste Crude oil | Eskander et al. (2012) and Olusola and Anslem (2010) |
| <i>Pleurotus sajor-caju</i> | Heavy metal Zn | Jibrán (2011) |
| | Indigo | Kamida et al. (2005) |
| <i>Pleurotus tuber-regium</i> | Heavy metals Nigerian trees; <i>Terminalia superba</i> , <i>Mansoniaaltissima</i> , <i>Holopteliagrandis</i> and <i>Miliciaexcelsa</i> Cotton waste, sawdust of <i>Khaya ivorensis</i> and rice straw | Oyetayo et al. (2012), Jonathan et al. (2008), and Kuforiji and Fasidi (2008) |
| <i>Pleurotus platypus</i> , <i>Agaricus bisporus</i> , <i>Calocybe indica</i> | Copper, Zinc, Iron, Cadmium, Lead, Nickle | Prasad (2013) |
| <i>Volvariella volvacea</i> | Banana leaves (<i>Musa sapientum lina</i>) Agro-industrial residues such as cassava, sugar beet pulp, wheat bran, and apple pomace | Belewu and Belewu (2005) and Akinyele et al. (2011) |

also react with white rot enzymes to perform the degradation process. Fungi continue to develop several survival mechanisms in limited and extreme conditions. These include glutathione *s*-transferases (GSTs) as an intracellular enzyme are also involved in stress responses and defense systems such as *Phanerochaete chrysosporium* (Morel et al., 2009).

The processes by WRF usually can be recognized when they are degrading the wood structures or any undesirable change in the properties of a wood material based on three stages: biodeterioration, biofragmentation, and assimilation (Luyt & Malik, 2019). They used wood as their source of carbon or nutrients, which will benefit other organisms when the particles of degradation process are distributed in soil. The biodeterioration process occurs on the polymer surface or inside the polymer, and it is also called a surface-level degradation that usually changes the mechanical, physical, and chemical properties of the wood material. This surface degradation occurs rapidly when it is exposed to abiotic factors (nonbiological agents) such as sunlight, temperature, and chemicals in the environment. The influence of abiotic factors allows the degradation process due to the weak wood material structure changes (mechanical), and potential WRF performance in a wide range of exposure conditions (Luyt & Malik, 2019; Lucas et al., 2008). Although biodeterioration is the first stage, it is possible that the process of biodegradation occurred parallelly with biofragmentation (Müller, 2005). The early stages of biodeterioration are often accompanied by a discoloration of the wood, which can be difficult to recognize until the late stages of biodegradation and are easily recognized by their changes in color and chemical properties.

The biofragmentation of complex polymers in the wood cell wall is a process in which the bonds in the polymer are cleaved, producing oligomers and monomers (Luyt & Malik, 2019). White-rot fungi carry out aerobic digestion, which uses oxygen to break down these polymeric materials and it produces carbon dioxide, water, some types of residues, and new biomass (Van Der Zee, 2011). Most of WRF are potentially used due to their aerobic digestive capabilities in waste management systems and as a local renewable energy source (Klinkner, 2014).

Meanwhile, the process of assimilation occurs from biofragmentation and the product is then integrated into microbial cells. The WRF used the mycelium to carry these products into the cell by membrane carriers. These biofragmentation products will enter the catabolic pathway, leading to the production of adenosine triphosphate (ATP) or elements of cell structure (Lucas et al., 2008).

The results of this biodegradation process can give ideas to researchers to produce potential products for entrepreneurs who want to overcome environmental problems. The ability of WRF to use wood material (substrate) as a food source with physical changes in the substrate is an important indicator. Also, potential in the mycoremediation of chemical pollutants when proven by changes in the chemical properties of wood material. This will provide an opportunity to produce many products related to mycoremediation to solve the problem of polluted environment.

3.2.2 Mycoremediation of Environmental Pollutants

Various types of environmental pollution have been produced in public areas including chemicals from various sectors such as agriculture, industry, and pharmacy. The mycoremediation process uses enzymes secreted by hyphae or mycelium, often producing a mass of fibers to whiten the substrate or wood at high levels of biodegradation. The process of developing this product can be combined with the use of potential WRF with engineering technology to clean up environmental pollution. The use of bead hydrogels is a technology that needs to be developed with a combination of mycelial fungi that have potential in mycoremediation. This technology was able to maintain the structure of the hydrogel and does not significantly reduce the quality of propagule within seven months. (Kuek et al., 1992).

The ability to use wood material (substrate) as a food source by WRF by producing physical changes to the substrate is an important idea for producing new products.

3.2.3 Capsule Technology of WRF Product

Hydrogel products constitute a group of polymeric materials, the hydrophilic structure of which renders them capable of holding large amounts of water in their three-dimensional networks. In another definition, hydrogel is a polymeric material that exhibits the ability to swell and retain a large amount of water in its structure (Ahmed, 2015), but will not dissolve in water (Ahmed, 2015; Oyen, 2014). Due to their cross-linked polymer networks, hydrogels exhibit elastic solid properties with deformation and softness (Liu et al., 2020; Hong et al., 2008).

Hydrogels can be classified into various groups based on their: physical structure (amorphous, semicrystalline, hydrogen bonded, or supramolecular); electric charge (ionic (charged or neutral); crosslink (physically or chemically cross-linked); responses to external effects (stimulus-sensitive and insensitive ones); and origin (synthetic and natural) (Deligkaris et al., 2010).

Extensive employment of hydrogel products in a number of industrial and environmental areas of application is considered to be of prime importance (Ahmed, 2015). Hydrogels have been widely used in applications that closely interact with biological organisms, such as tissue engineering, drug delivery, and biological research (Liu et al., 2020). Moreover, the unique properties of hydrogels, such as superior softness, wetness, responsiveness, biocompatibility, and bioactivity (Liu et al., 2020), indeed suggest the possibility of their crucial functions in devices and machines, such as sensors, actuators, coatings, optics, electronics, and water harvesters (Liu et al., 2020; Gerlach & Arndt, 2010).

The hydrogels for agriculture can be used as a water-repellent granule product as it is capable of swelling from its original size many times when in contact with water. It has been widely proposed for use in agriculture with the aim of improving root conditions as well as supplying water to crops, by improving the water-retaining

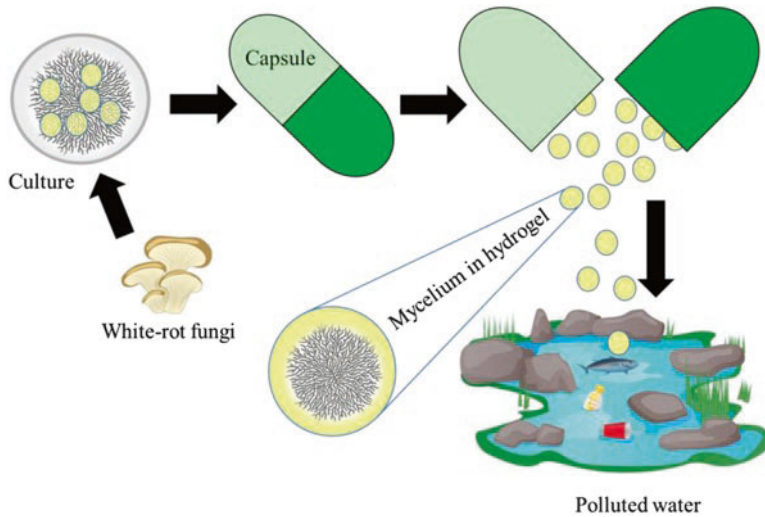


Fig. 3.4 Inoculated hydrogel by mycelium as a model product for decontamination of contaminated habitats

properties of growing media (soil or substrate without soil). Moreover, low price, high durability, and stability in swelling environment and during storage, gradual biodegradability without formation of toxic species characters encouraged scientists to apply hydrogel system in agriculture (Neethu et al., 2018).

According to the above definition, one of the emerging fields for both scientists and entrepreneurs is inoculation of hydrogels by mycelium in capsule format. This can solve the problem of pollution in high-risk areas such as contaminated wastewater pollution areas, which are close to pharmacies, food industry factories, and others. The type of hydrogel capsule inoculated with mycelium is able to retain the mycelium for a long-time during storage and control its release (i.e., once exposed to the wastewater) (Fig. 3.4).

3.2.4 *Cleaning and Removal Contaminations Product*

White-rot fungi have the ability to degrade the contaminated liquid, soil, sediments and demonstrated their ability to survive in extreme environment conditions (Stella et al., 2017). Tišma et al., 2010 investigated the possibility of degradation of the high polyphenolic compounds, dyes, and other xenobiotics. Also, WRF have a great range of different enzymes such as hydrolytic enzymes (cellulase, pectinase, xylanase) (Tišma et al., 2010; Teerapatsakul et al., 2007). Most of these enzymes are industrially important and have great potential in the processes of bioremediation, biodegradation, biopulping, degradation, and detoxification of recalcitrant substances (Tišma et al., 2010) Fig. 3.5.



Fig. 3.5 Summary of WRF role in the removal of contamination

A previous study by Stella et al. (2017) examined the ability of *Pleurotus ostreatus* and *Irpex lacteus*, as WRF, to degrade polychlorinated biphenyls (PCBs) in actual contaminated soils with varying chemical characteristics and autochthonous microflora. *Pleurotus ostreatus* had the best results, with PCB removal rates of 18.5, 41.3, and 50.5 percent from the bulk soil, top (surface), and rhizosphere after 12 weeks of treatment, respectively. They concluded that the extensive PCB degradation performance in a relatively short period of time and the exceptional competitiveness of *P. ostreatus* make its application in the remediation of soils contaminated with PCBs potentially transferred to a larger scale (Stella et al., 2017).

Polluted waters with phenol are formed from the production of olive oil in Mediterranean region (Gianfreda et al., 2006). Olive mill wastewater is highly toxic effluent obtained from the extraction process by the olive oil industry. Numerous studies were done on using WRF in the treatment of phenols and related compounds (Laconi et al., 2007; Justino et al., 2009, 2010; D'Annibale et al., 2006; Lu et al., 2009; Bollag et al., 2003; d'Acunzo et al., 2002). For instance, Justino and coworkers in 2009 and 2010 reported the efficiency of three different approaches on phenols removal from olive mill wastewater (Justino et al., 2009, 2010). They oxidized different types of phenols that were extracted from Olive mill wastewater via biological (i.e., treatment with two fungal species *Trametes versicolor* and *Pleurotus sajor*), enzymatic, and chemical treatment. They found that chemical treatment by photo-Fenton was more efficient than biological or enzymatic treatments.

3.3 Development of Potential WRF Product to Entrepreneurs

3.3.1 Natural Environment and Municipal

The municipal management organizations might be interested based on WRF product, such as mycobead to land and water pollution issues. This product can be used as a bioremediation to chemicals. It is clear that polluted chemicals that dissolved in

water cannot be treated by physical and chemical methods but rather by biological treatment. The efforts to produce WRF products in interaction with contaminants through bio-mediated and engineering techniques are required. Research and development of these products can benefit countries that produce pollutants, improve environmental policies and manage with appropriate procedures, future regulations, address bioremediation targets, identify pollutant availability, and potential threats to natural ecosystems and human health. This is considered an excellent effort for the country to protect us from pollution in natural systems and the level of threat to human health caused by various human activities.

3.3.2 Agriculture Industry

Isolation of potential species from the soil can benefit the agricultural industry and increase yields. This is through technology and products produced such as hydrogels, which can absorb chemicals such as herbicides or pesticides using the mycelial structure of the fungi. These chemicals can be converted to carbon dioxide and water by fungi. Toxins that are considered waste or used excessively can become pollutants in the soil and these will be absorbed by agricultural crop products and pose a risk to its users.

This innovative mycobeat product can also be used to store potential fungi for at least 7 months without losing capacity as propagules (Kuek et al., 1992). A number of these hydrogel products are also used by fungi that can function in symbiosis with plants.

3.3.3 Food Industry

As mentioned earlier in the previous section of cleaning and removal contaminations product, WRF can help to treat the contaminated soil, water, and sediments. It is suggested to the entrepreneurs to build the greenhouse and to use the WRF near to the contaminated wastewater. This strategy can fulfill two goals: (a) application of contaminated wastewater and (b) producing WRF for human consumption. In other words, eliminate the contaminated wastewater while producing a new product (Fig. 3.6).

3.3.4 Pharmaceutical Industry

Waste generated by the pharmaceutical industry has harmful implications for the environment and public health if disposed of untreated (Shah & Shah, 2020). Most of these wastes are released into the environment at low concentrations through routine pharmaceutical use, damage and expiration. The treatment performed by

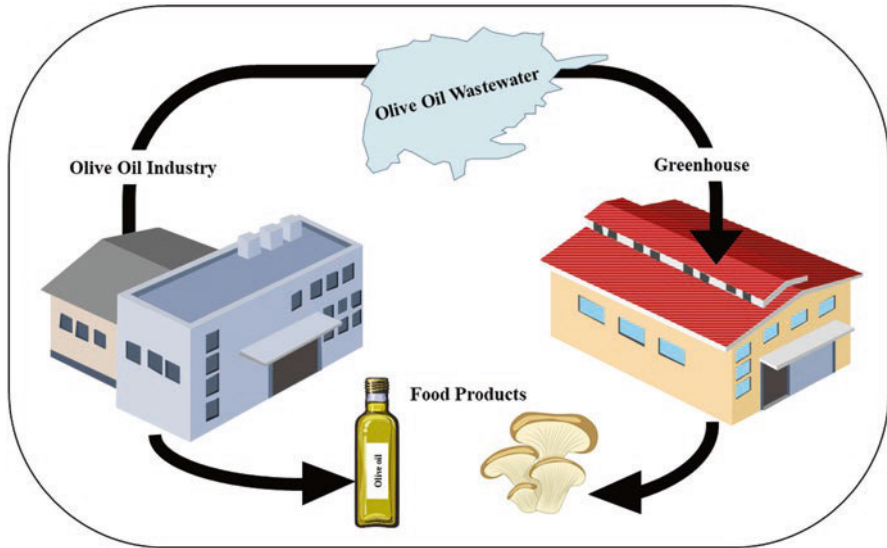


Fig. 3.6 Development of potential WRF product by entrepreneurs in the food industry

WRF can offer an environmentally friendly platform for the removal of pharmaceuticals and personal care products (PPCP) from wastewater as PPCP may have adverse effects on aquatic organisms as well as humans. This will be of great interest to the water industry. It has been reported that intact cell WRF or their extracellular lignin-converting enzyme (LME) can degrade PPCP efficiently. Treatment of PPCP mediated by WRF depends on several factors including the physicochemical properties of PPCP (e.g., hydrophobic and chemical structure) and wastewater matrix (e.g., pH, temperature, and dissolved constituents), the type of WRF species, and specific extracellular enzymes (Asif et al., 2017).

3.4 Economic Value of Mycoremediation Product in the Market

3.4.1 Benefit to Researchers and Entrepreneurs

Bioentrepreneurship is an activity that has a scientific element in the field of entrepreneurship. This is the most recent knowledge shift being discussed by academics toward industry. Biotechnology entrepreneurship is greatly aided by experienced academic research. Bio-entrepreneurship is all the activities required to set up a business that develops, manufactures, and markets biotechnology products to solve research problems. We can see that bio-entrepreneurship is more advanced now, where the biotechnology sector is producing real-world results that enable

technological advancement and integration with other fields. Also, it contributes to environmental conservation, disease research, new biomedical tools, product refining using biosynthetic processing, biofabrics and biofuels, and more (Afandi, 2018).

A biotechnology company is usually started by researchers to produce a potential product. This will provide and open up many business and employment opportunities. Through the manufacture of new hydrogel bead products, bioentrepreneurs will have a lot of assessment data to improve the quality and effectiveness of the product for the future. Products will also generate profits for entrepreneurs and product creators. In this way, technology not only benefits the environment but also society. These include ideas, the basics of biological science and materials engineering technology can be improved.

3.4.2 Income of Country

The modern transition of health, food, agriculture, and bio-based industries is referred to as bioeconomy. Pharmacy, biofuels, farming, enzymes, atmosphere and climate, bio-based polymers, biorefining, and well-being and nutrition are all examples of biotechnology assets (Fig. 3.7).

The invention products of white-rot fungi that able to be marketed and used to clean environment issue are very crucial to help the environmental management

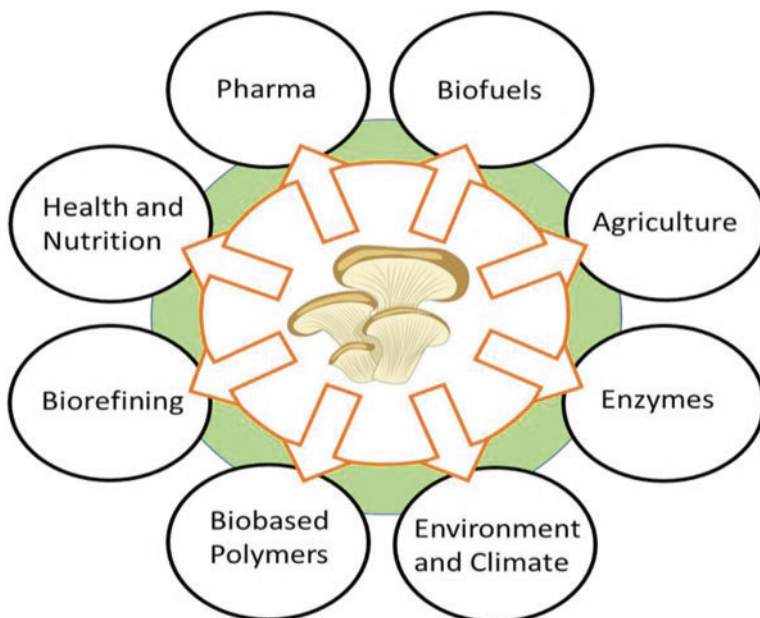


Fig. 3.7 The branch of biotechnology in the field of mushrooms

system. Products used to control or recover toxic waste are beneficial to society and the economy. In the twenty-first century, strategies to clean up toxic wastes and pollutants using affordable and efficient low costs are underway. Hydrogel bead products with capsules can actually provide a high opportunity to the income of the country, because now many environmental areas are polluted. The future of produced products will help all scientists to explore more opportunities to produce the best products. This will also benefit entrepreneurs as they will be able to market the product to the rest of the world. This will generate the income of all industries involved, including increasing economic income.

3.5 Conclusions

The environmental issues are the result of human decisions and their inappropriate activities that harm the environment implies within the communities. White-rot fungi, which are the most efficient lignin-degrading organisms to convert pollutants into carbon dioxide and water, are promising bioremediation agents. They have a high tolerance to toxic environment and the ability to transform a complex variety of aromatic pollutants and reduce their concentration in soil or water. In general, WRF are responsible for the biodegradation of complex biomass, including lignin and a wide variety of environmental pollutants. The degradation of surface structure occurred rapidly when it is exposed to abiotic factors. The integrated development of the use of hydrogel bead and mycelium fungi are potentially in mycoremediation and this technology does not significantly reduce the quality of propagule within seven months. This biodegradation process can be used by researchers to manufacture products that will benefit some entrepreneurs to solve problems related to environmental pollution. It will provide an opportunity for more mycoremediation-related products to be produced with the properties of WRF. Interestingly, hydrogel capsules inoculated with mycelium can maintain the mycelium for a long time during storage. The effort to produce WRF product in interaction with contaminants through the most appropriate bioremediation techniques and other relevant techniques are needed. These are considered very good efforts to protect us from contaminants in our environment caused by various human activities. Therefore, the WRF process can regenerate energy and save the environment. It is suggested to the entrepreneur in the food industry area to use the WRF product near the contaminated wastewater of the food and pharmaceutical factory to refine the wastewater.

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Chapter 4

Production, Partial Optimization, and Characterization of Keratinase Enzyme by Fungal Species Isolated from Soil of Bhopal



Renu Mishra and Shraddha Tamrakar

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4.1 Introduction

Soil encourages a large number of microorganisms and is one of the highest complicated microbial habitats that permit bacteria, fungi, and other micro-organisms to nurse their entire life. Soil is the shelter for several fungi, which are not noticed and explored in the environment (Pahare et al., 2018).

Keratinophilic fungi are a little, all around characterized and essential gathering of growths that colonize diverse keratinous substrates, and generate them to fragments of low atomic weight. These fungi are available on the earth with variable dispersion designs, which rely upon components, for example, human or animals (Kumar et al., 2013).

Keratinophilic means keratin-loving micro-organism. Keratinophilic fungi are natural colonizers of keratinous substrates. Few of them are keratinolytic and play a massive ecological character in the decay of insoluble fibrous proteins, keratins (Kushwaha & Guarro, 2000). Keratin has a place with the exceptionally heterogeneous group of fibrillar proteins of conservative structure. Countless extensions of disulfide bridges in keratins structure altogether impact its properties, specifically, high mechanical and substance obstruction. Keratin is highly stable and water insoluble. Common proteases like pepsin and papain are not capable of degrading keratin (Staroń et al., 2011).

Keratinase is a specific class of extracellular proteolytic inducible chemical with the ability of debasing insoluble keratin substrates. It is vital for hydrolyzing hair, plume, and collagen in the sewage framework amid wastewater treatment.

Human hair and nails are always in interaction with pollutants in the environment and waste exposed to fungal and bacterial contamination (Alghamdi et al., 2018). Recent studies have become an important clinical condition, which require attention to public health, because some of them are seriously harmful to human health. Health authorities must build up their health information campaigns, which cover not only serious disease prevention and treatment but also environmental hygiene.

4.1.1 *Keratinases: Applications*

Keratinolytic micro-organisms and keratinases may be utilized for multiple applications in a myriad of industries, like the detergent industry, leather production, food and feed industries, medical and pharmaceutical industries, waste management industries, etc. Compared to traditional microbial proteases, production of keratinases can be achieved at lower costs as the organisms are able to utilize the abundant keratinous waste as nitrogen and carbon source. Since elevated production cost is one of the main issues in the industrial production of enzymes, this methodology of utilization of abundant quantities of keratinous waste will make the process comparatively more economical while also increasing its application portfolio.

4.1.1.1 Detergent Industry

Proteases are widely utilized in the detergent industry as a safe alternative replacing harmful chemicals such as caustic soda. In addition to being used in washing detergents, keratinases may also be used as an additive in lens cleaning solutions, drain cleaning solutions, etc. (Nasaimento & Martin, 2006).

4.1.1.2 Leather Industry

The leather industry involves several steps of leather treatment with chemicals, such as sodium sulfide, sodium hydroxide, and lime. These chemicals are potent water pollutants along with solid wastes emerging due to these treatments (Sivakumar et al., 2012). Processing of leather with enzymes such as proteolytic, lipolytic, and glycolytic enzymes has emerged as a safe alternative (Bihari et al., 2010).

4.1.1.3 Food and Feed Industry

The National Research Development Corporation (NRDC), India, has recently utilized bifunctional chimeric keratinase from *Bacillus* sp. for the purpose of debittering amino acids in the processing of soy sauce. It was observed that this increased the taste of tea, and feather meal preparation along with other applications (www.nrdcindia.com). Usually, enzymes are useful in feed industries because of their part in the breakdown of the antinutritional elements in feed ingredients. In addition to that, this keratinase treatment improves the bioavailability of biomolecules in animal feed. It does so by assisting endogenous proteolytic enzymes of feeding animals to unwind the rigid structure of feed. However, often, this is restricted due to the high production costs, and therefore, low-cost keratinases can prove to be a useful alternative as a feed additive. Recently, many commercial enzymes, such as Versazyme, and Valkerase, etc., have been used as applications in animal feed preparations.

4.1.1.4 Fertilizer for Organic Farming

Composting of keratinous wastes using keratinolytic micro-organisms leads to gradual release of nitrogen fertilizers. This makes it a viable application in organic farming. This methodology has become one of the safest and most cost-effective technologies to resolve keratinous waste materials.

4.1.1.5 Biofuel Production

Poultry waste can be used as a viable source of biofuel. Keratinolytic micro-organisms and their keratinases can be employed in the utilization of these wastes for generation of natural gas (Puhl et al., 2009), methane gas fuel pellets (Gushterova et al., 2005), and biohydrogen (Saha et al., 2013).

4.1.1.6 Cosmetic and Pharmaceutical Applications

Non-collagenolytic keratinases have emerged as promising biocatalysts in the pharmaceutical as well as cosmetic industries. They have been illustrated as a useful ingredient in depilatory compositions for hair shaving formulations and skin lightening agents (Yang, 2012). Certain crude keratinases have also been demonstrated to elevate hair characteristics like weight, flexibility, brightness, softness, and strength; this illustrates that these can be applied in the creation of hair care products (Cao et al., 2012). Additionally, keratinases have shown the ability to degrade thickened layers of dead skin (hyperkeratosis) found around the toes and fingers, thereby promising to be an alternative to the conventional salicylic acid used for the treatment of such predilections (Gupta & Ramnani, 2006). In the same fashion, keratinases can be capable of peeling skin to remove acne, which is caused by the blockage of the sebaceous gland by keratins (Selvam & Vishnupriya, 2012).

4.2 Materials and Methods

4.2.1 Collection of Soil Samples

Soil samples of keratin containing waste were collected in polythene bags from the different sampling sites, brought to the laboratory, and analyzed.

4.2.2 Isolation of Keratinophilic Fungi

For the isolation of fungi, hair baiting technique was used (Vanbreuseghem, 1952).

4.2.3 Morphological Microscopic and Molecular Identification of Keratinophilic Fungi

Firstly, the fungi were indentified on the basis of morphological characteristics of the colony and microscopic examination. Molecular identification of isolated microorganisms was confirmed by the sequence of ITS1, 508 s, and ITS2 region of their DNA sequences.

4.2.4 Optimization of Culture Condition for Mass Production of Keratinase Enzyme

To optimize the culture conditions for mass production of keratinase enzyme by isolated soil microorganisms, a basal medium containing insoluble keratin was prepared with various pH (6, 7, 8, and 9) incubated on different temperatures (40, 45, 50, 55, and 60 °C) for various incubation periods (12, 15, 18, and 21 days) (Table 4.1).

4.2.5 Enzyme Production

The keratinase enzyme production was carried out in the basal medium by using a shaking flask. To begin the process, 5 mg of azokeratin was added in the basal medium: 2.5 ml of the all isolated fungal inoculums was added in 50 ml of medium and cultured on a rotary shaking incubator at 150 rpm and 37 °C for 72 h. After incubation, fermented broth was centrifuged at 5000 rpm for 20 min at 4 °C. The cell-free supernatant was collected and used for the assay of keratinase activity (Sivakumar et al., 2012).

Table 4.1 Composition of basal medium for keratinase production

| S. no. | Component | Gm/L |
|--------|--------------|------|
| 1 | Meat extract | 1 |
| 2 | Peptone | 1 |
| 3 | NaCl | 0.5 |
| 4 | Keratin | 5 |

4.2.6 Keratinase Assay

Keratinase activity was determined by taking 20 ml of 0.1 M Tris buffer (pH 8) containing 0.1% keratin and 40 μ l of enzyme solution and was incubated for 30 minutes at 55 °C. The reaction was stopped with 500 μ l of 0.1 mol-1 trichloroacetic acid (TCA) in 0.1 mol-1 Tris buffer, pH 8. The amount of protein present was assayed by the Lowry method using bovine serum albumin (BSA) as standard. The color developed was read at 660 nm (Jin et al., 2017).

4.3 Results

4.3.1 Morphological Characterization

Morphological characterization was performed and further confirmed by the National Center for Fungal Taxonomy (NCFT), New Delhi. All positive keratinophilic fungi were identified by colony morphology and lactophenol cotton blue microscopic staining method (Table 4.2).

Table 4.2 Morphological and microscopical characterization of isolated keratinophilic fungi

| S. no. | Sample ID | Species (morphological basis) | Color of mycelia | Pigmentation | Spore arrangement | Sporulation |
|--------|-----------|-------------------------------------|------------------|----------------|-----------------------|-----------------|
| 1 | S1 | <i>Fusarium solani</i> | White cottony | Pink | Long branched | Sexual |
| 2 | S2 | <i>Curvularialunata</i> | Blackish | Greenish brown | Branched septa | Sexual |
| 3 | S3 | <i>Chrysosporium keratinophilum</i> | Cream | Pale | Long lateral branched | Asexual |
| 4 | S4 | <i>Acremonium kiliense</i> | Cream | Colorless | Branched | Sexual/ asexual |
| 5 | S5 | <i>Aspergillus Niger</i> | Black | Pale | Septet | Sexual |
| 6 | S6 | <i>Acremonium restrictum</i> | White orange | Not appear | Septet | Sexual/ asexual |
| 7 | S7 | <i>Absidia californica</i> | Grayish brown | Colorless | Branched | Sexual |
| 8 | S8 | <i>Absidia repens</i> | White | Pale gray | Branched | Sexual |
| 9 | S9 | <i>Penicillium chrysogenum</i> | White | Colorless | Highly branched | Asexual |
| 10 | S10 | <i>Gibberella pulicaris</i> | Bluish | Light pink | Long branched | Sexual |
| 11 | S11 | <i>Aspergillus flavus</i> | Greenish | White | Branched | Sexual/ |

4.3.2 Molecular Identification of Fungi

The genomic DNA was isolated from selected keratinophilic fungi using the CTAB method. The DNA was visualized on agarose gel and quantified for an accurate amount of DNA isolated. We have observed an average of 2.86 mg/ μ l genomic DNA from all major keratinophilic fungi.

The gene-specific primers were designed using coding sequence available at National Centre for Biological Information (NCBI). The primers were designed based on in silico PCR validations. Primers are supplied in lyophilized form and dissolved in recommended volume of sterile water or TE buffer result in 100 μ M concentration.

The 100 μ M concentration primers are used for master stock and aliquoted into 10 μ M for PCR reactions. The targeted genes were amplified using thermal cycler and PCR reaction mixture. Different primers are required for identification of different set of genes in PCR Programme.

The primer may also vary in its melting temperature, and hence T_m values were pre-calculated and PCR program was designed based on T_m value for efficient annealing. The Tag Polymerase, a thermostable DNA polymerase, had a chain elongation capacity, 500 bases per second, and hence PCR program was designed based on gene length. Here, we have run several individual reactions and PCR trials for gene amplification and amplified gene was visualized on agarose gel with a standard DNA ladder. The primer used in the study was 550 bp in length.

The amplification products were sequenced for species identification. The DNA sequences of isolated strains were recited using basic local alignment search tool (BLAST) for species identification. The BLAST search showed that the sequence data of the isolated strain S1 shared 99% similarity with *Fusarium solani* and S2 shared 99% similarity with *Curvularia lunata*. The three isolated strains, identified as *Gibberella pulicaris*, *Chrysosporium keratinophilum* and *Penicillium chrysogenum* were S10, S3 and S9. DNA sequences of S5 and S11 showed 90% matching with *Aspergillus niger* and *A. flavus*. Similarly, S7 and TS8 fungi found close similarity with *Absidia* species. Amplification of each primer pair was tested on a panel of strains representing *Acremonium kiliense* and *A. restrictum*, along with the genomic DNA of the isolate from which the library was generated.

All isolated strains were submitted in the Gen-Bank databases for the purpose of phylogenetic study. The gene bank accession numbers for fungi from S1 to S11 are shown in (Fig. 4.1).

4.3.3 Phylogenetic Analysis

In the phylogenetic tree (Fig. 4.1), all major keratinophilics were reported to be of a distant origin and do not have any close resemblance. The phylogenetic tree was constructed by the neighbor-joining method and Jaccard coefficient. There are three

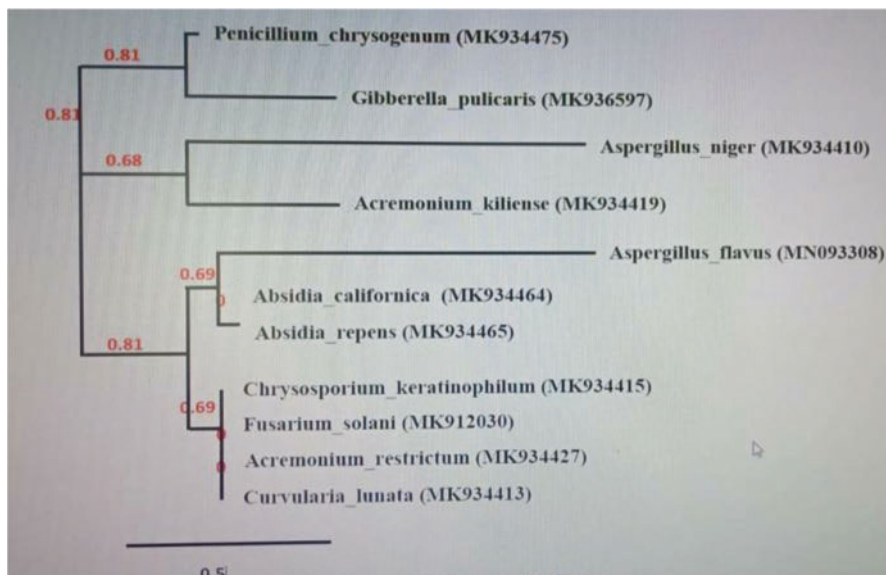


Fig. 4.1 Phylogenetic tree of the major keratinophilic fungi

ancestors of isolates found in this study. Group one has two species, that is, *Penicillium chrysogenum* and *Gibberella pulicaris*. *Aspergillus niger* and *Acremonium* are in another group, whereas the third has two subgroups, one including *Aspergillus flavus*, *Absidia californica*, and *Absida repens*. The other subgroup includes *Chrysosporium keratinophilum*, *Fusarium solani*, *Acremonium restrictum*, and *Curvularialuntana* (Fig. 4.1).

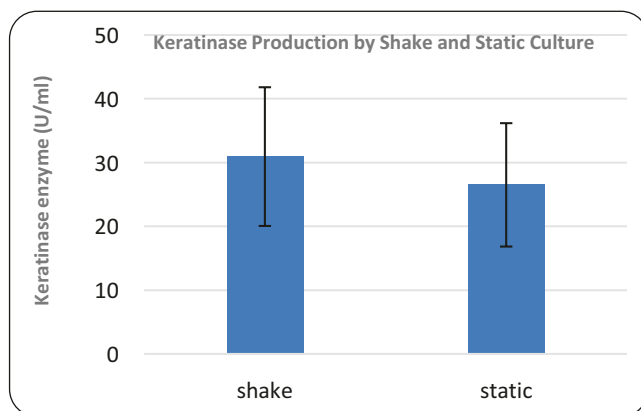
4.3.4 Optimization of Culture Media for Keratinase Enzyme Production

4.3.4.1 Effect of Fermentation on Keratinase Enzyme

Maximum keratinase production was recorded in the shake culture method compared to static culture. Table 4.3 and Fig. 4.2 show the results of keratinase production by both techniques. *Curvularialunata* produced a maximum 43.5 U/ml keratinase while minimum 17.5 U/ml was observed in the culture medium containing *Aspergillus flavus* during the shake culture method. The highest amount (38.6 U/ml) of keratinase was produced by *Fusarium solani* and minimum 14.6 U/ml produced by *Gibberellapulicaris* during static culture technique. Data represent the mean and \pm standard deviation for different methods. Although Table 4.3 and Fig. 4.2 show varied keratinase production in the two different methods, this

Table 4.3 Optimization of culture media for keratinase production by shake and static culture sample ID

| Sample ID | Species | Enzyme (U/ml) in shake culture | Enzyme (U/ml) static culture |
|------------------|------------------------------------|--------------------------------|------------------------------|
| S1 | <i>Fusarium solani</i> | 41.6 | 38.6 |
| S2 | <i>Curvularialunata</i> | 43.5 | 28.7 |
| S3 | <i>Chrysosporiumkeratinophilum</i> | 34.1 | 30.2 |
| S4 | <i>Acremonium kiliense</i> | 40.1 | 36.2 |
| S5 | <i>Aspergillus Niger</i> | 20.5 | 18.4 |
| S6 | <i>Acremonium restrictum</i> | 22.1 | 19.6 |
| S7 | <i>Absidia californica</i> | 19.6 | 16.9 |
| S8 | <i>Absidia repens</i> | 42.3 | 38.4 |
| S9 | <i>Penicillium chrysogenum</i> | 39.1 | 34.5 |
| S10 | <i>Gibberellapulicaris</i> | 19.8 | 14.6 |
| S11 | <i>Aspergillus flavus</i> | 17.5 | 15.4 |
| Mean ± SD | | 30.9 ± 10.8 | 26.5 ± 9.6 |

**Fig. 4.2** Keratinase production by shake and static culture (error bars show 1 st dev)

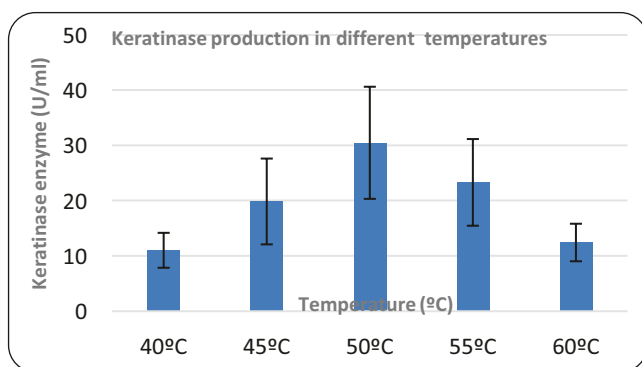
difference, however, is not statistically significant (p is 0.32 at t-test static value) and t-test is calculated using “Microsoft Excel Version 2019.”

4.3.4.2 Effect of Temperature on Keratinase Enzyme Production

Different temperatures, such as 40 °C, 45 °C, 50 °C, 55 °C, and 60 °C, were tested to obtain the optimum temperature for the maximum production of keratinase by all isolates. Out of the temperatures tested as above, the maximum production of keratinase was observed to be at 50 °C (Table 4.4 and Fig. 4.3). Maximum keratinase (43.5 U/ml) at optimum temperature 50 °C was produced by *Penicillium chrysogenum*, while minimum keratinase (5.9 U/ml) was recorded at 40 °C temperature

Table 4.4 Effect of temperature on keratinase enzyme production U/ml

| Sample ID | Species Temp (°C) | 40 °C | 45 °C | 50 °C | 55 °C | 60 °C |
|------------------|------------------------------------|-------------------|-------------------|--------------------|-------------------|-------------------|
| S1 | <i>Fusarium solani</i> | 8.6 | 12.9 | 19.4 | 11.8 | 7.9 |
| S2 | <i>Curvularialunata</i> | 10.3 | 14.6 | 22.3 | 15.9 | 9.4 |
| S3 | <i>Chrysosporiumkeratinophilum</i> | 9.9 | 14.9 | 23.5 | 14.9 | 8.6 |
| S4 | <i>Acremonium kiliense</i> | 12.4 | 24.8 | 32.1 | 34.2 | 14.6 |
| S5 | <i>Aspergillus Niger</i> | 13.3 | 28.5 | 35.1 | 28.4 | 14.2 |
| S6 | <i>Acremonium restrictum</i> | 11.4 | 29.1 | 40.1 | 28.9 | 18.4 |
| S7 | <i>Absidia californica</i> | 14.6 | 28.7 | 40.3 | 27.5 | 16.2 |
| S8 | <i>Absidia repens</i> | 14.9 | 21 | 41.7 | 30.1 | 11.4 |
| S9 | <i>Penicillium chrysogenum</i> | 13.8 | 24.9 | 43.5 | 30 | 14.8 |
| S10 | <i>Gibberellapulicaris</i> | 6.2 | 10.3 | 19.8 | 18.6 | 11 |
| S11 | <i>Aspergillus flavus</i> | 5.9 | 8.8 | 17.5 | 16.2 | 10.3 |
| Mean ± SD | | 11.0 ± 3.1 | 19.8 ± 7.7 | 30.5 ± 10.1 | 23.3 ± 7.8 | 12.4 ± 3.4 |

**Fig. 4.3** Keratinase production in different temperature

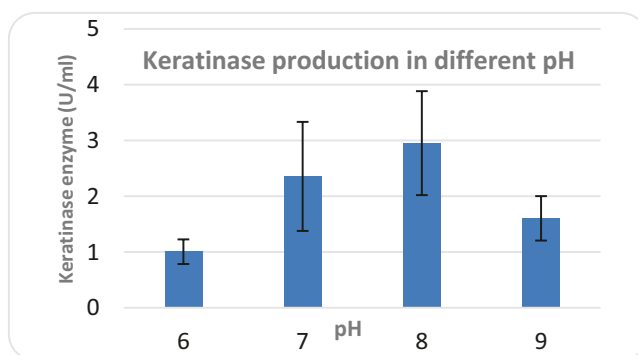
produced by *Aspergillus flavus*. Proteolytic activities of most organisms are seen to be at 30–37 °C, whereas some keratinolytic microbes show keratin-degrading capacities at higher temperatures. A nonparametric Kruskal-Wallis test applied on the data (the data fulfill the requirements for the test) shows that there was a statistically significant difference in the keratinase activity under different temperatures studied ($p < 0.0001$ for the test static $H = 31.08$).

4.3.4.3 Effect of pH on Keratinase Enzyme Activity

The medium adjusted on different pH values, (such as pH 6, 7, 8, and 9) was assayed to obtain the optimum pH for the maximum production of keratinase by all isolated keratinophilic microbes. From Table 4.5, it is clear that *Penicillium chrysogenum* is found to be the maximum enzyme-producing microorganism, which produced

Table 4.5 Effect of pH on Keratinase Enzyme Activity of Keratinophilic Fungi (U/ml enzyme produced)

| Sample ID | Species pH | 6 | 7 | 8 | 9 |
|---------------------------------|------------------------------------|---------------------------------|---------------------------------|---------------------------------|---------------------------------|
| S1 | <i>Fusarium solani</i> | 0.620 | 1.010 | 1.910 | 0.960 |
| S2 | <i>Curvularialunata</i> | 0.920 | 1.920 | 2.110 | 1.640 |
| S3 | <i>Chrysosporiumkeratinophilum</i> | 0.940 | 1.840 | 2.210 | 1.680 |
| S4 | <i>Acremonium kiliense</i> | 1.160 | 2.980 | 3.970 | 1.890 |
| S5 | <i>Aspergillus Niger</i> | 1.210 | 3.010 | 3.940 | 1.760 |
| S6 | <i>Acremonium restrictum</i> | 0.960 | 2.670 | 3.210 | 1.540 |
| S7 | <i>Absidia californica</i> | 1.280 | 3.240 | 3.860 | 2.140 |
| S8 | <i>Absidia repens</i> | 1.190 | 2.970 | 3.390 | 1.790 |
| S9 | <i>Penicillium chrysogenum</i> | 1.210 | 3.910 | 4.010 | 2.050 |
| S10 | <i>Gibberellapulicaris</i> | 0.840 | 1.230 | 2.010 | 1.130 |
| S11 | <i>Aspergillus flavus</i> | 0.720 | 1.120 | 1.840 | 1.050 |
| Mean \pm SD | | 1.0 \pm 0.2 | 2.3 \pm 0.9 | 2.9 \pm 0.9 | 1.6 \pm 0.4 |

**Fig. 4.4** Keratinase Production in different pH

4.01 U/ml keratinase at 8pH at 50 °C optimum temperature. pH 6 reduced the proteolytic activity of all isolates. The lowest enzyme (0.62 U/ml) production was found at pH 6 by *Fusarium solani*. There was a statistically significant difference in the keratinase activity under different pH values studied ($p = 0.00001$ for the test static $H = 29.25$) (Fig. 4.4) (Bewick et al., 2004).

4.3.4.4 Effect of Incubation Time

The effect of incubation period on keratinase production from keratinophilic fungi was studied for the incubation period from 12 days to 21 days as shown in Table 4.6 and Fig. 4.5. It was observed that the maximum enzyme production was attained between 15–18 days of incubation period. Incubation beyond the optimum time showed a rapid decline in the enzyme yield, as compared to the maximum (39.8 U/ml)

Table 4.6 Effect of incubation period on keratinase production

| Sample ID | Harvesting time species | 12 days /U/ml | 15 days /U/ml | 18 days /U/ml | 21 days/U/ml |
|------------------|------------------------------------|-------------------|-------------------|-------------------|-------------------|
| S1 | <i>Fusarium solani</i> | 19.8 | 39.8 | 38.4 | 30.6 |
| S2 | <i>Curvularialunata</i> | 18.6 | 24.3 | 27.6 | 26.4 |
| S3 | <i>Chrysosporiumkeratinophilum</i> | 17.2 | 20.1 | 29.8 | 27.4 |
| S4 | <i>Acremonium kiliense</i> | 18.7 | 35.6 | 34.6 | 30.2 |
| S5 | <i>Aspergillusniger</i> | 11.2 | 21.6 | 21.9 | 14.6 |
| S6 | <i>Acremonium restrictum</i> | 11.6 | 15.6 | 16.8 | 12.4 |
| S7 | <i>Absidia californica</i> | 10.3 | 16.1 | 17.8 | 12.3 |
| S8 | <i>Absidia repens</i> | 19.7 | 28.4 | 29.8 | 16.8 |
| S9 | <i>Penicillium chrysogenum</i> | 10.9 | 35.4 | 34.2 | 24.4 |
| TS10 | <i>Gibberellaputricaris</i> | 8.9 | 18.7 | 19.6 | 17.6 |
| TS11 | <i>Aspergillus.flavus</i> | 7.9 | 16.4 | 17.6 | 15.4 |
| Mean ± SD | | 13.5 ± 4.5 | 24.7 ± 8.7 | 26.2 ± 7.8 | 20.7 ± 7.1 |

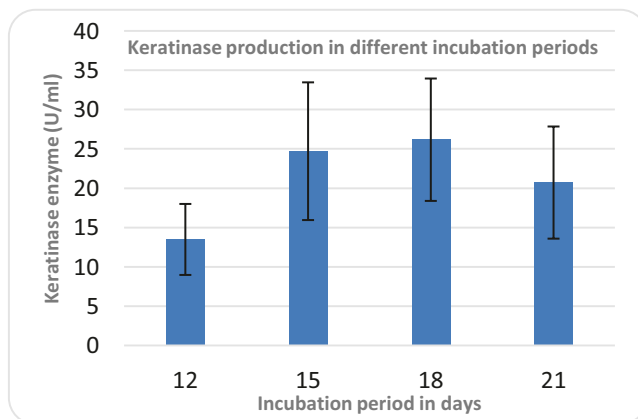


Fig. 4.5 Keratinase production in different periods

at 15 hrs. *Fusarium solani* produced maximum (39.8 U/ml) keratinase enzyme at the incubation of 15 days and minimum (7.9 U/ml) was produced by *Aspergillus flavus* after 12 days of incubation. The Kruskal-Wallis test showed that there was a statistically significant difference between the overall keratinase production by the fungi under the different incubation periods studied ($p = 0.004$ at H statistic 13.15).

4.4 Discussion

The molecular identification was carried out by DNA bar-coding using the ITS region sequencing. The DNA sequences were compared to those in the databases using NCBI-BLAST. Eleven species were identified using DNA bar-coding with an identical range between 97–99%. It is also proposed that DNA region sequence is one of the most important tools for the identification of the fungal species isolated from environmental sources. Hence, it has been widely used to detect the soil fungal community, and as an improvement of the classical identifications (Alsohaili & Bani-Hasan, 2018).

Pakshir et al. (2013) used the molecular technique for the first time in Iran to identify keratinophilic fungi. In the current investigation, 11 species of eight genera were identified by molecular marker-based sequence analysis (Fig. 4.1).

In the present examination, *Absidia californica* and *Absidia repens* were also found majorly in the soil sample of dumping sites. These fungi also have a great keratin-solubilizing potential. The current findings are in agreement with previous reports about *Absidia* as keratinophilic fungi. Morphological criteria and molecular marker for clear distinction of *Absidia* was used. Different species of clinically significant zygomycetes including 2 species of *Rhizopus*, 3 species of *Mucor*, 2 species of *Cunninghamella*, as well as 1 strain each of *Rhizomucorpusillus* and *Absidiacorymbifera* were identified by molecular techniques (Iwen et al., 2011).

Majority of reports on Keratinase production are under submerged shaking/static conditions (Cai et al., 2011). It is difficult to compare the production condition for Keratinase due to a variety of organisms and the methods of cultivation.

In comparison to the present findings, other studies reported that maximum enzyme activity was obtained within 96 h (150 U/ml and 90 U/ml for *Chrysosporium* and *Microsporum* respectively) of cultivation at pH 7.0, 30 °C. The optimal conditions for the keratinolytic activity of both enzymes were found to be at pH 9.0 and temperature 50 °C; however, the enzymes showed stability over a broad range of pH between 7.0 and 10.0 and temperature 30 °C–50 °C (Kanchana, 2013).

Similar studies were conducted by different scientists. Three factors, temperature, pH, and manitol, were used for keratinase optimization (Shankar et al. 2014). Similar work for keratinase production employing response surface methodology was performed (Ramnani & Gupta, 2004; Hashemet et al., 2018). Impact of different substrates, temperature, pH and protease inhibitors, lessening operators, and metal particle supplements in the generation medium on enzyme creation was studied (Yadav et al., 2011).

Previous reports show that the activity of keratinolytic enzyme produced by *Aspergillus flavus* effectively degraded feather substrate (Mini et al., 2015). Extremely high and low pH values showed complete loss of activity for most enzymes. Alkaline pH possibly supported keratin degradation as higher pH modified cysteine residues, making it accessible for enzyme action. Keratin degradation took incubation time for enzyme production from 24 h to several days. This is probably involved in the complex procedure of keratinolysis of these microbes (Kumawat et al., 2013). The harvesting time of 72 h was found to be ideal for maximum enzyme production.

These results showed similarity with our findings that keratinase enzyme production was maximum at 8 pH, between 15 and 18 days of incubation at 50 °C. Although the mean enzyme activity is maximum at 18 days of incubation period (Table 4.6), fungi such as *Fusarium solani* and *Penicillium chrysogenum* did show better activity after 15 days of incubation. Also, some fungi showed good activity after 21 days of incubation. For more precise optimization, enzyme activity could be observed on consecutive days between 15 to 21 days of incubation.

4.5 Conclusion

The findings concluded that total eleven keratinophilic fungi were isolated from the soil of Bhopal. All these fungi showed good keratinase production in shake culture method at 8 pH, 50 °C temperature, between 15–18 days of incubation period. These fungal isolates will be useful in eco-friendly and environment cleanup process of keratinic wastes. They may also be used for large-scale production of keratinase for industrial purposes. This research is suggestive of potential optimization scheme and its scope alongside the development of optimal medium compositions

and culture conditions, for the production of keratinolytic enzyme that can be useful for developing entrepreneurship.

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Chapter 5

Potential Impression of Arbuscular Mycorrhizal Fungi on Agricultural Growth, Productivity, and Environment Toward Global Sustainable Development for Green Technology



Kamal Prasad

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5.1 Introduction

The term *Green Technology* is an umbrella that designates the utilization of technology and science to make products that are additional eco-friendly. The chief goal of green technology is to shield the surroundings and repair past damage to the environment. Sustainable agriculture could be an important role in today's biosphere, because it offers the potential to fulfill our agricultural needs. A perfect agricultural system is sustainable, maintains and improves human health, surroundings edges, and produces sufficient food for the increasing current world population. Microbes play a crucial role in the rhizosphere and greatly have an effect on the availability of obtainable soil nutrients to plants and thereby modify the standard and quality of root exudates (Barea, 2000). The root exudates are rich sources of carbon, and therefore, the carbon fluxes are crucial determinants of rhizosphere operation. The discharge of root exudates and decaying plant material offer sources of carbon compound for the heterotrophic soil biota as either growth substrates or structural material for root-associated microbial flora (Werner, 1998). Numerous soil organisms act with one another to beat the limitation. Positive effects of arbuscular mycorrhizal fungal biofertilizer with and without combination with species of genus *Rhizobium*, *Azotobacter*, *Pseudomonas*, *Fraturia*, *Azospirillum* inoculants are reported for various plant species (Prasad et al., 2019). These microbes are usable in several post-harvest agricultural processes, in manure production or perhaps in control of pests/diseases. Extraction of pricey minerals from poor or leftover mines or production of raw materials for production of chemicals of huge utility or perhaps production of fuel gases are all modern achievements within the field of microorganisms. Presumably, thousands of different technologies could also be waiting behind the screen for their release in future agricultural development programmers for green technology. AM fungal biotechnology and its application are comparatively safe to be utilized in the field for all sorts of plants worldwide. Mycorrhiza are capable to provide micronutrients for effective plants growth, development and quality produce.

Food and nutrition security has long been one of the best challenges for humanity and given existing population and climate change eventualities is a progressively difficult task in current scenario. A number of the newest estimates predict the requirement to extend agricultural productivity by more than 70% by the year 2050, and therefore, the focus shifts progressively to the role of soil biodiversity in normally (Bender et al., 2016) and significantly AM fungi (Thirkell et al., 2017), in an exceedingly achieving in a sustainable methodology. Symbiosis could be a biological phenomenon involving dynamic changes within the genome metabolism and signaling network (Kawaguchi & Minamisawa, 2010). Most of the legumes possess two varieties of microorganism symbionts: mycorrhizal fungi and nitrogen-fixing bacteria, thereby establishing triple association, capable of supplying nitrogen and phosphorous contents to the plants (Prasad et al., 2019; Gautam & Prasad, 2001; Prasad, 2006a, b; Prasad, 2013; Prasad and Meghavanshi (2005). Each of the mycorrhizal fungi and *Rhizobium* act as microbial biofertilizer and have the distinctive

ability to convert nutritionally vital components from unavailable to available form through organic processes (Prasad et al., 2019; Vessey, 2003; Prasad, 2000, 2015, 2017, 2020; Harwani et al., 2009; Prasad & Pandey, 2012). The standard for mycorrhizal biofertilizers necessitates an in-depth examination of microorganism features, efficacy, consistency, as well as precautions and limitations, not only in the lab and in the factory, but also in the field. The AM fungal biotechnology is incredibly important, 100% natural, low cost, eco-friendly organic inputs, and has supplementary ability to the chemical fertilizers. The most effective quality AM fungal biofertilizers assist in providing all of the nutrients required by plants as well as maintaining the soil's standard in a natural environment. Mycorrhizal fungi have been used over the past fifty years to varied strategies for food safety; quality, environmental protection, and more practical treatment of agricultural crops give a most spectacular record of achievement. In recent years, however, mycorrhizal biotechnologies have been successfully applied to a variety of agricultural and environmental challenges. AM fungi are terribly helpful in eliminating issues related to the utilization of chemical fertilizers and pesticides; they are currently wide applied in natural farming and organic agriculture (Prasad et al., 2019; Prasad, 2015, 2017; Parr & Hornick, 1994). Soil microbiologists and microbial ecologists have well-tried to differentiate soil microbes as useful or harmful consistent with their functions. Microbes have the ability to effect soil quality, plant health, growth, and productivity. The present manuscript has been critically mentioned numerous aspects of mycorrhizae inputs application in agriculture and their economic importance for sustainable development for green technology.

5.2 Positive and Active Microbes

The mismanagement and extreme use of chemical fertilizers and pesticides have typically adversely affected the soil ecology, atmosphere and created food safety, and human and animal health issues. Consequently, there has been a growing interest in natural and organic agriculture by customers, microbiologists, and ecologists as potential alternatives to chemical-based mostly conventional agriculture. Agricultural systems that adapt to the principles of natural ecosystems are currently receiving an excellent deal of attention in each developed and developing country. A variety of articles has been published that deal with several aspects of natural and organic farming systems. Alternative and sustainable agriculture, soil quality, integrated pest and nutrient management, and even helpful microorganisms are being explored by the agricultural research analysis institution (Prasad et al., 2019; National Academy of Sciences (1989) Alternative Agriculture; Parr & Hornick, 1992; Prasad et al., 2005a, b; Meghavanshi et al., 2008). The use of single and consortium cultures of helpful microbes as soil inoculants relies on the principles of natural ecosystems that are sustained by their constituents; that is, by the standard quality and quantity of their inhabitants and specific ecological parameters, that is the bigger the variety and range of the inhabitants, the upper the order of their

interaction and also a lot of stable the ecosystem. The consortium culture approach is just an endeavor to use these principles to agricultural soils, and to shift the biological equilibrium in favor of increased plant growth, productivity, and disease protection (Prasad et al., 2019; Higa, 1994).

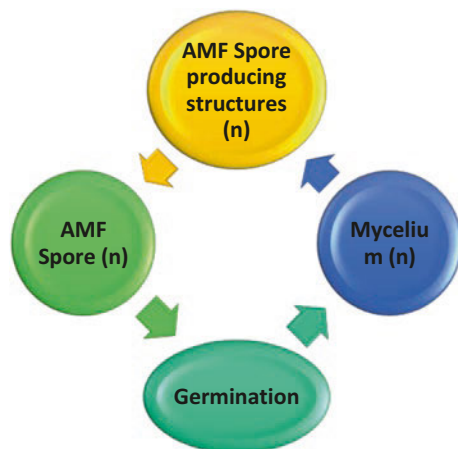
5.3 Arbuscular Mycorrhizal Fungi (AM Fungi)

An AM fungus is functionally an integral part of the plant community. AM fungi and therefore the plant community are interdependent. The term mycorrhiza was coined a century past to designate a mutual symbiotic association of plant roots and fungi. There are numerous forms of mycorrhizal fungi. Orchids and ericaceous plants have their own particular brand whole; however, those fungi that colonize *in vivo* and *in vitro* conditions are renowned as arbuscular mycorrhizal fungi. AM fungi are colonizing the root system of most native land plants. The AM fungal hyphae extend from root surfaces like little conduits out into the soil, greatly enhancing the absorptive surface area of root systems. These fungous colonize plants roots from a vital link between the plant and its soil environment, and work as an effective biological tool in making a healthy ecosystem (Prasad et al., 2005, 2019; Gautam & Prasad, 2001; Prasad, 1993, 2006a, b, 2010, 2011a, b, 2013, 2015, 2017; Prasad & Pandey, 2012; Prasad & Deploey, 1999; Prasad & Warke, 2018a, b). Approximately more than 90% of land plant species form association with AM fungi (Prasad, 2017). These plant species represent a broad variety of agricultural crops, agro forestry and horticultural crops, vegetables, oilseeds, grains crops, millets, and decorative and forage crops (Gautam & Prasad, 2001; Prasad, 2015, 2017; Trappe, 1987). The roots of most plant species from symbiotic association with specialized fungi (Genus *Glomus*, *Gigaspora*, *Acaulospora*, *Enterophospora* and *Sclerocystis*) are referred to as arbuscular mycorrhizal fungi (AMF). An AM plant is that the amplest kind of mycorrhizal fungi delineated as universal plant symbioses. Researches on mycorrhizal fungi conducted throughout the previous few decades envisaged their prevalence during a giant variety of hosts, completely different habitats and variability in quality and quantity (Gautam & Prasad, 2001; Prasad, 2015, 2017; Curl & Truelove, 1986; Deacon, 2019; Prasad & Kaushik, 2004). Prasad and Kaushik (2004) describe ecology, physiology, biochemistry, and taxonomy of mycorrhizal fungi. Previous few decades mycorrhizae fungi frequently used in several trees, agricultural crops, and decorative plants worldwide. AM fungi increase approximately 20–50% yield and enhance uptake of macro- and micronutrients, especially P, K, Zn, S, Fe, mg, Mn, Cu, boron, molybdenum, and water. Even it survived in harsh conditions in soil ecosystems. AMF is sometimes inoculated to seedlings, coated on seeds moreover as soil application. AM fungi also are applied as barren wasteland development, industrial contaminated and polluted land moreover as plant diseases and pest management (Prasad & Rajak, 2002; Johnson et al., 1997).

5.3.1 Development of Mycorrhizal Fungal Infection

The growths of AM fungal colonization within the host has been demonstrated in Figs. 5.1 and 5.2, and therefore the following steps: (1) The preinfection resting reproductive spore or hyphae can initiate infection during a host cell (Fig. 5.2). Germination of reproductive spore is primarily ruled by physical and chemical factors of soil such as temperature, CO₂, moisture, pH, presence of phosphate, etc. (Gautam & Prasad, 2001; Prasad, 1993; Smith & Read, 1997). Throughout the development of germ tubes, no divisions of nuclei occur, rather get disturbed in newly fashioned germinating structures. Once germ tubes encounter an appropriate living host, additional development of infection proceeds, whereas in absence of appropriate host, the infection potential of germinated fungal propagule is lost. (2) The AM fungal hyphae penetrate the root between epidermal cells and create an entry (Gautam & Prasad, 2001; Prasad, 1993; Li et al., 2006). The appressorium is the first structure formed within the host cells, and it signifies the start of a successful infection. After this, the autotrophic growth of fungus ceases. (3) The AM fungal hyphae grow inter- and intracellular and develop infection segment spread either by the expansion of hyphae within the root (internal spread) and/or by the formation of secondary entry points adjacent to the primary (Gautam & Prasad, 2001; Prasad, 1993; Li et al., 2006). Within the deeper root cortex, the living intracellular hyphae are positioned paralleled to the long axis of the root system. Shortly when hyphae penetration, arbuscules formation can be initiated in cortical region. Arbuscules are powerfully branched hyphae, and its formation is accompanied by stretching of host cell wall and invagination of plasma membrane. Most intensive affiliation between fungus and plants are maintained through arbuscular and it is believed to be the particular site of phosphate transfer (Prasad et al., 2019; Gautam & Prasad, 2001; Prasad, 1993; Adesemoye et al., 2009). The arbuscules have distinct wall morphology that is improving up to ten times thinner than extraradical hyphae and fibrillar nature of wall be usually lost and becomes amorphous (Prasad, 2007). When

Fig. 5.1 Sketching diagram of asexual life cycle of arbuscular mycorrhizal fungi



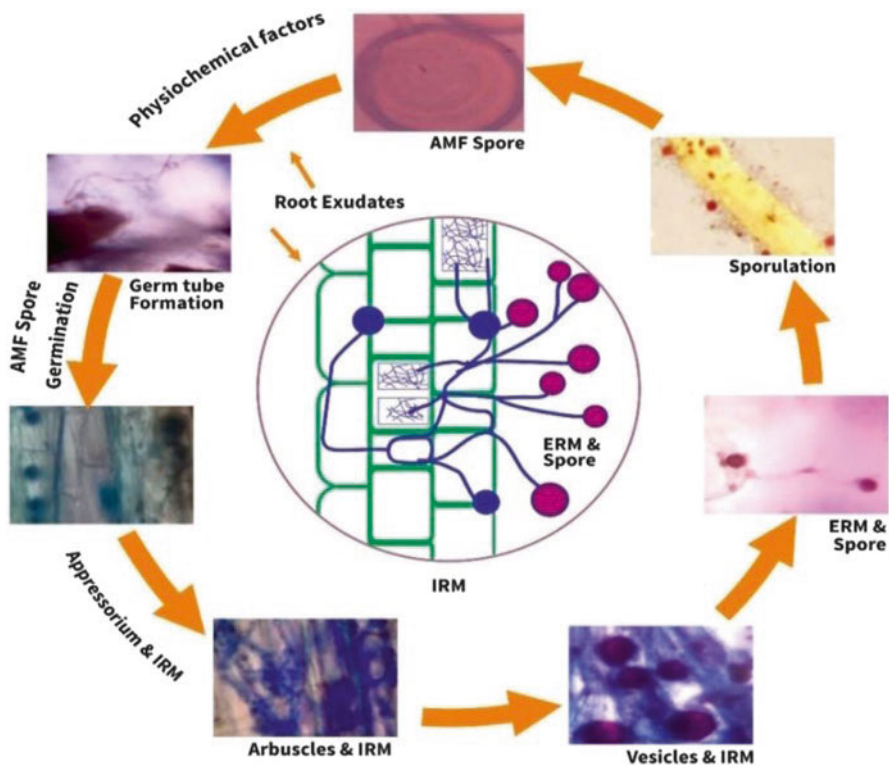


Fig. 5.2 Structural characterization and development of AM Fungal infection within the host cell and their life cycle in soil ecosystem

colonization in root tissue, apical or intercalary swelling of hyphae known as vesicle containing lipid. They have relatively thinner wall than spore. In conjunction with the interior spread of infection, the fungus ramifies the rhizospheric soil and generally extends as over eight cm from the root (Gautam & Prasad, 2001; Prasad, 1993). (4) AMF forms resting spores to external hyphae known as arbuscular mycorrhizal fungal spores. Spore formation can begin as early as 2-4 weeks when primary infection or can take up to three–six months. Short-lived fine hyphae were primary concerned in nutrient acquisition and ne’er colonized roots, whereas a cluster of thicker hyphae were exclusively concerned in colonizing roots (Gautam & Prasad, 2001; Prasad, 1993). As the host plant grows, new roots are produced that become successively infected by runner hyphae or new developed spores. Arbuscules and vesicles are unceasingly fashioned and degraded. Spore germination and spore formation occur at a similar time. Thus, at any stage, numerous phases of infection can be observed (Prasad, 1993).

5.3.2 *Characteristics of Arbuscular Mycorrhizal Fungi*

The vesicles and arbuscules formation within the root cortex are most vital characteristics of mycorrhizal fungi (Fig. 5.2). The root colonization is directly linked to external AM fungal mycelium that spreads and ramifies within the soil. The appressorium of AM fungal-infected plants forms on the host root surface and penetrates the epidermal and exodermal cells, with hyphal branches passing into the root's middle and inner cortex. So, they sometimes form coil-like structures. The intercellular distributive hyphae grow parallel to the root axis. The lateral branches then penetrate root cells leading to the formation of arbuscules (haustoria-like structure) that develop further by repeated dichotomous branching among the host cells to yield a cluster of fine filaments. Within the next stage (after 2–4 days), the arbuscules disintegrate (digestion stage) to produce a granular mass. At the tip or within the middle of distributive hyphae swellings develop inner or intracellular to create vesicles. AM fungal species of *Gigaspora* and *Scutellospora* are don't seem to be forming vesicle, they solely form arbuscules; thus, presently vesicular arbuscular mycorrhizal fungi (VAMF) are known as arbuscular mycorrhizal fungi (AM fungi). AM fungi colonization is nearly no modification in external morphology of the root. The size and shape of the vesicles are storage and reproductive structure. With relevancy to transfer of nutrients, it had been thought to require place through arbuscule digestion. AM fungi uninterrupted in soil as spores or propagules for sizable amount over a long time. Spores can germinate within the soil with appropriate host plants. Completely different species of AM fungi endophyte have different temperature limits and optimum for germination. Mycelial growth forms interconnections between interconnected plants. It survives in dry soil environment for a long time.

5.4 Positive Impact of AM Fungi on Agricultural Crops and Environments

5.4.1 *AM Fungi and Plant Growth*

Mycorrhizae play positive role in phosphate (P) uptake by almost all terrestrial plants. The matter for P₂O₅ nutrition comes from heterogeneity of soils; low-quality ends increase in native depletions. In most soils, massive a part of P is held in mineral lattices and so isn't at once exchangeable with ions in soil solution. The huge synthesis of cytoplasm is proportional to extend in plasmalemma area in arbuscular region. AM fungi are capable for convert insoluble P to soluble form. Roots of AM fungi-mediated plants uptake again and again additional P from soil. Presently it's been created clear that replenishment of supply of P from distant soil (beyond the eight cm distance from roots) to root surfaces being low, the mycorrhizal hyphae, having organic connection with the root, extend to these distance and absorb macro and micro nutrients (Prasad et al., 2005a, b; Prasad, 1993). This AM

fungi-mediated phosphate transport turn up in three phases specifically uptake by fungal hyphae in soil, movement to hyphae within root cortex, and release in host plant. Typically, the fungus uptake P from soil as orthophosphate ions from the labile pool and its translocation to root is believed to take place mainly by cytoplasm streaming of polyphosphate granules within the vacuoles of the AM fungi. Presently AM fungi are being thought of as a system of various importance in hemisphere. Their effects in inflicting shifts in chlorophyll pigment and phytohormone gradients, resistance to water stress, and attenuate escape of electrolytes from injured or pathologic cells are gaining importance and drawing attention of scientists (Prasad et al., 2019; Gautam & Prasad, 2001; Prasad, 2011a; Prasad & Warke, 2018a, b; Prasad & Rajak, 2002). Scientists have determined that growth of varied crops once infected with completely different species of AM fungi was increased, thereby improving crops growth and productivity. Rotations of crops influence the stability of mycorrhizal flora in soil ecosystems. The importance of AM fungi as a tool for improving the expansion and productivity in various groups of plants was recognized solely once the work of Baylis (Baylis, 1959), and Mosse (Mosse, 1962). However, throughout last five decades, plenty of documents have been gathered concerning the taxonomy, ecology, physiology, and anatomy of AM fungi and their relation with their hosts particularly with relation to uptake of water, enhance root absorption capability, uptake of macro- and micronutrients, growth hormone production, antibiotic secretion, barren wasteland management, biotic and abiotic management, environmental protection, and management of diseases and pests (Fig. 5.3), (Prasad et al., 2019; Prasad & Kaushik, 2004; Prasad & Rajak, 2000, 2001, 2002; Rewari & Tilak, 1988; Warke et al., 2018; Bitterlich et al., 2019; Mathur et al., 2019). Trees are being cultivated in order to use them in a variety of plant production methods (Prasad et al., 2019; Prasad, 1993; Prasad & Warke 2018a; Rahimzadeh & Pirezad, 2019; Ji et al., 2019; Ghorchiani et al., 2018; Mayer et al., 2019).

5.4.2 AM Fungi and Fertilization

The most widely appreciated contribution of AM fungi to plant performance is their ability to extend nutrient uptake, notably of P (Prasad, 2002a, b; Smith & Smith, 2011). Harnessing the nutrient provides by AM fungi, the amount of applied fertilizer, and therefore the energy linked to its production can be reduced. A significant issue in optimizing economical fertilization is reducing the amount of nutrients lost to the system via leaching. AM fungi decrease nutrients leaching not solely increasing the nutrient interception zone due to the development of a mycorrhizosphere; however, additionally increased nutrients uptake increased soil structure and fostering of the microbial community with associated nutrient immobilization (Cavagnaro et al., 2015). Kohl and van der Heijden (Kohl & van der Heijden, 2016) confirmed that different AM fungal species differ in their ability to decrease nutrient leaching. AM fungi increase nutrients leaching in extremely fertile agro ecosystems due to

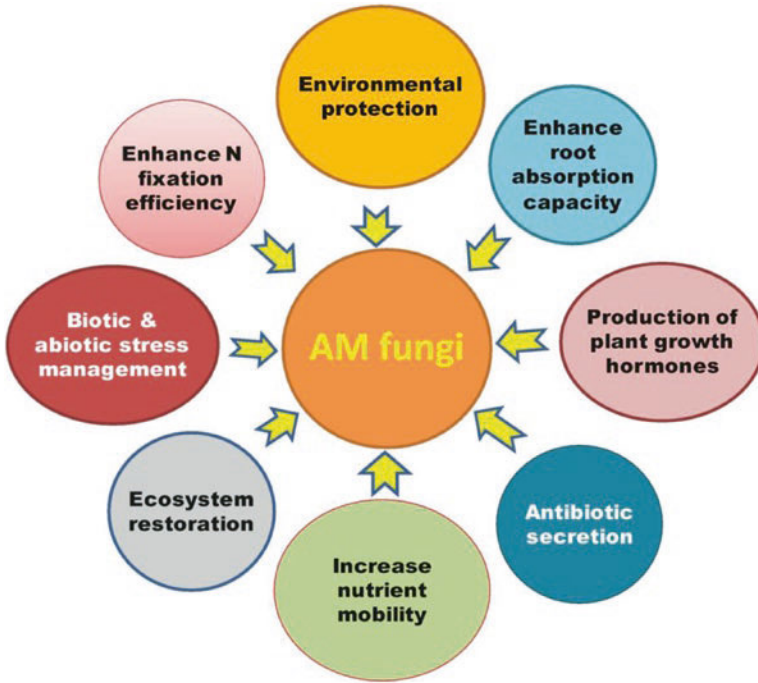


Fig. 5.3 A diagrammatic representation of arbuscular mycorrhizal fungi functions to regulate various processes in the ecosystem and plant growth promotion and protections

larger soil nutrient content. (van der Heijden, 2010). AM fungi are shown to stabilize community productivity across gradients of nutrient availability and to reduce plant tissue nutrient content variability on such gradients in grassland (Yang et al., 2016). They transferable to agricultural systems, these effects would be crucial in achieving food and nutrition security notably in regions wherever access to fertilizers could be restricted or irregular. Furthermore, extending the available soil nutrient pool to deeper soil might improve resistance, for example, maintaining plant growth under dry situations, where minerals in the soil may be there but are positional untouchable for the roots. Altogether, with the endlessly increasing costs of chemical fertilizers and their foretold deficiency in a very close to future, creating the foremost out of those resources is that the solely approach forward and soil and AM fungi could prove necessary in agriculture industry.

5.4.2.1 AM Fungi and Nitrogen Absorption and Translocation

Nitrogen (N) may be a premier growth-limiting issue for higher plants and agricultural crops. Varied studies have processed that AM fungi have the power to absorb and transfer N to the proximate plants as well as host plants (Prasad, 1993; Hodge

& Storer, 2015; Battini et al., 2017; Turrini et al., 2018). Nitrogen applied in agricultural fields is lost via leaching or in type of gaseous emissions. Leachate N happens principally in form of dissolved nitrate (NO_3^-), a particularly mobile type of N in soil. AM fungi promote soil aggregation (Zhu et al., 2012; Leifheit et al., 2014; Nouri et al., 2015) by improving soil structure and thus increasing soil water-holding capability. Moreover, AM fungi take up nitrogen preferentially within the type of ammonia (NH_4^+), reducing the pool of N out there for nitrification and consequently reducing the mobility of nitrogen. In soil, AM fungi might intercept N that migrated down the profile and immobilize it or deliver it to the plant, so avoiding N losses. Moreover, the proportion of NH_4^+ to different N sources can increase in soil (Prasad, 2007; Meghavanshi et al., 2010; Kautz et al., 2013), increasing the potential role of AM fungi in mobilizing and delivering N to the plant, reassuring access to an antecedently untouchable pool and reducing the requirement for N fertilization. Zhang et al. (2018) have verified AM fungi-mediated plants increased allocation of shoot biomass to panicles and grains through increased N and P redistribution to panicles significantly below low fertilizer levels. Translocation of N into seeds is increased from heading to maturity. AM fungal interdependency turns out in-depth underground extraradical hyphae starting from the roots increase to the encompassing rhizosphere, thereby serving to in improving the uptake of nutrients specifically nitrogen (Battini et al., 2017). The interaction of salinity stress and AM fungi considerably affects the concentrations of P and N and also the N: P magnitude relation in plant shoots (Wang et al., 2018). Recently, it has been reported that native AM fungi treatments produce significant alterations within the N contents of crop plants (Turrini et al., 2018). It has been widely accepted that fungi have the power to require substantial quantity of N from dead and decomposed material that later can increase their fitness to grow and keep alive. With the exception of this, massive biomass and increased N needs for AM fungi render them the most stakeholder of global N pool that's equivalent in scale to fine roots. Thus, they play a crucial role within the N cycle (Hodge & Fitter, 2010). The AM fungi extraradical hyphae can absorb and assimilate inorganic N (Jin et al., 2005). Many studies have shown that approximately 20–75% of the overall N uptake of AM plants is transferred by the AM fungi to their hosts (Tanaka & Yano, 2005; Govindarajulu et al., 2005; Ahanger et al., 2014; Hameed et al., 2014; Hashem et al., 2018). Increased N in AM fungal-mediated plants evidently results in higher chlorophyll pigments, as chlorophyll molecules can effectively trap N (Prasad, 1993; de Andrade et al., 2015). AM fungal immunization improves C and N accumulation and N assimilation below ambient and elevated CO_2 concentrations (Zhu et al., 2012).

5.4.2.2 AM Fungi and Phosphorus Translocation

Phosphorus (P) is typically assumed low quality in soils. P action is important, and most effort has been spent on avoiding P loss and P-mediated eutrophication via wearing. However, excessive manuring, existence of advantageous pathways, or a sandy soil texture can cause vital P action (Djordjic et al., 2004; Schoumans, 2015),

with its associated economic and environmental consequences. The role of AM fungi in P uptake has been extensively researched (Prasad, 2000; Smith & Smith, 2011), which they are going to cut back the need of significant manuring attributable to exaggerated economical P uptake. AM fungi increase water-holding capability, reducing the danger of action. AM fungi might intercept P that has migrated down the profile and deliver to the plant. Uptake of organic P from soil, mainly via roots but to boot with direct injection of organic matter, can keep inaccessible to the plant attributable to attenuated decomposition and mineralization rates. Similarly, in another study, it was recorded that AM fungi in soil might contribute further to plant P nutrition than soil AM fungi, below serious P fertilization (Wang et al., 2017). Consequently, soil AM fungi have potential to be of nice association inside the rejection of P loss, notably in sandy soils or once the soil is P saturated. Phosphorus is usually assumed low quality in soils. P leaching is importance and most effort has been spent on avoiding P loss and P mediated eutrophication via soil erosion. However, excessive manuring, existence of discriminatory pathways, or a sandy soil texture can lead to significant Pleaching (84–85), with its associated economic and environmental consequences. The role of AM fungi in P uptake has been extensively researched (Prasad, 2000; Smith & Smith, 2011), and that they can scale back the necessity of heavy manuring due to increased economical P uptake. AM fungi are increase water holding capability, reducing the chance of leaching. AM fungi may intercept P that has migrated down the profile and deliver to the plant. Uptake of organic P from soil, principally via roots however conjointly with direct injection of organic matter, can remain inaccessible to the plant due to decreased decomposition and mineralization rates. Wang et al. (2017) found some proof that AM fungi in soil would possibly contribute a lot of to plant P nutrition than soil AM fungi, under heavy P fertilization. As a result, soil fungi have the potential to play an important role in the rejection of P loss, particularly in sandy soils or when the soil is P saturated.

5.4.2.3 AM Fungi and Potassium Nutrition

The role of potassium remains poorly investigated in mycorrhizal analysis. Plant potassium nutrition is improved by mycorrhization in crops, particularly beneath K⁺ limiting conditions as found in forest ecosystems. Moreover, this improvement may act on abiotic stress tolerance, P equilibrium maintenance, or exclusion of soil contaminants. Moreover, different reports have shown increased activity of a K⁺ transporter within the mycorrhizal roots of *Lotus japonicus* (Guether et al., 2009; Berruti et al., 2016).

5.4.2.4 AM Fungi and Other Mineral Nutrition

AM fungal immigration is stimulating nutrients uptake in overall agricultural crops. AM fungi symbiotic association is to transfer organic carbon within the kind of lipids and sugars (Jiang et al., 2017; Luginbuehl et al., 2017). AM fungal vaccination in plant can enhance the concentration of assorted macro- and micronutrients considerably, that ends up in exaggerated photosynthate production and therefore exaggerated biomass accumulation (Prasad, 2017, 2020; Mitra et al., 2019). AM fungi have the flexibility to extend the uptake of inorganic nutrients in nearly all plants. Except for macronutrients, AM fungal colonisation has been done in order to increase the provision of micronutrients like zinc and copper (Smith & Read, 1997). AM fungi develop in a living extension with root system and dramatically increasing access to get essential nutrients such as N, P, K, Ca, Zn, Mn, Mg, Cu, B, S, and Mo from the soil surroundings. Consequently reciprocally, the host plant feeds the sugar and different organic substance that the AM fungi need to survive and grow. Therefore, AMF give nutritional support to the crop plants even beneath inappropriate conditions within the root cells. AM fungus produce arbuscules, which aid in the interchange of inorganic minerals and thus carbon and phosphorus compounds, giving host plants a significant boost in vigour (Prasad, 2017, 2020; Li et al., 2016). Increased photosynthetic activities and different leaf functions are directly associated with higher growth frequency of AM fungi colonization that is directly linked to the uptake of nutrients and promote the alteration of crop yield. AM fungi interdependency absolutely exaggerated the concentrations of N, P, and Fe in *Pelargonium graveolens* L. beneath drought stress (Amiri et al., 2017). Improved levels of P, Ca, and K were reported in AM fungal associated *Euonymus japonica* beneath salinity stress condition (Gomez-Bellot et al., 2015). It's supposed that AM fungi increase the uptake of virtually all essential macro and micro nutrients and contrarily decrease the uptake of Na and Cl, resulting in growth stimulation (Evelin et al., 2012). The extraradical hyphae can effectively improve nutrient uptake, thereby improving plant growth and development (Lehmann & Rillig, 2015). Numerous scientists have conferred the role of AM fungi in uptake of soil essential nutrients and effectively promote the expansion of host plants (Prasad, 2020; Smith et al., 2011). Improvement of plant nutrition and maintenance of Ca²⁺ and Na⁺ ratio are the significant dynamic attributes that facilitate improving useful aspects of AMF colonization on overall plant performance (Evelin et al., 2012; Abdel Latef & Miransari, 2014). Increased growth and levels of protein, Fe, and Zn were recorded in AM fungal-treated chickpea (Pellegrino & Bedini, 2014). Asrar et al. (2012) reported that the required fungal association increased the contents of macronutrients such as N, P, K, Ca, and Mg of *Antirrhinum majus* beneath drought condition. AM fungi additionally well tried to be effective in limiting the high accumulation of Na, Mn, Mg, and Fe in roots (Bati et al., 2015). Many studies unconcealed that AM fungous (*Glomus mosseae* and *Rhizophagus irregularis*) exhibited improved heavy metal translocation within the shoot (Zaefarian et al., 2013; Ali et al., 2015). Micronutrients such as Zn and Cu being diffusion restricted in soils are absorbed by plants with the assistance of mycorrhizal hyphae.

5.5 AM Fungi and Greenhouse Gas Emission

Modern agriculture is responsible for around 12% of global anthropogenic greenhouse gas emissions (Linguist et al., 2012). Most of the emissions are related to chemical fertilizer production and therefore the use of significant heavy machinery; however, most of them occur within the type of direct emissions from the field. AM fungi is having potential for reducing emissions from chemical fertiliser (Prasad, 2017). The role of AM fungi in reducing the discharge of carbon dioxide and nitrous oxide greenhouse gasses related to agriculture.

5.6 AM Fungi and Carbon Farming

The traditional, terribly stable carbon found in soil is problematic. Stable soil carbon is also promptly decomposed once contemporary carbon is added. AM fungi have the potential to counteract this development due to their function in soil structure and within the capture of nutrients (Prasad, 2017). AM fungi contribute in the plant to soil organic matter by making a sink demand for plant carbon and distributing to hyphal biomass. The role of AM fungi is essentially unnoticed in terrestrial carbon cycling and global climate change models despite their larger involvement in net primary productivity augmentation and any accumulation of this extra photosynthetic fastened carbon within the soil. However, this buffering mechanism against elevated carbon dioxide condition to sequester additional carbon by AM fungal interaction.

5.7 AM Fungi and Soil Management

Soil is also restricted by physical properties, just like the existence of a troublesome plow pan that stops root growth. The advantages of deep tillage and various soil tillage management options are also debatable and contextdependent; yet, assuming the presence of a plough pan, they are very frequent. Yields are also significantly increased once deep ploughing (Schneider et al., 2017). The existence of AM fungal mediated plant phylotypes and their inability to survive soil mixture events, however, entails precaution and conjointly the overall turning of any methodology that inverts the profile (Sosa-Hernandez et al., 2018). Intensive tillage has been referred to as a major issue reducing AM plant abundance and variety in agriculture (Kabir, 2005). Recently, Sale et al. (2015) has examined the outcomes of reduced and traditional tillage, down to 40cm within the profile exploitation reproductive structure based on community analysis. Their results make sure the expected shifts in reproductive structure abundance and variety in soil but those effects weren't necessary in deeper layers, despite a shift in community composition. The absence of AM

fungi reproductive structure abundance shifts do not basically imply an absence of impact on hyphal abundance or settlement rates, but changes in soil community composition highlight that tillage can have an impact on AM fungi in deeper layers, with unknown consequences for its usefulness. AM fungal immunisation has managed the soil structure and performance.

5.8 AM Fungi and Heavy Metals

AM fungi extensively support plant establishment in soils contaminated with heavy metals, due to their potential to strengthen defense of the AM fungal-mediated plants to promote growth and development. Heavy metals could build up in food crops, fruits, vegetables, and soils, posing a health risk to the common people (Liu et al., 2013; Yousaf et al., 2016). AM fungi association with wheat completely multiplied nutrient uptake aluminum stress (Aguilera et al., 2014). Plants well-grown on soils enriched with Cd and Zn exhibit significant suppression in shoot and root growth, leaf chlorosis, and even death (Moghadam, 2016). There are several reports within the literature on uncovering the AM fungi elicited effects on the build-up of metals in plants (Souza et al., 2012). Heavy metals are immobilized within the fungal hyphae of internal and external origin (Ouziad et al., 2005) that have the power to repair heavy metals within the cell wall and store them within the vacuole or could chelate with other substances within the cytoplasm (Punamiya et al., 2010) and therefore reduce the metal toxicity within the plants. The strong impacts of AM fungus on plant development and growth under harsh conditions have commonly reported (Kanwal et al., 2015; Miransari, 2017). It is additionally believed that increased growth or chelation within the rhizospheric soil can cause metal dilution in plant tissues (Kapoor et al., 2013; Audet, 2014). AM fungi reportedly bind Cd and Zn within the cell membrane of mantle hyphae and cortical cells, thereby limiting their uptake and leading to improved growth, yield, and nutrient status (Andrade & Silveira, 2008; Garg & Chandel, 2012) AM fungal mycelia having a high cation exchange capability and absorption of metals (Takacs & Voros, 2003). AM fungi are believed to manage the uptake and accumulation of some key inorganic nutrients and increased uptake of Si has been reportable in mycorrhiza-inoculated plants in soybean and *Zea mays* (Clark & Zeto, 2000). Hammer et al. (2011) additionally recorded significant uptake of Si in spores and hyphae of *Rhizophagus irregularis* and its transfer to the host roots. It is pertinent that low Cd mobility and toxicity can even be addressed with AM fungi by increasing soil pH (Shen et al., 2006), restoring Cd within the additional radical mycelium (Janouskova & Pavlikova, 2010), and binding Cd to glomalin, a glycoprotein. AM fungi are very effective in lowering the level of Cd in the vacuoles and cell wall, leading to Cd detoxification (Li et al., 2016). Wang et al. (2012) determined that AM fungal-mediated alfalfa (*Medicago sativa* L.) improved Cd tolerance because of the modification of chemical form of Cd in numerous plant tissues. Various processes that occur through the AM fungi are immobilization/restriction of metal compounds, precipitation of polyphosphate

granules within the soil, adsorption to fungal cell wall chitin and heavy metal chelation within the fungus.

5.9 AM Fungi and Environmental Temperature

Soil temperature increases, plant community reactions are also captivated with AM fungal interactions for sustainable yield and production (Bunn et al., 2009). Heat stress has a significant impact on plant growth and development, causing loss of plant vigour, seed germination suppression, scrubby growth rate, and lower biomass production, wilting and burning of leaves and generative organs, abscission and senescence of leaves, injury additionally as discoloration of fruit, cell death, reduction in yield (Wahid et al., 2007; Hasanuzzaman et al., 2013), and increased oxidative stress.

Generally, AM fungal-immunized plants show higher growth below heat stress compared to non-AM-fungal-inoculated ones (Gavito et al., 2005). Maya and Matsubara (Maya & Matsubara, 2013) have reported the association of AM fungi with plant growth and development, resulting in positive changes in growth below the conditions of height temperature (Fig. 5.4).

AM fungi can increase plant tolerance to cold stress (Liu et al., 2013; Birhane et al., 2012; Chen et al., 2013). Furthermore, according to the majority of findings, plants inoculated with AM fungus at low temperatures grow faster, whereas plants not inoculated with AM fungi grow slowly (Liu et al., 2013; Chen et al., 2013; Zhu et al., 2010a; Abdel Latef & Chaoxing, 2011a). AM fungi support plants in

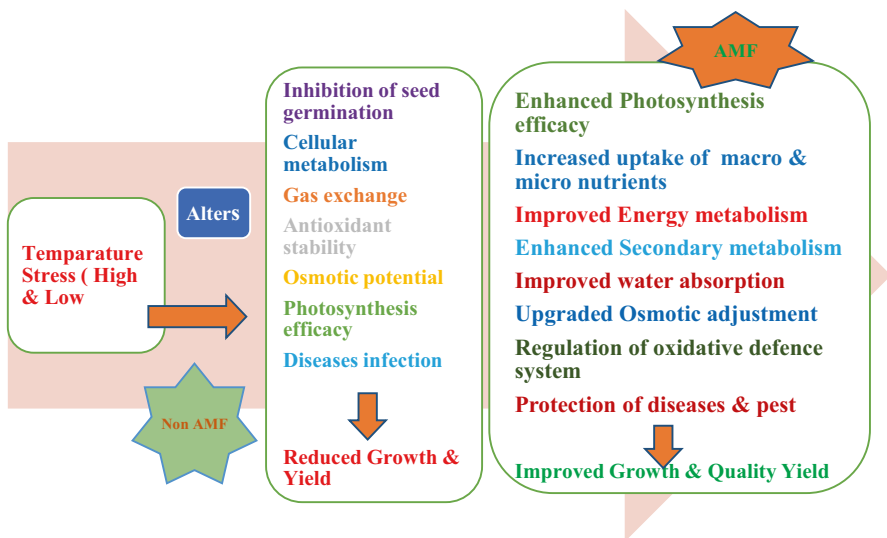


Fig. 5.4 AM Fungi mediation alleviates temperature stress in plants

combating cold stress and eventually improve plant development (Birhane et al., 2012; Gamalero et al., 2009). Moreover, AM fungi can also retain moisture within the host plant (Zhu et al., 2010), increase plant secondary metabolites resulting in strengthen plant immune system, and increase protein content for supporting the plants to combat cold stress conditions (Abdel Latef & Chaoxing, 2011a). AM fungal symbiotic relationship improves water and plant relationships and can increase gas exchange potential and osmotic adjustment (Zhu et al., 2012). AM fungi improve the synthesis of chlorophyll, resulting in a major improvement within the concentrations of varied metabolites in plants in cold stress conditions (Zhu et al., 2010; Abdel Latef & Chaoxing, 2011a). The role of AM fungi throughout cold stress has conjointly been according to change protein content in tomato and different vegetables (Abdel Latef & Chaoxing, 2011a).

5.10 AM Fungi and Abiotic Stresses

It is widely accepted that AM fungi may improve various stresses or combinations of stresses condition such as drought, salinity, temperature, nutrients, and heavy metals. Exposure of plants to a combination of drought and salinity causes an increased production of reactive oxygen species, which might be extremely injurious to plants (Bauddh & Singh, 2012). Detoxification of reactive oxygen species is finished by the enzymes that included commonly superoxide dismutase, catalase, peroxidase, and glutathione enzyme (Ahanger & Agarwal, 2017). AM fungi are improving plant growth and yield beneath stress (Abdel Latef & Chaoxing, 2011; Abdel Latef & Chaoxing, 2011; Abdel Latef, 2011; Abdel Latef & Chaoxing, 2014). AM fungal interdependency protects plants against a range of abiotic stresses exploitation varied processes such as improved photosynthetic rate, uptake and accumulation of mineral nutrients, accumulation of osmoprotectants, regulation of antioxidant enzyme activity, and alter within the rhizosphere ecosystem (Barzana et al., 2015; Calvo-Polanco et al., 2016; Yin et al., 2016). Numerous studies have shown improved nutritional status of AM fungal-mediated plants beneath osmotic stress conditions (Lehmann & Rillig, 2015; Auge et al., 2014; Lehmann et al., 2014), ensuing from deficit irrigation or salinity. Similarities among the tolerance mechanisms could occur in response to AM fungal-mediated combined stress diversifications. AM fungal colonization alterations in growth regulator profile, mineral uptake and assimilation, accumulation of compatible osmolytes and secondary metabolites, and upregulation of antioxidant system can be the common mechanisms induced throughout different stresses. However, specific mechanisms like compartmentation and sequestration of toxic ions, production of phytochelators, and macromolecule (protein) expression are specific and exhibit a significant amendment with stress type and also the AM fungal concerned. Changes in root characteristics improve the osmotic stress tolerance to extended levels (Evelin et al., 2009). The AM fungi could elevate the nutraceutical quality of crops and could be

of extended scientific agronomic importance for the production and management of various potential crops.

5.11 AM Fungi: A New Dimension for Green Technology

Universal microbiologists take into account that the number of soil microbes can be improved by applying organic amendments to the soil. The majority of soil bacteria are heterotrophic, thus this is usually true. They need complex organic molecules of carbon and nitrogen element for metabolism and biogenesis. The use of AM fungi and various organic inputs (seaweed extract, organic microbial fertilisers, fishmeal, crushed crab shells, and so on) not only helps to balance the micronutrient content in soil, but it also helps to enhance the population of important antibiotic producing actinomycetes. It is changing the soil to a disease suppressive condition over a comparatively short period. All useful microbes predominant in organic farming or natural farming ways can rely upon the ecosystem and environmental conditions. The ultimate goal is to pick microorganisms that are physiologically and ecologically compatible with one another, which can be introduced as single or consortium cultures into soil wherever their useful effects can be realized (Prasad et al., 2019; Prasad & Pandey, 2012; Prasad, 2015, 2017, 2020; Higa, 1994).

5.12 Conclusion and Future Prospects

With an intensive literature assessment, it is often all over that the AM fungi are utilized in biological science, agriculture, and forestry for numerous resolutions. A major deliberation in their application of valuable AM fungi to soils is that the improvement of their synergistic properties. AM fungi are to be effective once immunization into soil or agricultural crops. Various research analysis reports have already recognized the helpful role of AM fungi in improving plant growth and productivity even nerve-wracking environments. Therefore, during this manuscript, the prevailing information associated with the important role of AM fungi has been combined during a coherent manner for beneath standing of AM fungal dependent relationship with a spread of plants under stress environments. Mostly, AM fungi are principally mentioned as helpful entities for macro- and micronutrients uptake from soil ecosystem; but recently, it has been clearly pictured that plants inoculated with AM fungi can effectively combat numerous environmental such as salinity, drought, nutrient stress, alkali stress, cold stress, and extreme temperatures, and therefore helps increase yield in numerous crops. Encouragement of the use of AM fungus is critical for the long-term viability of modern international agriculture systems. AM fungal application in agricultural improvement can considerably scale back the utilization of artificial fertilizers, pesticides, insecticides, and alternative chemicals, thereby promoting biologically healthy agriculture. Truly AM fungi live

in the soil facilitate plants to absorb most macro- and micronutrients for overall performances. AM fungi help plant growth, productivity, management of weeds, pests, and diseases and assist the plant to require increased essential energy sources. In return, plants donate their waste byproducts for the AM fungi to use as food. Phosphate, nitrogen, and potassium are necessary macronutrients for the expansion of plants. These compounds exist naturally within the surroundings environment; however, plants have a restricted ability to extract them. Phosphate plays a crucial role in crop stress tolerance, maturity, quality and directly or indirectly, in biological nitrogen fixation. AM fungal vaccinated plants boost growth and output, and they are frequently used to meet the increasing consumption demands of the world's population. These environments friendly technologies shall be extremely inspired due to their widespread uses in agriculture, forestry, floriculture, green land development, polluted land, and bioremediation and wasteland development. Typically, AM fungi elicited modulations within the tolerance mechanisms and also the cross-talk triggered to manage plant performance can facilitate improve crop productivity. It absolutely was all over that AM fungal biotechnology is cost effective, higher substitute of chemical fertilizers and better quality and productivity and nutritive crops. These approaches can be appropriate for agriculturalists and organic alimental crops growers with their low inputs farming practices. This technology has the potential to improve soil health while also preserving the environment for current and future generations.

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Chapter 6

Mycometabolites in Industrial Applications with Emphasis on Bioherbicide Production



Puja Ray and Mayukh Ghosh

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6.1 Introduction

Weeds are considered a perpetual menace leading to reduction of crop yield by limiting growth-associated resources (water, nutrient, sunlight, space, etc) and being alternate and alternative hosts of crop pathogens and insect pests (Kumar et al., 2021). Many weeds and nonnative invasive plant species, due to their large seed bank, fast growth rate, lack of natural control agents, allelopathic potential, etc., negatively impact native biodiversity and hence ecosystem goods and services

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(Duncan et al., 2004). Present agricultural systems target to achieve maximum yield while minimizing the costs. This leads to several weed management options with more emphasis on using chemical herbicides, which are effective, fast, and relatively cheap means of weed control. In a recent study (Qi et al., 2020), it was observed that nontarget impacts of herbicides' resulted in changing the species composition, reduced the number of plant species, ecological adaptability of plants and thus reducing community diversity even at sublethal doses not only in the agricultural fields where herbicides are being applied but also in non-crop situations including field edges, fallow land, and other seminatural habitat. This herbicide further drifts into water bodies and protected areas causing a range of negative impacts on the ecosystems worldwide. Thus, owing to nontarget herbicide impact more emphasis is being made on alternate control strategies, including bioherbicides based on microorganisms (Ray et al., 2008).

Kingdom fungi are a diverse group of microorganisms with possibly over five million species (Blackwell, 2011), of which only a small fraction (5% approx.) have been identified. They are associated with some of the crucial ecological processes, like decomposition, recycling, and transportation of nutrients in different environments. Fungi produce a wide array of low molecular weight, non-nutritive metabolites that are exuded into the environment and play a key role in biotic interactions across species. The astounding structural diversity of these natural compounds has attracted the researchers globally to search for lead novel structures and find their possible utilization. The discovery of penicillin in 1928 by Alexander Fleming from the culture plate of bacterium *Staphylococcus* contaminated with the filamentous fungi *Penicillium rubens* has led to a revolutionized substantial research on fungi and their metabolites and simultaneous isolation of thousands of new fungal metabolites with a diverse range of biological and pharmacological effects. Many of these novel and natural bioactive compounds have potential bioprospecting applications in industries including pharmacy, cosmetics, nutraceuticals, agriculture, etc., Some of the well-known mycotoxins include ergot alkaloids, aflatoxins, and fumonisins. There is an increasing interest in the isolation and identification of a range of unusual fungal metabolites with interesting and innovative mechanisms of action. Such compounds have been found to possess a range of properties including anti-diabetic (H. Hussain et al., 2021a), anticancer (Singh & Datta, 2020), antioxidant (Ujam et al., 2021; Vitale et al., 2020), antimicrobial (Al-Fakih et al., 2019; Dalinova et al., 2020; Jakubczyk & Dussart, 2020), immunosuppressive (Duan et al., 2020; Ujam et al., 2021), and several other activities (Hyde et al., 2019). Recent studies by Nguyen et al. (2020) indicate *Monascus purpureus* and its secondary metabolites are capable of reducing cholesterol content in vitro and hence may have a potential application in probiotics. Thus, kingdom fungi have demonstrated to be an exceptional but comparatively less explored source of biologically active compounds with diverse potential. Rising global concerns about environmental consequences of chemical compounds including pesticides have further boosted an era of research in using naturally occurring compounds for our everyday use through industrial production. In this chapter, we discuss the avenues of using fungal metabolites as

bioherbicides against invasive weed species, their associated challenges, and way forward.

6.2 Ecological Role of Fungi and Their Metabolites

Due to wide differences in their chemical structures, the fungal metabolites have different ecological and environmental roles and mechanisms of action. Here we discuss some of the major roles played by fungi and their secondary metabolites in the ecosystems of their occurrence.

6.2.1 Decomposition and Cycling of Organic and Inorganic Materials

Fungi are well known for being decomposers in many ecosystems. Fungal secondary metabolites have been favoured by natural selection for their use in critical environmental conditions like intraspecific and interspecific interactions, growth and development, and abiotic stress (Rohlf & Churchill, 2011). Coprophilous fungi produce secondary metabolites to not only reduce interspecific competition but also to metabolize very complex molecules (Sarrocco, 2016). They also make chemical changes to minerals and make them usable for plants and other organisms; the process is also known as bioweathering. Fungi also help in the breakdown of xenobiotics and detoxification of the environment (Gadd, 2007; Tang et al., 2013).

6.2.2 Interactions with Other Organisms

Fungi also have direct interactions with many different organisms like insects, molluscs, and various other organisms including many plant species, which can be mutualistic or antagonistic (Boddy et al., 2007; Masters & Brown, 1997). Many fungivore organisms obtain high amounts of nutrients from decomposing organic materials or directly from fungal mycelia (Maraun et al., 2003).

6.2.3 Mutualistic Symbiosis with Plants

Many fungi are associated with plant hosts in mutualistic symbiosis, where they increase fitness for each other's survival. Arbuscular mycorrhizal fungi help symbiont plants to accumulate more vitamins and minerals with antioxidant properties as

well as help the plants in obtaining bioavailable nitrogen from mineral associated proteins, where the source of nitrogen is scarce (Baslam et al., 2013; Wang et al., 2020). *Curvularia* sp. was found to increase high temperature tolerance in *Dichanthelium lanuginosum* plant species (Redman et al., 2002). Peramine is a secondary metabolite produced by *Neotyphodium* sp. that deters herbivore insects from feeding on its host plants such as ryegrass (Tanaka et al., 2005).

6.2.4 Competitive Interactions with Different Organisms

There are various strategies to prevent insect competitors for the same resources as fungi by means of chemical warfare. Fungal secondary metabolites were found to reduce immune response in insects and rendered them susceptible to infectious diseases (Fiolka, 2008; Pal et al., 2007). Saprophagous insect larvae are strong competitors to fungi like *Aspergillus*, depending on the same food source. Mycotoxins like Kojic acid and Ochratoxin A were found to be effective in causing insect-specific responses in *Drosophila* spp. and larval death, which helped in the survival of the fungi (Rohlf's & Obmann, 2009; Trienens et al., 2010). *Ophiocordyceps unilateralis* fungus is known to manipulate behaviour of infected *Camponotusleonardi* ants to bite onto plants before the death of the ant at optimal zones for fungal growth (Andersen et al., 2009; Bekker, 2019).

6.3 Secondary Metabolites Biosynthesis

Secondary metabolites are generally synthesized from genes of a single cluster that usually contain one or more central genes for nonribosomal peptide synthetase (NRPS) or polyketide synthase (PKS) and other genes for modifications of the NRPs or PKs (Hertweck, 2009; Strieker et al., 2010; Yu & Keller, 2005).

The NRPSs are multinodular enzymes that catalyze the formation of nonribosomal peptides (NRPs) with very specific function of each domain in a series where each domain activates and adds a single monomer covalently to the substrate to form the NRP. So knowing the assembly and functions of the domains of the NRPs can help in determining the type of NRP to be formed (Fischbach & Walsh, 2006). The NRPs can be functionality divided into three domains: activation or adenylation (A) domain: activates the amino acids, thiolation, or peptidyl carrier protein (PCP) domain: binds the cofactor 4'-phosphopantetheine (4'PP), to which activated amino acids are covalently attached, and propagates the peptide chain being synthesized, condensation (C) domain: peptide bonds are formed, and at the terminal end there is a thioesterase (TE) domain: catalyzes peptide release by either hydrolysis or macrocyclization (Strieker et al., 2010).

PKSs also contain three main domains that are acyltransferase domain: selection and acylation of extender substrate molecules such as malonyl-CoA or

methylmalonyl-CoA, acyl-carrier domain: loading extender molecules, and keto-acyl synthase domain: decarboxylative condensation of extender molecules (Brakhage, 2013).

6.3.1 *Transcriptional Regulation of Secondary Metabolite Gene Clusters*

The secondary metabolite gene clusters are regulated by complex interactions of transcription factors (Pan & Liu, 2018). The transcription factors for such gene clusters are mostly very specific to environmental conditions like carbon and nitrogen sources, pH, temperature, biofilm, light, hypoxic conditions, interspecific signalling, etc. (Bruns et al., 2010; Vödisch et al., 2011; Yin & Keller, 2011). There are two types of transcriptional signalling mechanisms, that is, global regulation: the transcription factor gene is not a part of the gene cluster and transcription factor is also involved in the regulation of other genes that are not part of secondary metabolite gene clusters; on the other hand, there is pathway-specific regulation: the transcription factor is very specific to the promoter sequence of genes from the cluster that are involved in the pathway that is required for the synthesis of the secondary metabolite, and the transcription factor gene may reside within the specific cluster itself or it can be a part of another gene cluster even on a different chromosome (Bergmann et al., 2010; Brakhage, 2013).

6.3.2 *Signalling and Regulations for pH, Light Intensity, Iron Starvation, and Oxidative Stress During Production of Secondary Metabolites Are as Follows*

I. *Ambient pH*

PacC transcription factor (contains zinc finger motifs) in *Aspergillus nidulans* is inactive in ambient acidic pH. In ambient alkaline pH, pal genes (pal A, B, C, F, H, and I) are expressed and the pal genes' products activate the PacC transcription factor by cleaving C-terminal domain and making it accessible to protease and then signalling protease cleaves PacC between residues 493 and 500. Active PacC works as transcriptional activator of alkaline expressed genes such as palD (alkaline phosphatase) and prtA (alkaline protease), and the penicillin biosynthesis genes N-(5-amino-5-carboxypentanoyl)-l-cysteinyld-valine synthase (acvA)40 and isopenicillin N synthase (ipnA) by the interaction of zinc finger motifs to ipnA promoter element 5'GCCAAG3' (Denison, 2000; Díez et al., 2002; Espeso & Penalva, 1996; Then Bergh & Brakhage, 1998).

II. Light Intensity

In *A. nidulans*, phytochrome FphA and other light sensor proteins LreA, LreB interact with LaeA protein in dark condition and nuclear localization of LaeA occurs. LaeA is a methyl transferase that removes methyl group from ninth lysine residue of histone H3 and heterochromatin protein 1 (HepA), and promotes euchromatinization, which positively regulates various secondary metabolites gene clusters like penicillin and sterigmatocystin by epigenetic modifications (Bayram et al., 2008; Sarikaya et al., 2010).

III. Iron Starvation

HapX is a basic region leucine zipper protein that is active in iron-deficient condition in *A. fumigatus* where it binds to CCAAT-binding complex (CBC) made of HapB, HapC, and HapE, and induces siderophore biosynthesis genes that contain a central NRPS gene (Brakhage, 2013; Furukawa et al., 2020).

IV. Oxidative Stress

In presence of H₂O₂, Yap1 is activated by forming disulfide bond with Gpx3 with the help of Ybp1 and the nuclear export signal at the C-terminal is inaccessible. The activated Yap1 binds to Yap1 response element (YRE) TTA(G/C)TAA of anti-oxidant genes like GSH1 (γ -glutamylcysteine synthetase), GPX2 (glutathione peroxidase), TRX2 (thioredoxin), TSA1 (thioredoxin peroxidase) and induces their transcription (Simaan et al., 2019).

6.4 Fungal Formulation–Based Early Bioherbicides

Bioherbicides are weed-control living agents, mostly microorganisms and plants or products derived from them. Mycoherbicides are weed control formulations derived from live fungi or their products. Early research and field success of bioherbicides, particularly mycoherbicides, suggested that a large number of spores or mycelial mass of native, target-specific fungi could be used to transform an ordinarily endemic pathogen into a severe epidemic (Templeton et al., 1979). Commercial mycoherbicides first appeared in the market in the United States in the early 1980s with Devine as the first registered bioherbicide, developed by the Abbott Laboratories, United States, for controlling the milkweed vine. It consisted of liquid formulation containing chlamydospores of fungi *Phytophthora pulmivora*, which was capable of infecting roots of the weed leading to wilting. This was followed by the release of several other products, including Collego and BioMal (TeBeest, 1991), and subsequently several others. Mycoherbicides have in particular attracted global attention due to their weed host specificity, low environmental impact, and mostly cost-effectiveness (Ash, 2010; Ray & Hill, 2013). Development of microbial herbicides can be especially beneficial against the herbicide-resistant weeds.

However, the efficacy of these products is often restricted as the growth, sporulation, dissipation of infection, disease development, and disease-causing potential of

the pathogen can be overblown by daily changes in the environment under field conditions impacting weed management drastically (Ray & Hill, 2016). Several studies (Auld & Say, 1999; Hetherington et al., 2002; Zhang & Watson, 1997) show that the dew period requirement in terms of duration, temperature, timing, and frequency in most fungi is perhaps the predominant impediment to the development and deployment of mycoherbicides. Furthermore, after application, most fungal spores die within a short span of time: the infection process gets aborted midway due to this phenomenon. Also challenges due to increasing herbicide resistance in weeds (Bourdôt et al., 2007; Hussain et al., 2021b) and nontarget environmental effects of chemical herbicides have led to increasing research in finding their biological and eco-friendly alternatives (Bourdôt et al., 2007; Gurusubramanian et al., 2008; Ray et al., 2008). Thus, synthesis of novel agrochemicals like mycoherbicides has thus attracted much attention from researchers preferring eco-friendly pest control options and sustainable organic farming.

6.5 Fungal Metabolites as Bioherbicides

Fungal metabolites also provide a prosperous resource for developing environmentally viable and sustainable mycoherbicides. Decades of research and biotechnological advances have resulted in a surfeit of knowledge on extraction of interesting compounds through fractionation and characterization from a range of fungi with herbicidal potential (McLean, 1996; Ndam et al., 2014). The fairly high degradability of these microbial metabolites causes less environmental problems in the soil ecosystem or bioaccumulation of toxic compounds in the food. Thus, the primary goal of these extensive researches into these naturally existing microbial phytotoxins is to curtail the use of environmentally damaging chemical herbicides and facilitation of nonchemical farming, enhancing agricultural productivity and ecological systems in a sustainable way.

A few potential toxins of fungal origin exploited for the management of weeds include A-AL toxin from *Alternaria alternata* against prickly sida and range of other broadleaf weed species (Abbas et al., 1995; Meena & Samal, 2019), Fumonisin from *Fusarium moniliforme* against Hemp sesbania (Ismaiel & Papenbrock, 2015; Nelson et al., 1993), Ophiobolins from *Drechslera oryzae* against sickle pod (Evidente et al., 2006; Fumio Sugawara et al., 1988), Pyrenophorol from *Drechslera avenae* against *Avena sterilis* L (Kastanias & Chrysayi-Tokousbalides, 2000) etc. A list of some interesting secondary metabolites has been given in Table 6.1. Zhao et al. (2021) while clarifying the role of phytotoxic substances of fungi *Bipolaris setariae* observed several ophiobolins associated with strong phytotoxicity toward green foxtail (*Setaria viridis*). Among these one of the ophiobolins (6-epi-Ophiobolin A) interestingly showed cytotoxicity against five kinds of human cancer cells. This study opens avenues for parallel development of potential compound both as herbicide as well as an antitumor drug.

Table 6.1 List of toxins of fungal origin exploited for the management of some weed

| Toxin | Source fungal species | Target weed | References |
|-------------------|--|--|--|
| A-AL toxin | <i>Alternaria alternata</i> | Prickly sida (<i>Sida spinosa</i>), <i>Lemna minor</i> L., Northern Jointvetch, <i>Amaranthus retroflexus</i> , <i>Xanthium strumarium</i> | Abbas et al. (1995) |
| Alternariol | <i>Alternaria alternata</i> | <i>Amaranthus retroflexus</i> | De Souza et al. (2013) and Tang et al. (2020) |
| Ascaulitoxin | <i>Ascochyta caulina</i> | <i>Chenopodium album</i> L. | Evidente et al. (1998) |
| AT-toxin | <i>Alternaria longipes</i> | Tobacco (<i>Nicotiana tabacum</i>) | Nishimura and Kohmoto (1983) |
| Bipoloroxin | <i>Bipolariscyanodontis</i> | <i>Cynodondactylon</i> | Sugawara et al. (1985) |
| Brefeldin A | <i>Alternaria zinniae</i> | <i>Xanthium occidentale</i> | Vurro et al. (1998) |
| Colletotrichinin | <i>Colletotrichum tabacum</i> | <i>Nicotiana tabacum</i> | Duke et al. (1991) |
| Curvulin | <i>Dreschslera indica</i> | <i>Amaranthus spinosus</i> L. and <i>Portulaca oleraceea</i> | Kenfield et al. (1989) |
| Eremophilanes | <i>Bipolariscyanodontis</i> | Bermudagrass (<i>Cynodondactylon</i>) | Sugawara et al. (1985) |
| Fumonisin | <i>Fusarium moniliforme</i> | Anoda cristata, Sickle pod (<i>Cassia obtusifolia</i>), Hemp sesbania (<i>Sesbania exaltata</i>) | Nelson et al. (1993) |
| Maculosin | <i>Alternaria alternata</i> | Spotted Knapweed (<i>Centaurea maculosa</i>) | Meena and Samal (2019) and Stierle et al. (1988) |
| Ophiobolins | <i>Dreschsleraoryzae</i> , <i>D. sorghicola</i> <i>D. gigantea</i> <i>Bipolaris setariae</i> | <i>Sorghum</i> , Sickle pod (<i>Cassia obtusifolia</i>) crabgrass (<i>Digitariasanguinalis</i>) Green foxtail (<i>Setaria viridis</i> (L.) P.Beauv.) | Sugawara et al. (1988), Evidente et al. (2006), and Zhao et al. (2021) |
| Porriolide | <i>Alternaria porri</i> | Lettuce and Stoneleek seedling | Suemitsu et al. (1993) |
| Radicinin | <i>Cochliobolusaustriensis</i> <i>Pyricularia grisea</i> | Buffelgrass (<i>Cenchrus ciliaris</i>) | Masi et al. (2019) |
| Tentotoxin | <i>Alternaria alternata</i> | Johnson grass (<i>Sorghum halepense</i>), Jimson weed (<i>Datura stramonium</i> L), <i>Sida spinosa</i> | Duke (1986) |
| Tentuaazonic acid | <i>Alternaria alternata</i> | <i>Datura innoxia</i> | Umetsu et al. (1973) |

(continued)

Table 6.1 (continued)

| Toxin | Source fungal species | Target weed | References |
|--------------|----------------------------|---|------------------------------------|
| Pyrenophorol | <i>Drechsleraavenae</i> | <i>Avenasterilis</i> L | Kastanias and Tokousbalides (2000) |
| Viridiol | <i>Gliocladium virens</i> | <i>Amaranthus retroflexus</i> | Jones et al. (1988) |
| Zinniol | <i>Alternaria cichorii</i> | Russian knapweed (<i>Acroptilon repens</i>) | Stierle et al. (1993) |

6.6 Future Prospects of Fungal Metabolites as Bioherbicides and Entrepreneurship Possibilities

Numerous mycotoxins with phytotoxic potential with the possibility of being developed as bioherbicide have been identified over the last four decades. Most of the fungal metabolites have different modes of actions, biochemical reactions, and signalling mechanisms leading to the development of diseases symptoms and possible death in their respective host plants (Meena & Samal, 2019). These provide opportunities for elucidating new mechanisms of herbicide action not yet discovered by traditional pesticide discovery methods, thereby providing tools for combating environmental issues and herbicide resistance. Further commercial interest in these metabolites has mostly caused researchers to largely look into these natural compounds for structural elucidation and industrial application, while very little effort has been given to understand the causes and consequences or the primary role and impact of these metabolites. Also, several challenges exist in developing these biologically active metabolites into bioherbicides. One of the major challenges include dealing with biosynthesis of low quantities of the bioactive compound naturally or under the influence of growing conditions including, impact of media, and environmental factors like temperature (Rai et al., 2021). Thus, more studies shall be interesting to understand their ecological role in nature. However, rapid advent and magnitude of possible utilization of techniques, like metabolomics, meta-proteomics, meta-genomics, meta-transcriptomics, next-generation sequencing, etc., have vastly increased our ability to untap the infinite fungal resources. Though over the last few decades, metabolic engineering of fungi has led to understanding biosynthetic pathways leading to the production of large quantities of secondary metabolites for commercial application, yet an extensive scope of finding bioherbicidal agents still remains less explored. The continuing search for herbicidal agents from fungi is extremely necessary to find the possible ways to have safe and more effective weed management while reducing the use of chemical pesticides. With the advancement of expanding knowledge on fungi and their metabolites, formulation development research and field trials are needed. More researchers of the present and future generations need to take up these technological and scientific challenges for a pesticide pollution-free world.

6.7 Conclusion

The problem of invasive weeds in contemporary intensive agriculture under increasing anthropogenic population pressure and changing climatic conditions makes plant protection inevitable. Further aggravating environmental issues have ushered in extensive research needs into eco-friendly bio-based solutions to invasive weed management. Fungal diversity offers a rich natural resource with possibilities of providing for human needs, in a sustainable way, with immense possibilities. Fungal metabolites thus have promising potential as bioherbicides.

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Chapter 7

Lichenized Fungi as Significant Source of Pharmaceuticals: Possibilities and Limitations for Entrepreneurship Development



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7.1 Introduction

The practices related to the traditional use of plant-based medicines pertained to many parts of the world among which China is at the top followed by India and other nations (Liu, 2010). This could be attributed to their efficacy without any side effect; hence, the demand for herbal medicines at a high rate never declined. The trend continues at present and is expected to upsurge in the near future too (Verma & Singh, 2008).

Lichenized fungi or lichens are one such dual-natured biological entities composed of fungi and algae (cyanobacterium or green alga), known as mycobiont and photobiont, respectively. They are the rich repository of secondary metabolites, which are produced exclusively in them to perform special functions. Some of the important secondary metabolites reported in lichens are amino acid derivatives,

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sugar alcohols, aliphatic acids, macrolytic lactones, monocyclic aromatic compounds, quinines, chromones, xanthenes, dibenzofurans, depsides, depsidones, dapsones, terpenoids, steroids, carotenoids, and diphenyl ethers (Bhattacharyya et al., 2016). These exhibit a range of pharmaceutical activities, including antimicrobial, antiproliferative, antioxidant, antiviral, anti-inflammatory, and further allelopathic, antiherbivore, and photoprotective activities (Goga et al., 2018; Wink, 2018).

7.2 Applications of Lichenized Fungi

7.2.1 Pharmaceuticals

Due to the presence of secondary metabolites, lichens are being explored in pharmaceutical industries. Streptomycin and 2-propanol are lichen derivatives isolated from *Parmotrema tinctorum*. These exhibit antibacterial and antifungal activities against human pathogenic bacterial strains and also against a wide range of fungi (Anjali et al., 2015).

Different solvent extracts of lichens (*Cladonia furcata*, *Parmelia caperata*, *Parmelia pertusa*, *Hypogymnia physodes*, *Umbilicaria polyphylla*, *Lasallia pustulata*, *Parmelia sulcata*, *Umbilicaria crustulosa* and *Umbilicaria cylindrica*) showed antibacterial and antifungal activities. However, antioxidant, antipyretic, antiviral, antitumor, and analgesic effects of these lichens have also been reported (Mitrović et al., 2011).

Chharila, an Indian lichenized drug composed of *Parmelia perlata*, *Parmelia perforata*, and *Parmelia sancti-angelii*, is reported to be effective against dyspepsia, spermatorrhoea, amenorrhoea, calculi, diseases of blood and heart, stomach disorders, enlarged spleen, bronchitis, bleeding piles, scabies, leprosy, excessive salivation, soreness of throat, tooth-ache, wounds, and general pain; it can also be used as astringent, resolvent, laxative, and carminative (Nayaka et al., 2010).

Usnea longissima is known for its wounds healing properties and is effective against tumor (Crockett et al., 2003).

In Brazil, since ancient times, *Parmelinella salacinifera*, *Heterodermia galactophylla* and *Parmotrema wrightii* were used to treat common digestive track problems (Londoño-Castañeda et al., 2017).

Usnea ghattensis shows antimicrobial activity against gram-negative and gram-positive pathogenic bacteria (*Staphylococcus aureus*, *Streptococcus faecalis*, *Bacillus cereus*, *Escherichia coli*, *Pseudomonas aeruginosa*, and *Salmonella typhimurium*). *Lecanora muralis*, *Peltigera polydactyla*, *Ramalina farinacea*, and *Xanthoria elegans* also possess antibacterial activity (Srivastava et al., 2013).

Lichen compounds such as Barbatic Acid, Barbatolic Acid, Caperatic Acid; Diffractic Acid, Divaricatic Acid, Gyrophoric Acid, Hypostictic Acid, Lobaric Acid, Lecanoric Acid, Lichexanthone, Lobastin, Nortictic Acid, Olivetoric Acid,

Physodic Acid, Perlatolic Acid, Protolichesterinic Acid, Protocetraric Acid, Psoromic Acid, Potassium Usnate, Retigeric Acid, Ramalin, Salazanic Acid, Sekikaic Acid, Usnic Acid, and Vulpinic Acid show anticancer, antimicrobial, antioxidant, anti-inflammatory, antiproliferative and anti-diabetic properties (Solárová et al., 2020).

Evernia furfuracea used for medicinal purposes is reported to be effective against jaundice, rabies, and cough. *Cladonia*, *Evernia*, *Lobaria*, *Parmelia*, *Peltigera*, *Pertusaria*, *Physica*, *Rocella*, *Usnea*, and *Xanthoria* are used as herbal medicines (Zambare & Christopher, 2012).

Usnea barbata is effective against *Bacillus subtilis* and *Pseudomonas fluorescens*. Besides, it also possesses antioxidant properties (Bazarmova et al., 2018).

Anaptychia ciliaris, *Cetrelia olivetorum*, *Lecanora muralis*, *Peltigera polydactyla*, *Peltigera praetextata*, *Ramalina farinacea*, *Rhizoplaca melanohphthalma*, *Umbilicaria vellea*, *Xanthoria elegans*, *Xanthoria parietina*, *Xanthoparmelia tinctoria* are known for their antibacterial potential against *Escherichia coli*, *Pseudomonas aeruginosa*, *Bacillus subtilis*, *Klebsiella pneumonia*, *Staphylococcus aureus*, and *Staphylococcus* (Karagoz et al., 2009).

Antimicrobial activity against some bacteria viz. *Escherichia coli*, *Pseudomonas aeruginosa*, *Enterococcus faecalis*, *Staphylococcus aureus*, and a fungus *Candida albicans* was shown by *Melanohalea exasperata*, *Physcia aipolia*, *Usnea florida*, *Usnea subfloridana* and *Xanthoria parietina*. Antimicrobial activity by lichens was reported both against gram positive and gram-negative strains (Çobanoğlu et al., 2016).

Usnic acid a derivative of *Cladonia* sp. is proved to be effective against lung cancer. On the other hand, many lichens are used as antibiotics against a number of bacteria (Shrestha & St. Clair, 2013).

A total of 153 active chemical constituents have been reported in 118 different species of genus *Ramalina*, which have antimicrobial, antioxidant, antiviral, antitumor, cytotoxic, anti-inflammatory, antifungal, fungicidal, anti-helminth, anti-Schistosoma, and larvicidal properties (Moreira et al., 2015).

7.2.2 Pesticides

Chemical pesticides are used to kill pests and insects to protect crops from destroying. On the other hand, many natural products have substituted these chemical-based pesticides without affecting soil, crops, and beneficial insects. Lichens are also being used for this purpose, and crop is treated either with the extract of single species or more than one species in different concentrations with different solvents. *Lecanora muralis* (Schreb.) Rabenh. *Letharia vulpina* (L.), Hue, and *Peltigera rufescens* are used as insecticides against *Sitophilus granarius* (Emsen et al., 2016).

7.2.3 Dyes

Dye extracted from lichens is used to color attires, food and also applied on body parts. Production of dye in lichens is due to many secondary metabolites present in them like gyrophoric, lecanoric acid, umbilicic acids, usnic acid, atranorin, chloroatranorin, salazinic acid, and parietin. Some of the dye-producing lichens are *Flavopunctelia*, *Flavoparmelia*, *Cladonia*, *Parmelia*, *Umbilicaria*, *Xanthoria*, *Ochrolechia*, *Hyperphyscia*, *Hypogymnia*, *Dermatocarpon*, and *Parmotrema* (Shaheen et al., 2019).

7.2.4 Bioremediation

Many reports are available showing that the lichens are used for uptake of heavy metals and air pollutants from the atmosphere.

Numerous scientific records are available based on multiple uses of lichens (Table 7.1 and Fig. 7.1). This can be referred to while looking for new avenues to explore possibilities regarding the establishment of lichen-based enterprises.

7.3 Exploring Lichens for Entrepreneurship Development

The numerous benefits associated with lichens make them promising entities to be explored commercially. Entrepreneurship development focusing on these aspects of lichenized fungi is needed so as to tap these important bioresources.

As they are harvested from their natural habitat, their distribution in nature is already under threat. Therefore, lichen farming or cultivation can be a suitable option for business for which only a little space is required. It can be profitable with low capital investment. But to ensure good returns, both study and experience are required. Technical training both at national and international levels can be imparted to gain expertise. Then, it can even be perfect option for the unskilled as well.

For sustainable entrepreneurship, budgetary requirements need to be worked out, focusing clean and green products and target market such as domestic or export.

Certain limitations such as their slow growth rate poses a challenge, which needs to be worked upon. Ensuring safe environment for workers and development of harmless lichen products for end users also holds importance.

Genetic engineering techniques leading to industrial large-scale production of lichen-based natural products by modifying natural pathways to increase productivity to meet demands both globally as well as locally for lichen-based products.

However, researching the mainstream market and strategic use of these inconspicuous forms can definitely reap maximum benefits and fulfill demand for these pharmaceutical products in the market commercially.

Table 7.1 Bioactive metabolites and uses associated with lichenized fungi

| S. no. | Lichen | Bioactive metabolites | Uses | References |
|--------|---|---|--------------------------|---|
| 1. | <i>Bacidea stipitata</i> | Atranorin | Anticancer | Shrestha and St. Clair (2013) |
| 2. | <i>Bryoria lactinea</i> (Nyl.) Brodo and D. Hawksw. | Fumarprotocetraric acid | Biofuel | Shukla et al. (2014) |
| 3. | <i>Bryoria</i> sp. | | Immunosuppressive | Hwang et al. (2017) |
| 4. | <i>Bulbothrix setschwanensis</i> (Zahlbr.) Hale | Salazinic acid | Biofuel | Shukla et al. (2014) |
| 5. | <i>Cetraria braunsiana</i> (Mull. Arg.) W. Culb. and C. Culb. | Alectoronic and α -collatolic acid | Biofuel | Shukla et al. (2014) |
| 6. | <i>Cladia retipora</i> | Usnic acid | Antiviral/ Cytotoxic | Perry et al. (1999) |
| 7. | <i>Cladonia deformis</i> | Zeorin | Bioremediation | Hämäläinen et al. (2015) and Pawlik-Skowrońska and Bačkor (2011) |
| 8. | <i>Cladonia furcata</i> | Depside, depsidones, dibenzofurane | Antibacterial/Antifungal | Mitrović et al. (2011) and Kosanic et al. (2018) |
| 9. | <i>Cladonia lepidophora</i> | Usnic acid | Anticancer | Shrestha and St. Clair (2013) |
| 10. | <i>Cladonia rangiferina</i> (L.) Nyl | Atranorin Fumarprotocetraric acid, Isousnic acid | Antibacterial | Smith (1932), Richardson (1974), Yoshimura et al. (1994), and Fraser (2006) |
| 11. | <i>Cladonia substellata</i> | Usnic acid | Insecticidal | Sachin et al. (2018) |
| 12. | <i>Cladonia sulphurina</i> | Usnic and Squamatic acid | Bioremediation | Hämäläinen et al. (2015) and Pasternak et al. (2015) |
| 13. | <i>Cornicularia aculeata</i> | Protolicheterinic acid | Anticancer | Shrestha and St. Clair (2013) |
| 14. | <i>Evernia divaricata</i> (L.) Ach. | β -pinene, α -pinene, limonene, α -phellandrene, camphene, and p-cymene | Antibacterial | Yuan et al. (2010) and Goga et al. (2018) |

(continued)

Table 7.1 (continued)

| S. no. | Lichen | Bioactive metabolites | Uses | References |
|--------|--|---|--------------------------|---|
| 15. | <i>Evernia prunastri</i> (L.) Ach | Atranorin, Chloroatranorin Evermic acid, Usnic acid | Astringent | Lopez et al. (2006) and Yoshimura et al. (1994) |
| 16. | <i>Everniastrum nepalense</i> (Taylor) Hale | Salazinic and protolichesterinic acid | Biofuel | Shukla et al. (2014) |
| 17. | <i>Flavopunctelia soredica</i> (Nyl.) Hale | Lecanoric acid | Biofuel | Shukla et al. (2014) |
| 18. | <i>Heterodermia diademata</i> (Taylor) D. D. Awasthi | Zeorin | Biofuel | Shukla et al. (2014) |
| 19. | <i>Heterodermia leucomela</i> (L.) Poelt | 3,6-Dimethyl-2-hydroxy-4-methoxybenzoic acid, Zeorin, norstictic, salazinic acid, and triterpenoids | Insecticidal/biofuel | Shukla et al. (2014) and Sachin et al. (2018) |
| 20. | <i>Hypogymnia physodes</i> | Depsidones, depsides, usnic acid | Antibacterial/antifungal | Mitrović et al. (2011) and Ranković et al. (2009, 2014a, b) |
| 21. | <i>Lasallia pustulata</i> | gyrophoric acid, arabitol, mannitol, and umbilicarin | Antibacterial/antifungal | Ranković et al. (2007) and Mitrović et al. (2011) |
| 22. | <i>Lecanora muralis</i> | Phenolics and flavonoid | Antibacterial | Rankovic et al. (2014a) and Srivastava et al. (2013) |
| 23. | <i>Leparia tomentosa</i> | (+)-Usnic acid | Insecticidal | Sachin et al. (2018) |
| 24. | <i>Leproloma sipmanianum</i> | 3,6-dimethyl-2-hydroxy-4-methoxybenzoic acid | Insecticidal | Sachin et al. (2018) |
| 25. | <i>Letharia vulpina</i> | Atranorin and vulpinic acid | Insecticidal | Sachin et al. (2018) |
| 26. | <i>Loberia retigera</i> (Bory) Trev. | Triterpenoids and thelephoric acid | Biofuel | Shukla et al. (2014) |
| 27. | <i>Ochrolechia deception</i> | Variolaric acid | Anticancer | Shrestha and St. Clair (2013) |
| 28. | <i>Pseudocyphellaria homoeophylla</i> | Usnic acid | Antiviral/ Cytotoxic | Perry et al. (1999) |
| 29. | <i>Parmelia subthomsonii</i> D. D. Awasthi | Atranorin, alectronic and α -collatolic acid | Biofuel | Shukla et al. (2014) |

| | | | | |
|-----|--|---|--|--|
| 30. | <i>Parmelia caperata</i> | salazinic acid, usnic acid, depsidone, atranorin, chloroatranorin | Antibacterial/antifungal/antioxidant/cytotoxic | Manojlović et al. (2013) and Mitrović et al. (2011) |
| 31. | <i>Parmelia petrusa</i> | atranorin and salazinic acid | Antibacterial/antifungal | Mitrović et al. (2011) and Ranković et al. (2009) |
| 32. | <i>Parmelia sulcata</i> | Salazinic acid | Antibacterial/antifungal | Mitrović et al. (2011) and Candan et al. (2014) |
| 33. | <i>Parmelinella wallichiana</i> (Taylor) Elix and Hale | Salazinic and consalazinic acid | Biofuel | Shukla et al. (2014) |
| 34. | <i>Peltigera polydactyla</i> | usnic acid, phenolic compounds, triterpenes, steroids, anthraquinones, depsides, depsidones, and dapsones | Antibacterial | Srivastava et al. (2013) and Kosanić and Ranković (2014) |
| 35. | <i>Protousnea magellanica</i> | Diffractic acid | Anticancer | Shrestha and St. Clair (2013) |
| 36. | <i>Protousnea malacea</i> | Divaricatic acid | Anticancer | Shrestha and St. Clair (2013) |
| 37. | <i>Pseudocyphellaria glabra</i> | Usnic acid | Antiviral/cytotoxic | Perry et al. (1999) |
| 38. | <i>Psoroma dimorphum</i> | Vicamicin | Anticancer | Shrestha and St. Clair (2013) |
| 39. | <i>Psoroma pallida</i> | Vicamicin | Anticancer | Shrestha and St. Clair (2013) |
| 40. | <i>Pyxine consocians</i> | Cabraleadiol monoacetate, 4-O-methylerythrochlorophaeic acid, lichexanthone | Insecticidal | Sachin et al. (2018) |
| 41. | <i>Ramalina conduplicans</i> Vain. | Usnic acid, sekikaic acid aggregate and salazinic acid | Biofuel | Shukla et al. (2014) |
| 42. | <i>Ramalina farinacea</i> | usnic acid, phenolic compounds, triterpenes, steroids, anthraquinones, depsides, depsidones, and dapsones | Antibacterial | Srivastava et al. (2013) and Kosanić and Ranković (2014) |
| 43. | <i>Rhizoplaca melanophthalma</i> | Protolichesterinic acid | Anticancer | Shrestha and St. Clair (2013) |
| 44. | <i>Stereocaulon alpinum</i> | Lobaric acid | Anticancer | Shrestha and St. Clair (2013) |

(continued)

Table 7.1 (continued)

| S. no. | Lichen | Bioactive metabolites | Uses | References |
|--------|-------------------------------|--|---|--|
| 45. | <i>Umbilicaria crustulosa</i> | methyl orsellinate, lecanoric acid, crustinic acid, gyrophoric acid and atranorin | Antibacterial/antifungal | Mitrović et al. (2011) and Zlatanović et al. (2017) |
| 46. | <i>Umbilicaria cylindrica</i> | norstictic acid, methyl- β -orcinol carboxylate, ethyl haematommate, atranorin and usnic acid | Antibacterial/antifungal | Manojlovic et al. (2012) |
| 47. | <i>Umbilicaria hirsuta</i> | Gyrophoric acid | Anticancer | Shrestha and St. Clair (2013) |
| 48. | <i>Umbilicaria polyphylla</i> | Phenol, flavonoid | Antibacterial/antifungal | Kosanic and Rankovic (2011) and Mitrović et al. (2011) |
| 49. | <i>Usnea ghattensis</i> | Usnic acid, norstictic acid | Antibacterial/antioxidant | Verma et al. (2008) and Srivastava et al. (2013) |
| 50. | <i>Usnea longissima</i> | Usnic acid, barbatic acid, diffractaic acid, glutinol, longissiminone A, longissiminone B, salazinic acid, protocetraric acid, evernic acid, 4-O-demethyl-barbatic acid Diffractaic acid and usnic acid | Antibacterial/antifungal/ antitumor/insecticidal | Sachin et al. (2018), Crockett et al. (2003), and Zlagic et al. (2018) |
| 51. | <i>Xanthoria elegans</i> | Anthraquinone | Antibacterial | Stocker-Wörgöter (2008) and Srivastava et al. (2013) |
| 52. | <i>Xylopsora caradocensis</i> | - | Bioremediation | Hämäläinen et al. (2015) |
| 53. | <i>Xylopsora friesii</i> | Friesic acid | Bioremediation | Hämäläinen et al. (2015) and Elix (2005) |

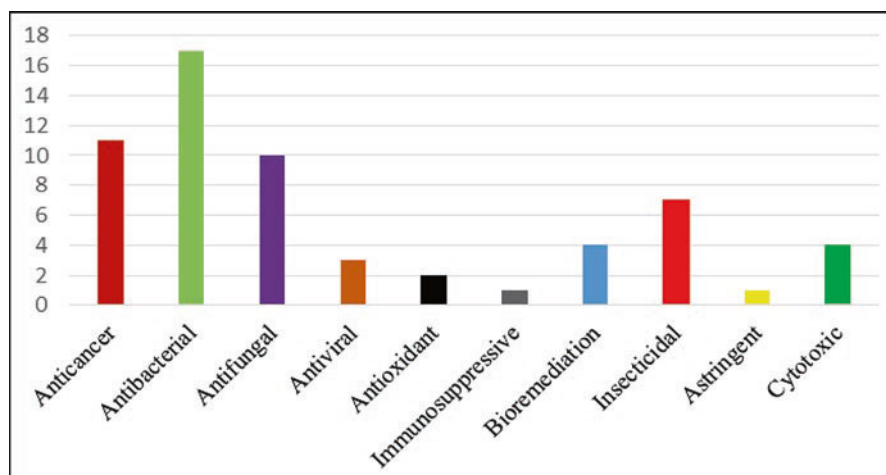


Fig. 7.1 Pharmacological uses of lichens

7.4 Conclusions

The multiple roles that lichenized fungi play in our environment are unmatched with the other living organisms. Every best effort must be done so as to harness their potential. The fundamental and applied science offers the best solution to the lichen-based economy. It opens new avenues and in the light of their contribution to 7 out of 17 United Nations Sustainable Development Goals, their role becomes more important for future, which needs to be realized, researched, tapped, and explored.

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Chapter 8

Protocols for Extraction, Isolation, and Purification of Secondary Metabolites of Mushroom and Its Applications



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8.1 Introduction

Mushrooms, the macroscopic fungi, harbor miraculous potentials in their filamentous fruiting bodies that significantly contribute to the nutrition of both animal and plant kingdoms (Feeney et al., 2014). Thousands of secondary metabolites of agricultural, medicinal, nutritional, and industrial application have been identified from fungi (Keswani et al., 2014). Basidiomycetes contribute about 23% of the total natural bioactive metabolites known today (Bérdy, 2012). These mushroom-derived metabolites are indubitable stars of modern medicine unveiling massive numbers of

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bioactivities like antifungal, antiviral, antibacterial, antitumor, and immunomodulating activities (Gargano et al., 2017).

These secondary metabolites are at present the topic of great research interest, although their part of extraction in phytochemical or biological researches presents specific challenges that are worth mentioning. Successful extraction and isolation of desired compounds begin with careful selection and preparation of mushroom samples. Further, literature review will be helpful to choose the appropriate protocol for a particular class of secondary metabolite or mushroom species. During the extraction process, it is vital to limit interference from components that may coextract with the target compounds and must take care to prevent decomposition of important metabolites and so on. This chapter presents an outline of the process of extraction, isolation, and identification of mushroom secondary metabolites. In addition, some bioactive secondary metabolites derived from *Ganoderma* spp., *Lentinula edodes*, and *Flammulina velutipe* are mentioned, along with their applications.

8.2 Protocol for Extraction, Isolation, and Purification of Secondary Metabolites of Mushroom

8.2.1 Selection, Collection, and Identification of Mushroom Material

The reproducibility of a phytochemical research is directly affected by the selection, collection, and identification of mushroom material. Any imprecision at this stage of an investigation might reduce the scientific value of the research when searching for a particular compound or metabolite. Specific secondary metabolites differ both qualitatively and quantitatively among members of single species, as well as among members of a population. Apart from the phytochemical considerations, norms related to intellectual property rights are to be taken care of when samples or extracts will cross international borders (Cantley, 1997).

8.2.2 Drying and Grinding

To begin with, mushrooms should be dried at temperatures below 30 °C in order to avoid decay of thermolabile compounds. Similarly, direct sunlight should be avoided to minimize the chemical transformations ensuing from exposure to ultraviolet radiation. When a material is required for research, it is recommended to extract it instantly using organic solvents, such as ethanol in order to deactivate enzymes present in the material. Small quantities of fungal material can be crushed using a mortar and pestle, or electric grinder. In the case of large quantities, milling

can be best carried out in industrial-scale comminution apparatus. Proper grinding decreases the quantity of solvent required for extraction and improves the efficacy of extraction.

8.2.3 *Extraction, Isolation, and Purification*

Some conventional processes of extraction and isolation of secondary metabolites include Soxhlet extraction, hydro-distillation, maceration, pressing, percolation, and infusion. In order to isolate a specific compound in a purified form, it is essential to add chemicals while extraction. Most of the water-insoluble compounds are extracted with organic solvents, such as chloroform, methanol, or ethanol. However, the requirement of large quantities of solvents for efficient extraction along with the high temperatures, sometimes, causes degradation of target molecules with fractional loss of volatile compounds (Bubalo et al., 2018). Consequently, the advanced methods of extraction are envisioned to make use of nontoxic solvents, like ethanol, water, and carbon dioxide. Some most promising and competent extraction techniques include ultrasound-assisted extraction, microwave-assisted extraction, supercritical fluid extraction, subcritical water extraction, and accelerated solvent extraction (Roselló-Soto et al., 2016) (Fig. 8.1). The use of standard extraction techniques to obtain valuable extracts rich in secondary metabolites from different mushroom species has been expansively described by various workers (Ruthes et al., 2015; Kaewnarin et al., 2016; Boonsong et al., 2016). Kimatu et al. (2017) obtained a protein-rich extract with conventional techniques in order to produce mushroom protein hydrolysates. Hassainia et al. (2018) isolated chitin from the fruiting body of *A. bisporus* with alkaline treatment of 1 M NaOH solution at 80 °C for 2 h under optimum conditions. Maeng et al. (2016) prepared the phenolic-rich extracts of mushroom *Coriolus versicolor* means of microwave-assisted extraction and optimized a central composite experimental design.

Smiderle et al. (2017) compared the two powerful extraction techniques, microwave-assisted extractions and pressurized liquid extractions, to obtain β -D-glucans from *Pleurotus ostreatus* and *Ganoderma lucidum* fruiting bodies. Although both methods were efficient, quick, and easy, but its postextraction treatments required some extra separation steps. Another recent advancement in extraction strategies involves molecular imprinting. Hashim et al. (2016) applied this technique to extract ergosterol from *Ganoderma tsugae*. Moreover, they reported an indistinct increase in the ergosterol amount extracted from *G. tsugae* as compared to that obtained from conventional extraction techniques. Overall, representation of extraction, isolation, and identification of metabolites can be summarized as represented in Fig. 8.2.

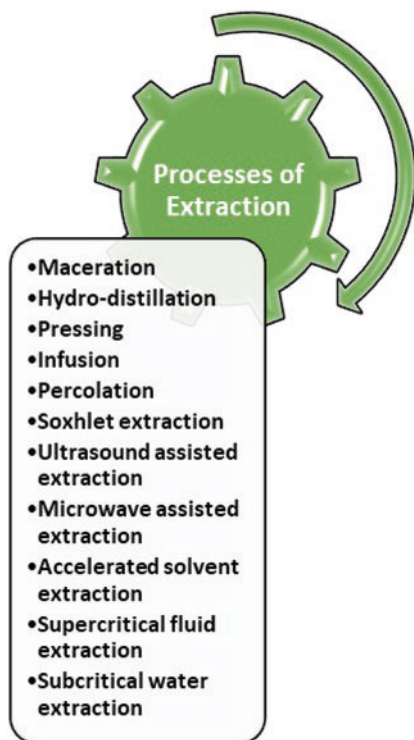


Fig. 8.1 Methods of extraction process. (Roselló-Soto et al., 2016)

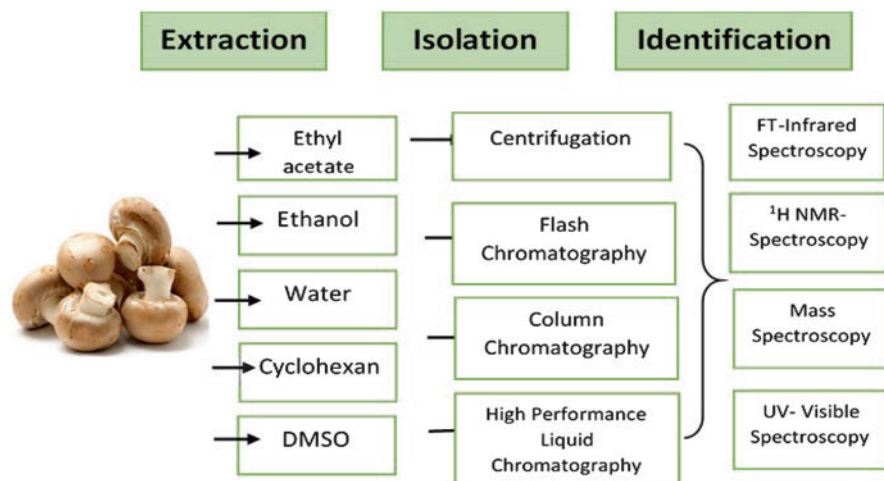


Fig. 8.2 Mushroom metabolites: extraction, isolation, and identification methods

8.2.3.1 Sequential Extraction Methods to Obtain Different Extracts from Shiitake Mushrooms (Morales et al., 2018)

A schematic outline of the extraction methods utilized to obtain different extracts from shiitake mushrooms is given in Fig. 8.3 (Morales et al., 2018):

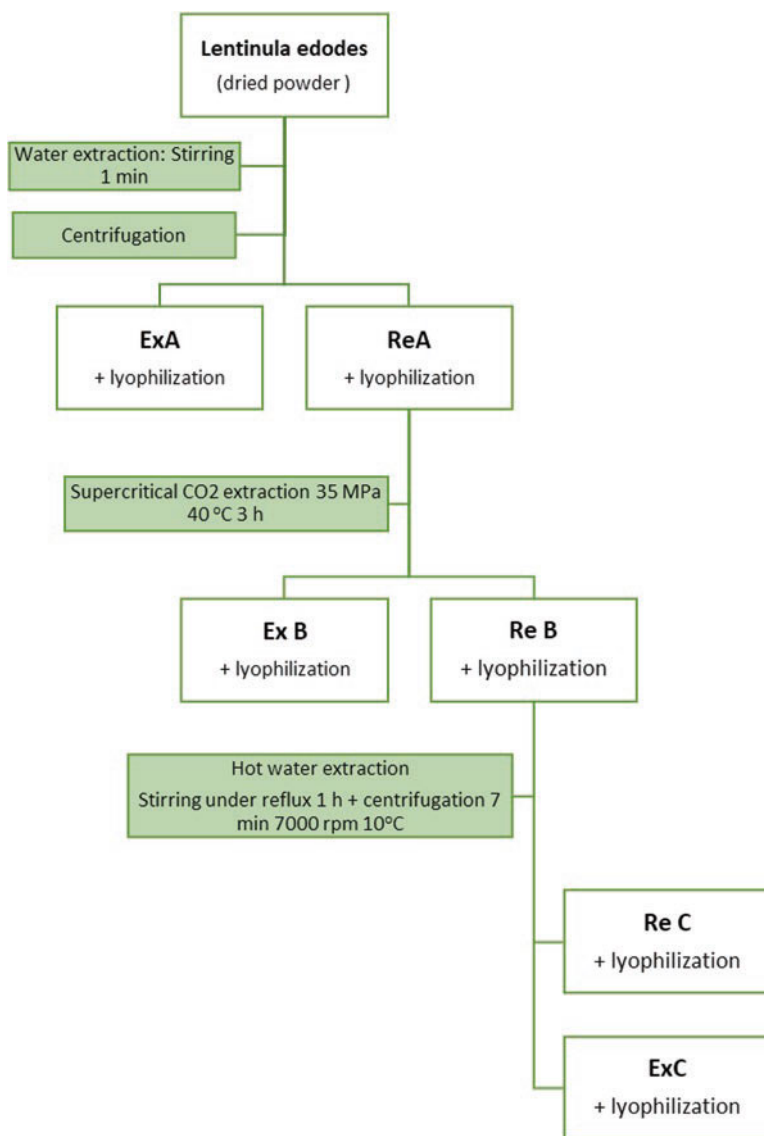


Fig. 8.3 Outline of the sequential extraction method utilized to obtain different extracts from shiitake mushrooms. (Morales et al., 2018).

- Step 1.** Collect mushroom sample and dry it.
- Step 2.** Mix mushroom powder with water (50 g/L) at room temperature and stir vigorously.
- Step 3.** Centrifuge the mixture, thus obtained (7 min, 7000 rpm, 10 °C) and separate the supernatant (ExA) from the residue (ReA). Now, freeze both fractions lyophilizer.
- Step 4.** Grind freeze-dried ReA and sieve until particles size <0.5 mm and submit it to
- Step 5.** Supercritical fluid extraction (SFE): Mix ReA (253 g) with 1.9 kg of 5 mm diameter stainless steel spheres (a ratio 1:1 (v/v) extract: spheres) in extraction cell.
- Step 6.** Collect the extracted material in two different separators.
- Step 7.** Precipitate both the extracted compounds.
- Step 8.** At the end of the extraction process, drag the fractions with ethanol and immediately submit them to concentration until dryness on a rotary vacuum evaporator.
- Step 9.** Dried extract (ExB) will be stored at −20 °C till further analysis and separate the nonextracted residual material (ReB) from steel spheres by sieving.
- Step 10.** Hot water extraction (98 °C) of ReB (100 g/L) in a flask under vigorous stirring.
- Step 11.** Separate the soluble (ExC) and nonsoluble (ReC) fractions by centrifugation and freeze-dry.

8.3 Extraction and Isolation of Metabolites from *Tapinella atrotomentosa*

Steps involved in the extraction and isolation of secondary metabolites from *T. atrotomentosa* can be summarized as follows (Béni et al., 2018):

- Step 1.** Collect the mushroom material, that is, *T. atrotomentosa*.
- Step 2.** Store the fruiting bodies at −20 °C until processing.
- Step 3.** Make Methanol extract (11.5 L) of the fruiting bodies of *T. atrotomentosa* (2 kg) and concentrate it.
- Step 4.** Dissolve the dry methanol extract (90.0 g) in 50% aqueous MeOH (600 mL).
- Step 5.** Perform solvent-solvent partition with n-hexane and chloroform (5 × 500 mL each) that will yield n-hexane, chloroform, and aqueous MeOH-soluble phases.
- Step 6.** Then evaporate the chloroform-soluble phase and roughly separate the residue (8.56 g) with flash column chromatography on silica gel column by means of n-hexane–acetone gradient system.
- Step 7.** Next, combine the obtained fractions on the bases of TLC into several fractions, for example, fractions I–VII in this case.
- Step 8.** Then, elute one fraction with n-hexane–acetone 85:15 and further separate it by multiple flash chromatography.

Step 9. The resulting two fractions can be finally purified with normal-phase HPLC by means of cyclohexane–isopropanol–water isocratic eluent system to result in compounds 1 and 2.

Step 10. Another fraction (fraction IV) will be subjected to flash column chromatography using n-hexane–acetone gradient system as mobile phase, and further subjected to fractions

Step 11. Further, purification of resulting fractions will be performed by HPLC that will lead to the isolation of compound 3 and so on (Fig. 8.4).

8.4 Extraction and Isolation of Secondary Metabolites from *Agaricus macrosporus*

Major steps involved in extraction and isolation of secondary metabolites from *Agaricus macrosporus* are summarized in Fig. 8.5. (Stadler et al., 2006)

Step 1. Collection of mushrooms.

Step 2. Propagation of mushroom in flask with culture media (for small-scale isolation) or in stirring fermenters (for large-scale isolation).

Step 3. Separate the mushroom mycelium from the culture broth by centrifugation.

Step 4. Filtration and extraction with acetone.

Step 5. Remove acetone in vacuo (ca. 40 °C, 250 mbar) to yield an aqueous residue.

Step 6. Dilute the resulting residue with water and subsequently extract it three times with EtOAc.

Step 7. Dry it over Na₂SO₄ and evaporate the combined organic phase in vacuo to yield oily residue.

Step 8. Divide this crude extract into portions and subject it to preparative HPLC and different compounds can be eluted at their respective retention times.

8.5 Secondary Metabolites of Some Mushrooms and Their Applications

8.5.1 *Ganoderma spp.*

Ganoderma, a medicinal mushroom, is splendidly known for its reservoir of abundant secondary metabolites and their biological significance (Sanodiya et al., 2009; Ríos et al., 2012; Baby et al., 2015). In context to Chinese medical manuscripts, various strains of *G. lucidum* have been medicinally applied in the treatment of chronic ailments such as cancer, diabetes, bronchitis, hypertension, arthritis, hepatopathy, insomnia, and nephritis (Fatmawati et al., 2010). Thus far, more than 200 secondary metabolites have been reported essentially from *G. lucidum* (Chen et al., 2012).

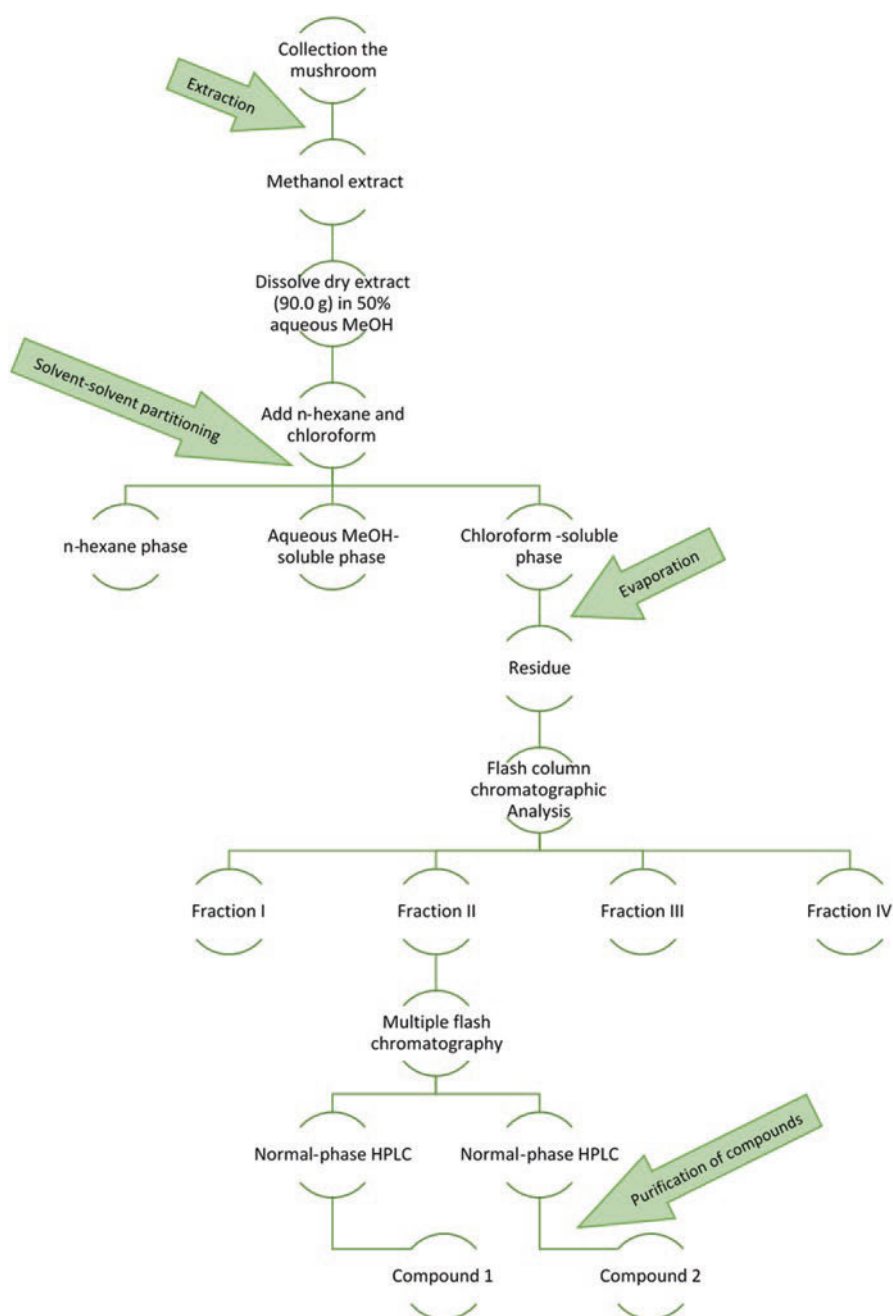


Fig 8.4 Schematic representation of extraction and isolation of metabolites from *Tapinella atro-tomentosa*. (Beni et al., 2018)

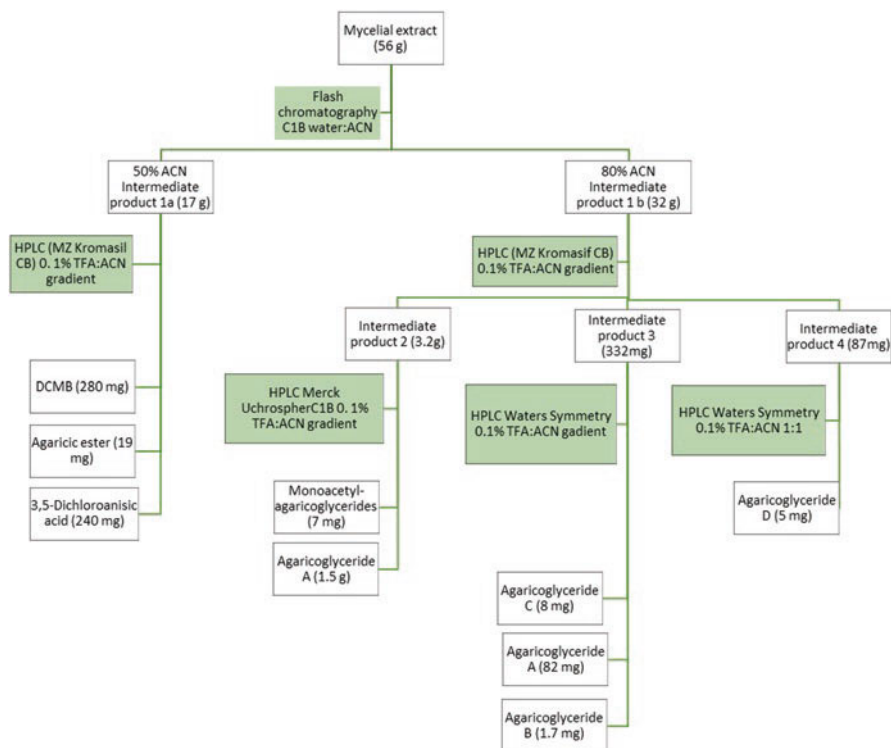


Fig. 8.5 Schematic representation of the isolation of secondary metabolites from mycelial crude extract of *Agaricus macrosporus* (200 litres scale; Q6/2 medium). (Stadler and Hoffmeister, 2015)

According to El Dine et al. (2008a), *Ganoderma colossus* is reported to have anti-HIV-1 protease activities. *Ganoderma applanatum* is very well known for its antitumor activities (Boh et al., 2000), antibacterial properties (Smania et al., 1999), inhibition of Epstein–Barr virus activation (Chairul & Hayashi, 1994), and inhibition of aldose reductase. Smania et al. (2007) reported *Ganoderma austral* to possess antimicrobial activities. Ngai and Ng (2004), described *Ganoderma capense* to be mitogenic and according to El Dine et al. (2008a), *Ganoderma colossus* was testified to embrace anti-HIV-1 protease activities.

Since the last 40 years, various phytochemical studies have led to the identification and isolation of more than 400 bioactive metabolites from several *Ganoderma* species. Some foremost secondary metabolites isolated include alkaloids, meroterpenoids, steroids, C30 lanostanes (aldehydes, ketones, alcohols, glycosides, esters, lactones), C15 sesquiterpenoids, C27 lanostanes (esters, alcohols, lactones), C24 and C25 lanostanes, C27 lanostanes (lucidenic acids), C30 lanostanes (ganoderic acids), farnesyl hydroquinones (meroterpenoids), C30 pentacyclic triterpenes, prenyl hydroquinone, benzopyran-4-one derivatives, benzofurans, and benzenoid derivatives (Baby et al., 2015) as represented in Fig. 8.6. Besides, some common

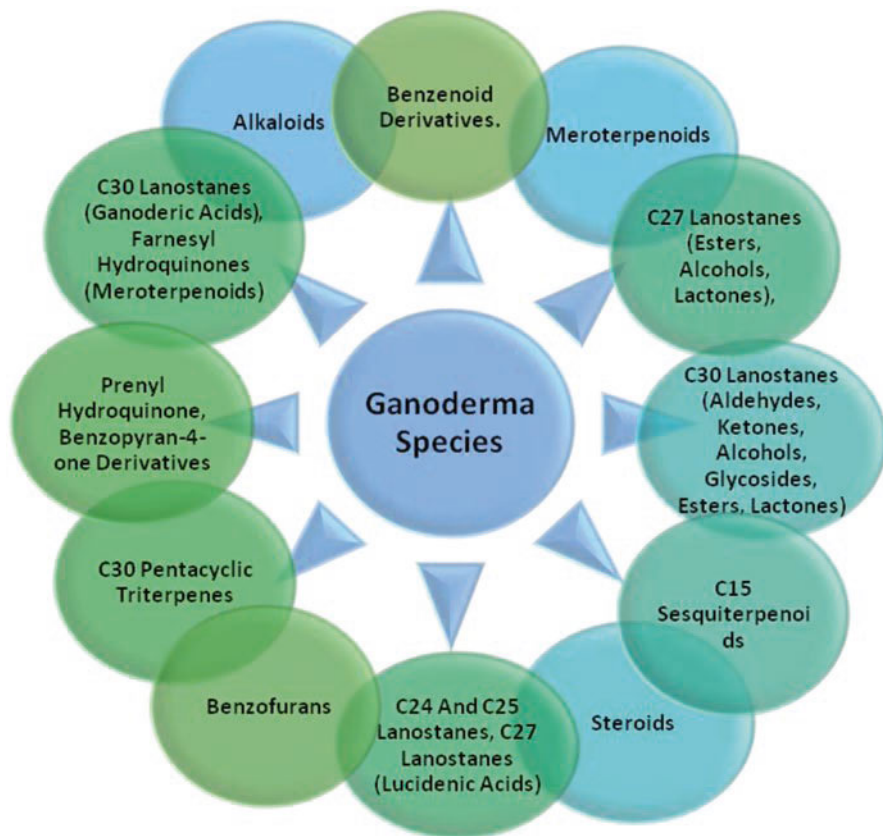
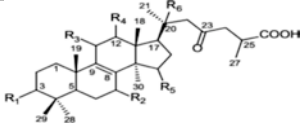
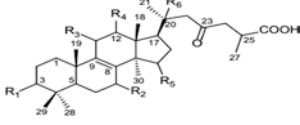
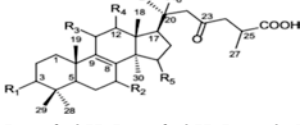
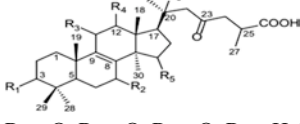
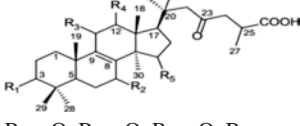
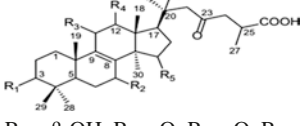
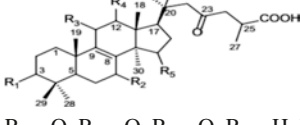


Fig. 8.6 Major groups of secondary metabolites found in *Ganoderma* spp

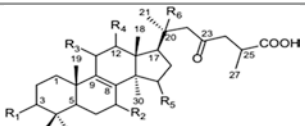
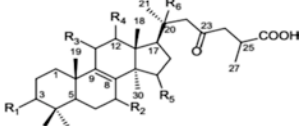
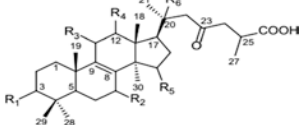
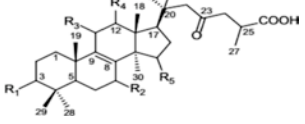
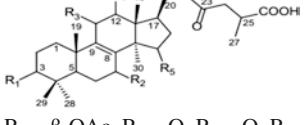
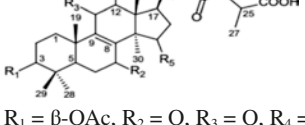
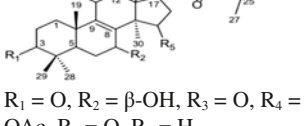
Ganoderma species that are exclusively subjected to phytochemical analysis include *Ganoderma annulare*, *Ganoderma australe*, *Ganoderma amboinense*, *Ganoderma applanatum*, *Ganoderma boninense*, *Ganoderma colossum*, *Ganoderma capense*, *Ganoderma concinna*, *Ganoderma cochlear*, *Ganoderma fornicatum*, *Ganoderma hainanense*, *Ganoderma lipsiense*, *Ganoderma neo-japonicum*, *Ganoderma mastoporum*, *Ganoderma orbiforme*, *Ganoderma pfeifferi*, *Ganoderma resinaceum*, *Ganoderma sinense*, *Ganoderma tsugae*, *Ganoderma tropicum*, and *Ganoderma theaecolum*. Various secondary metabolites isolated from *Ganoderma* species are listed in Table 8.1 for better understanding.

Table 8.1 Secondary metabolites isolated from *Ganoderma* species

| S. no | Compound name | Compound structure | <i>Ganoderma</i> species | References |
|-------|--------------------|---|--------------------------|---------------------|
| 1. | Ganoderic acid A |  <p>$R_1 = O, R_2 = \beta\text{-OH}, R_3 = O, R_4 = H, R_5 = \alpha\text{-OH}, R_6 = H$</p> | <i>G. sinense</i> | Liu et al. (2012b) |
| 2. | Ganoderic acid C1 |  <p>$R_1 = O, R_2 = \beta\text{-OH}, R_3 = O, R_4 = H, R_5 = O, R_6 = H$</p> | <i>G. lucidum</i> | Seo et al. (2009) |
| 3. | Ganoderic acid C2 |  <p>$R_1 = \beta\text{-OH}, R_2 = \beta\text{-OH}, R_3 = O, R_4 = H, R_5 = \alpha\text{-OH}, R_6 = H$</p> | <i>G. lucidum</i> | Min et al. (2000) |
| 4. | Ganoderic acid E |  <p>$R_1 = O, R_2 = O, R_3 = O, R_4 = H, R_5 = O, R_6 = H$</p> | <i>G. sinense</i> | Liu et al. (2012b) |
| 5. | Ganoderic acid F |  <p>$R_1 = O, R_2 = O, R_3 = O, R_4 = \beta\text{-OAc}, R_5 = O, R_6 = H$</p> | <i>G. amboinense</i> | Yang et al. (2012) |
| 6. | Ganoderic acid AM1 |  <p>$R_1 = \beta\text{-OH}, R_2 = O, R_3 = O, R_4 = H, R_5 = O, R_6 = H$</p> | <i>G. lucidum</i> | Cheng et al. (2010) |
| 7. | Ganoderic acid AP3 |  <p>$R_1 = O, R_2 = O, R_3 = O, R_4 = H, R_5 = \alpha\text{-OH}, R_6 = \xi\text{-OH}$</p> | <i>G. applanatum</i> | Wang and Liu (2008) |

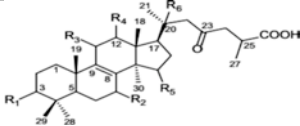
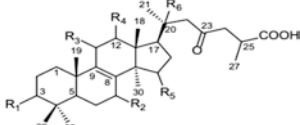
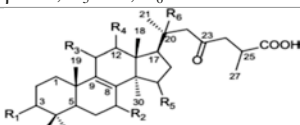
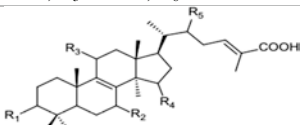
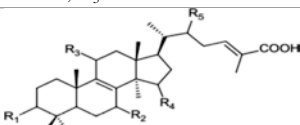
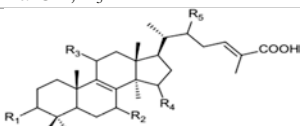
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Table 8.1 (continued)

| S. no | Compound name | Compound structure | <i>Ganoderma</i> species | References |
|-------|-------------------------------|--|--------------------------|-------------------------|
| 8. | Ganoderic acid Df |  <p>$R_1 = O, R_2 = \beta\text{-OH}, R_3 = \beta\text{-OH}, R_4 = H, R_5 = O, R_6 = H$</p> | <i>G. lucidum</i> | Fatmawati et al. (2010) |
| 9. | 12-Hydroxy ganoderic acid C2 |  <p>$R_1 = \beta\text{-OH}, R_2 = \beta\text{-OH}, R_3 = O, R_4 = OH, R_5 = \alpha\text{-OH}, R_6 = H$</p> | <i>G. lucidum</i> | Yang et al. (2007) |
| 10. | 20-Hydroxy ganoderic acid AM1 |  <p>$R_1 = \beta\text{-OH}, R_2 = O, R_3 = O, R_4 = H, R_5 = O, R_6 = \xi\text{-OH}$</p> | <i>G. theaecolum</i> | Liu et al. (2014) |
| 11. | 3-O-Acetyl ganoderic acid B |  <p>$R_1 = \beta\text{-OAc}, R_2 = \beta\text{-OH}, R_3 = O, R_4 = H, R_5 = O, R_6 = H$</p> | <i>G. lucidum</i> | Li et al. (2009) |
| 12. | 3-O-Acetyl ganoderic acid H |  <p>$R_1 = \beta\text{-OAc}, R_2 = O, R_3 = O, R_4 = \beta\text{-OAc}, R_5 = O, R_6 = H$</p> | <i>G. lucidum</i> | Yang et al. (2007) |
| 13. | 3-O-Acetyl ganoderic acid K |  <p>$R_1 = \beta\text{-OAc}, R_2 = O, R_3 = O, R_4 = H, R_5 = \alpha\text{-OH}, R_6 = H$</p> | <i>G. lucidum</i> | Li et al. (2009) |
| 14. | 12-Acetoxy ganoderic acid D |  <p>$R_1 = O, R_2 = \beta\text{-OH}, R_3 = O, R_4 = OAc, R_5 = O, R_6 = H$</p> | <i>G. lucidum</i> | Yang et al. (2007) |

(continued)

Table 8.1 (continued)

| S. no | Compound name | Compound structure | <i>Ganoderma</i> species | References |
|-------|---|--|--------------------------|---------------------|
| 15. | Ganolucidic acid B |  <p>$R_1 = \beta\text{-OH}$, $R_2 = \text{H}$, $R_3 = \text{O}$, $R_4 = \text{H}$, $R_5 = \alpha\text{-OH}$, $R_6 = \text{H}$</p> | <i>G. sinense</i> | Liu et al. (2012b) |
| 16. | 12 b-Hydroxy-3,7,11,15,23-pentaoxo-5a-lanosta-8-en-26-oic acid |  <p>$R_1 = \text{O}$, $R_2 = \text{O}$, $R_3 = \text{O}$, $R_4 = \beta\text{-OH}$, $R_5 = \text{O}$, $R_6 = \text{H}$</p> | <i>G. lucidum</i> | Cheng et al. (2010) |
| 17. | 12,15-Bis(acetyloxy)-3-hydroxy-7,11,23-trioxo-lanost-8-en-26-oic acid |  <p>$R_1 = \text{OH}$, $R_2 = \text{O}$, $R_3 = \text{O}$, $R_4 = \text{OAc}$, $R_5 = \text{OAc}$, $R_6 = \text{H}$</p> | <i>G. lucidum</i> | Yang et al. (2007) |
| 18. | Ganoderic acid V |  <p>$R_1 = \text{O}$, $R_2 = \alpha\text{-OH}$, $R_3 = \text{H}$, $R_4 = \alpha\text{-OAc}$, $R_5 = \text{H}$</p> | <i>G. orbiforme</i> | Isaka et al. (2013) |
| 19. | Ganoderic acid GS-2 |  <p>$R_1 = \text{O}$, $R_2 = \beta\text{-OH}$, $R_3 = \text{O}$, $R_4 = \alpha\text{-OH}$, $R_5 = \text{H}$</p> | <i>G. sinense</i> | Sato et al. (2009) |
| 20. | Ganoderic acid GS-1 |  <p>$R_1 = \text{O}$, $R_2 = \beta\text{-OH}$, $R_3 = \text{O}$, $R_4 = \text{O}$, $R_5 = \text{H}$</p> | <i>G. sinense</i> | Sato et al. (2009) |

8.5.2 *Lentinula edodes* (*Shiitake Mushroom*)

Lentinula edodes has recently garnered international recognition because of its high nutritional value and due to the presence of invaluable bioactive compounds, although it has been cultivated across Asian countries from time immemorial (Royse et al., 2017; Raut, 2019). Studies related to Chinese traditional medicine reveal that Shiitake mushroom had been working as therapeutic in the form of tonics to treat aging-associated sicknesses like pain and fatigue; moreover, they had beneficial effect on heart health and lung diseases (Money, 2016). Current research findings have elucidated the presence of high-molecular-weight compounds like peptides and polysaccharides (Finimundy et al. 2014) as well as small secondary metabolites (Emi Fukushima-Sakuno, 2020) that impart these mushrooms with health benefits avowed in traditional medicine. Nevertheless, some secondary metabolites with significant pharmacological possessions have been listed in Fig. 8.7 and are briefly explained below.

8.5.2.1 Polyacetylenes

L. edodes are found to be rich in with various polyacetylenes including octa-2,3-diene-5,7-diyne-1-ol (Lentinamycin), a compound with potent antimicrobial potential tested against *Trichoderma spp.* (Komemushi et al., 1996; Tokimoto & Komatsu, 1995). Another polyacetylene, 4-Hydroxyundeca-5,6-diene-8,10-diyneic acid (Cortinellin), has been isolated from a culture filtrate of *Cortinellus shiitake*, which is known for its antimicrobial action against mycobacteria, gram-positive as well as gram-negative bacteria, and fungi (MIC 1.6 μ M) (Herrmann, 1962; Dembitsk & Maoka, 2007). Apart from these, several research findings have revealed the abundance of numerous polyacetylenes, as mentioned in Fig. 8.7 that possess potential bioactivity.

8.5.2.2 Sulfurous Compounds

Various sulfurous compounds occurring in *L. edodes* include 1,2,3,5,6-pentathiepane (Lenthionine), 1,2,4,6-tetrathiepane, 1,2,3,4,5,6-hexathiepane (Morita & Kobayashi, 1967), that embrace potent antimicrobial properties. A study recorded that lenthionine had inhibitory effect on liver damage and could be effectively used in treating thrombosis (Kumagai et al., 2013). Chen et al. (2015) also recognized several sulfur compounds in chloroform extracts of *L. edodes* including 1,3,5-Trithiane, 1,2,3,5-Tetrathiane, Dimethyl tetrasulfide Dimethyl disulfide, and others.

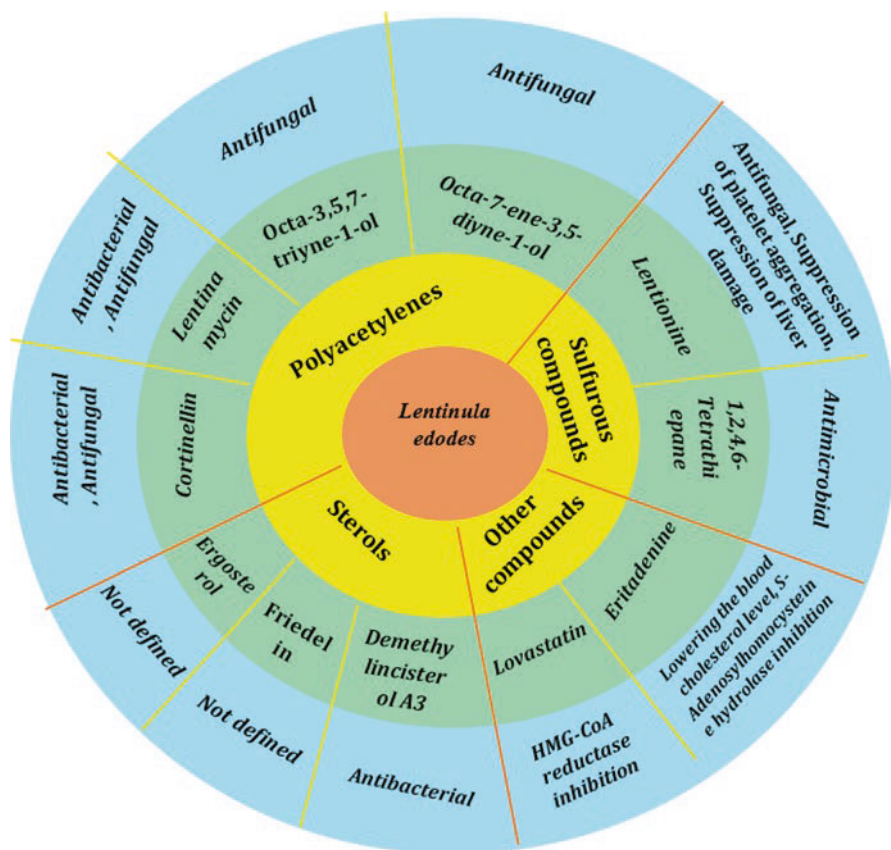


Fig. 8.7 Bioactive secondary metabolites and their related compounds from *Lentinula edodes*

8.5.2.3 Sterols

Ergosterol, a precursor of vitamin D₂, is another group of abundant secondary metabolite occurring in *L. edodes*. However, various other terpenes and sterols have also been reported from *L. edodes*. This is known to possess Friedelin, Demethylincisterol A3, Ergosterol, Methylcholesta 7,22-diene-3 β ,5 α ,6 β -triol, and Methylcholesta 6,22-diene-3 β ,5 α ,8 α -triol are some sterols isolated and identified from acetone extracts of fruiting bodies of *L. edodes* (Chen et al., 2015).

8.5.2.4 Other Compounds

Eritadenine, a nucleic acid derivative isolated from the fruiting body of *L. edodes*, is shown to inhibit angiotensin-converting enzyme and S-adenosyl-L-homocysteine hydrolase (Afrin et al., 2016). Various researches have also revealed the presence of

lovastatin in many mushrooms including *L. edodes* (Kała et al., 2020). Lovastatin is an explicit inhibitor of hydroxymethylglutaryl-CoA reductase enzyme involved in catalysis of cholesterol synthesis in the liver; hence, it lowers the concentration of blood cholesterol (Lin et al., 2013; Kała et al., 2020).

8.5.3 *Flammulina velutipe* (Golden Needle mushroom)

Flammulina velutipes, also called enokitake, is worldwide cultivated edible mushrooms, best known for its flavorsome taste and prodigious nutritional profile (Royse et al., 2017; Raut et al., 2019). Owing to the presence of diverse bioactive metabolites, they exhibit numerous pharmacological activities, like antioxidant, antitumor, immunomodulatory, and cholesterol-lowering properties. Many secondary metabolites of great pharmacologic significance have been isolated and classified from *F. velutipes*, as given in Fig. 8.8 (Tang et al., 2016).

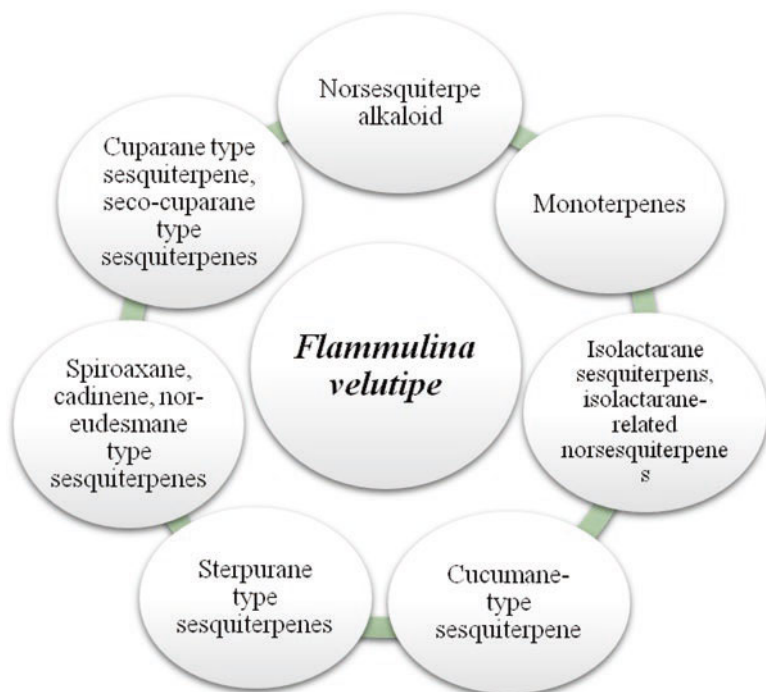


Fig. 8.8 *Flammulina velutipes*: Secondary metabolites of pharmacologic significance

8.5.3.1 Cuparane Type Sesquiterpenes, Seco-Cuparane Type Sesquiterpenes

Flammulin velutipes are known to possess as sesquiterpenes with highly oxygenated cuparane. Ishikawa et al. isolated Enokipodins A, B, C, and D from a culture filtrate of *F. velutipes*. Tabuchi et al. studied the bioactivity of these three compounds in bioassays using liquid medium, where they were found to be antibacterial against *B. subtilis* (IC₅₀ of 13–45 μM) (Tabuchi et al., 2020). Enokipodins B is also known to suppress spore germination of a few plant-pathogenic fungi (Tabuchi et al., 2020). Moreover, Enokipodins B and D also have strong antitumor activity and antiproliferative activity against several transformed cell lines (tsFT210, HL60, HeLa; IC₅₀ of 4.9–18.7 μM) (Tabuchi et al., 2020). Furthermore, enokipodins E, F, G, H, I, and, J were also isolated from rice substrate fermented by *F. velutipes*. Enokipodin F, G, and I have shown antifungal activity against highly pathogenic fungi *Aspergillus fumigatus* (IC₅₀ of 229–235 μM) (Wang et al., 2012). Flamvelutpenoids E & F (with cuparane skeleton) and flammufuranones A & B (secocuparane sesquiterpene) were also isolated from rice fermented *F. velutipes* (Tao et al., 2016).

8.5.3.2 Spiroaxane, Cadinene, Nor-Eudesmane-Type Sesquiterpenes

Flammuspirone A, 7,13,14-Trihydroxy-4-cadinen-15-oic acid methyl ester, Flammuspirone C and, 1,2,6,10-Tetrahydroxy-3,9-epoxy-14-nor-5(15)-eudesmane have been isolated from *F. velutipes* and they are known to show inhibitory activity against HMG-CoA reductase (Tao et al., 2016). Flammuspirone D, E, and H isolated from the same showed DPP4 inhibitory activity, with IC₅₀ 83.7, 70.9, and 79.7 respectively (Tao et al., 2016).

8.5.3.3 Sterpurane-Type Sesquiterpenes

Wang et al. (2012) isolated sterpuric acid and sterpurane sesquiterpenes, sterpurols A and B from *F. velutipes*. However, as per the literature, these three compounds have not shown any antimicrobial activity or antioxidative activity so far. Although sterpuric acid had shown weak antibacterial activity against MRSA and *B. subtilis* (Wang et al., 2012).

8.5.3.4 Cucumane-Type Sesquiterpene

Another cucumane-type sesquiterpene being isolated from rice fermented by *F. velutipes* is Flamulinol A. It is known to exhibit weak antibacterial activity against MRSA and *B. subtilis* (Wang et al., 2012).

8.5.3.5 Isolactarane Sesquiterpens, Isolactarane-Related Norsesquiterpenes

An isolactarane type sesquiterpene, flammulinolide, and a few isolactarane-related norsesquiterpenes, flammulinolides B–G have been reported from rice substrates fermented by *F. velutipes* (Wang et al., 2012). Flammulinolide A is known to be strongly cytotoxic against KB cells (IC₅₀ of 3.9 μ M) and moderately cytotoxic against HepG2 cells (IC₅₀ of 34.7 μ M), as per the literatures. Also, all of these compounds are known to possess antibacterial potentials against MRSA and *B. subtilis*.

8.5.3.6 Norsesquiterpe Alkaloid

A norsesquiterpe alkaloid, (R)-8-hydroxy-4,7,7-trimethyl-7,8-dihydrocyclopenta[e]isoindole-1,3 (2H, 6H) -dione has been reported from *F. velutipes* and it has shown to exhibit cytotoxicity against KB cells at an IC₅₀ of 16.6 μ M (Xu et al., 2013).

8.5.3.7 Monoterpenes

Some growth-promoting low-molecular-weight monoterpenetriols have been successfully isolated from the fruiting body of *F. velutipes* viz., (1R,2R,4R,8S)-(-)-p-Menthane-2,8,9-triol and (1R,2R,4R,8R)-(-)-p-Menthane-2,8,9-triol (Hirai et al., 1998). Some important bioactive secondary metabolites isolated from *F. velutipes* have been listed in Table 8.2.

Table 8.2 Some important bioactive secondary metabolites isolated from *F. velutipes*

| S. N. | Compound | Source | Bioactivity |
|-------|--------------|-------------------------|--|
| 1. | Enokipodin A | Liquid culture filtrate | Antibacterial, antifungal, antimalarial, cytotoxic |
| 2. | Enokipodin B | Liquid culture filtrate | Antibacterial |
| 3. | Enokipodin D | Solid culture substrate | Antifungal cytotoxic ntioxidative |
| 4. | Enokipodin E | Solid culture substrate | Not known |
| 5. | Enokipodin F | Solid culture substrate | Antifungal |
| 6. | Enokipodin I | Solid culture substrate | Antibacterial, antifungal |
| 7. | Enokipodin J | Solid culture substrate | antibacterial |

(continued)

Table 8.2 (continued)

| S. N. | Compound | Source | Bioactivity |
|-------|--|-------------------------|---|
| 8. | 2,5-Cuparadiene-1,4-dione | Solid culture substrate | Cytotoxic antioxidative |
| 9. | Flamvelutpenoid A | Solid culture substrate | Weak antibacterial |
| 10. | Flammuspironone A | Solid culture substrate | HMG-CoA reductase inhibition |
| 11. | 1,2,6,10-Tetrahydroxy-3,9-epoxy-14-nor-5(15)-eudesmane | | HMG-CoA reductase inhibition DPP-4 Inhibition |
| 12. | 7,13,14-Trihydroxy-4-cadinene-15-oic acid methyl ester | | |
| 13. | Flammuspironone D | Solid culture substrate | DPP-4 Inhibition |
| 14. | Sterpuric acid | Solid culture substrate | Cytotoxic Weak, antibacterial |
| 15. | Flammulinol A | Solid culture substrate | Weak antibacterial |
| 16. | Flammulinolide A | | Cytotoxic |
| 17. | Flammulinolide B | | Weak antibacterial |
| 18. | (R)-8-Hydroxy-4,7,7-trimethyl-7,8-dihydrocyclopenta[e]isoindole 1,3(2H,6H)-dione | | Cytotoxic |
| 19. | (1R,2R,4R,8S)-(-)-p-Menthane-2,8,9-triol | Fresh fruiting bodies | Growth promoting and inhibitory activity against stipe of enokitake |
| 20. | (1R,2R,4R,8R)-(-)-p-Menthane-2,8,9-triol | Fresh fruiting bodies | Growth promoting and inhibitory activity against stipe of enokitake |

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Chapter 9

Fungi as Nutraceutical: Present to Future



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9.1 Introduction

“Nutraceutical” has been described as medicinally active or nutritional food supplement since the late 1980s (DeFelice, 1995). The term “nutraceutical” was coined by Stephen De Felice from two different branches of science as “nutrition” and “pharmaceutical” in 1989 (Maddi et al., 2007). “Nutraceuticals” means a combination of “nutrient” and “ceutical,” where “nutrient” refers to functional foods having

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medicinal value and “ceutical” refers to therapy (Borodina et al., 2020). Nutraceuticals generally have some pharmaceutical and antiaging properties and the capability of preventing or reducing symptoms of many long-term diseases. Nutraceuticals can also improve overall wellness, along with boosting immunity (Sharma, 2009). Nutraceutical food products are categorized into different subcategories like polyunsaturated fatty acids, probiotics, prebiotics, dietary fibers, and antioxidants. Industries also define any nontoxic functional food as nutraceuticals that have vital benefits on health issues and also have some potential bioactive compounds to treat/prevent many diseases (Das et al., 2012). These are food products having pharmaceutical properties. Nutraceuticals are used as supplements in developed countries as they are associated with many potential bioactive phytochemicals that also have health benefits (Al-Obaidi et al., 2021). There are various nutraceutical compounds also found in some higher plants, bacteria, and algae. There are many studies on functional foods to discover some bioactive compounds that can be used as a nutraceutical and also to make some beverages having great health benefits.

In nutraceutical development, fungi play an important role. Several species of fungi are nutritionally rich and are a good source of bioactive compounds that not only have nutritional values but also drug-like properties. Among these, mushrooms nowadays have become a hot topic among scientists searching for new therapeutic alternatives (Barros et al., 2008). Mushrooms are rapidly finding a place in our regular diet due to nutritional value, low fat, and high protein content (Barros et al., 2007). Other than having nutritional values, fungi also have antiviral, antibacterial, anticancer, immune-booster, and medicinal importance values (Fig. 9.1).

9.2 Fungi: As Nutraceuticals

Modern science considers and also uncovered some new fungi sources of natural nutraceuticals. There are various fungal species that are being used as medicine since a long time in different parts of the world. Modern scientific researchers have found many fungi species that are very rich sources of nutraceutical (Al-Obaidi, 2016). There are some species that have traditional medicinal values, including Chinese *Cordyceps* (Paterson, 2008) and *Ganoderma lucidum* (Wachtel-Galor et al., 2011). This includes mushrooms, a type of fungi having potential bioactive compounds with some functional properties like antioxidants (Kalaras et al., 2017), anticancer compounds (Ajith & Janardhanan, 2007), antibiotics (Kumar & Kaushik, 2012), immunomodulators (Mallard et al., 2019), hypocholesterolemics (Kim et al., 2009), and hypoglycemics (Giavasis, 2014). This also includes *Cordyceps*, a fungi with medicinal properties that possesses several bioactive molecules and is also used as traditional medicine, with cordycepin (Soltani et al., 2018). Cordycepin is a potential bioactive molecule due to some ability to fight diseases like cancer, cardiovascular disease, and diabetics, which makes it an important mushroom (Ashraf et al., 2020). Fungi like *Ganoderma* were also showing activity to prevent various kidney diseases and also used as traditional medicine. Recently, *Ganoderma* has

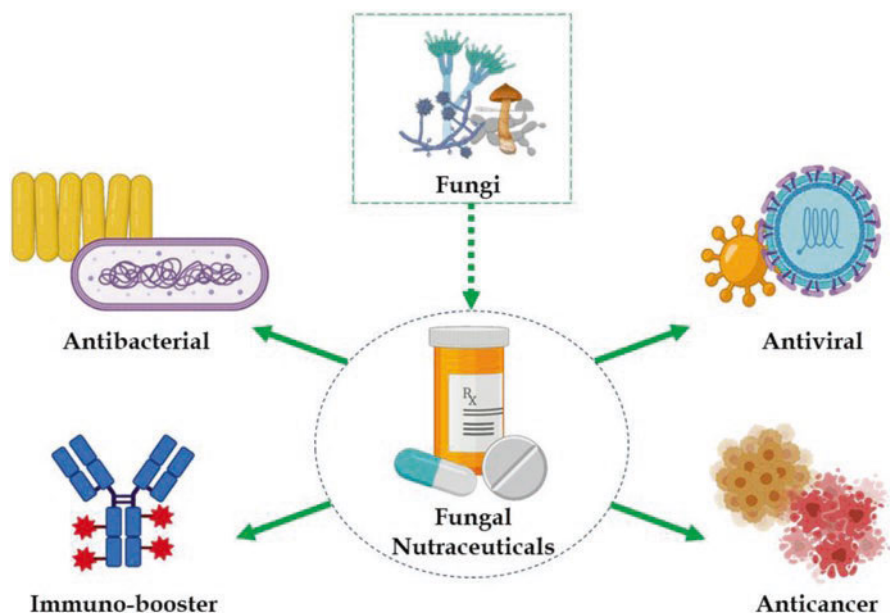


Fig. 9.1 Fungi as nutraceutical and its function

shown some activity having several bioactive compounds for different biological reactions through mass spectrometry studies and spectroscopy studies (Garuba et al., 2020). Research has shown some potential candidates of nutraceuticals that were derived from mushroom and can also be used to ease neurological disorders like mental dysfunction (dementia) and ataxia (Yadav et al., 2020). Beta-glucan is found in various fungal species, which is a common component for nutraceuticals (Kaushik et al., 2019). Cell walls of some plants like oats and barely possess the natural polysaccharide, that is, beta-glucan (Ciecierska et al., 2019). Beta-glucans also show some anticancerous properties that can promote apoptosis in cancer cells (Kobayashi et al., 2005). The antibacterial properties of some fungi are also advantageous to use fungal compounds in different nutraceutical industries. For example, cultured mycelia extracts of *Ophiocordyceps sinensis* has shown some antibacterial potential (Kaushik et al., 2019). Maitake and shiitake are well-known medicinal mushrooms with potential bioactive nutraceutical compounds; however, some other mushrooms like *Cordyceps*, *Agaricus*, and oyster mushrooms are also considered as potential species since the past decade. Some examples of nutraceutical mushrooms are as follows: *Grifola frondosa*, known as maitaka; *Ganoderma lucidum*, known as reishi; *Lentinula edodes*, known as shiitake; *Cordyceps sinensis*, known as *Cordyceps*; *Trametes versicolor* known as turkey tail; and *Hericium erinaceus*, known as lion's mane. The nutraceutical activity of various fungal species and their bioactive compounds are shown in Table 9.1. Structure of some bioactive compounds from fungal origin used as nutraceuticals shown in Fig 9.3. Future research on fungal nutraceuticals could highlight the natural nutraceuticals from traditionally

Table 9.1 List of different fungal species and its bioactive compounds as nutraceuticals

| Sl. no. | Fungal species | Family | Bioactive compounds | Nutraceutical activity | References |
|---------|--|------------------|--|--|--|
| 1 | <i>Clitocybe maxima</i> | Tricholomataceae | Quercetin, gallic acid | Antihyperglycemic and antioxidant, α -glucosidase inhibitory activity | Tsai et al. (2009) |
| 2 | <i>Catathelasma ventricosum</i> | Tricholomataceae | Quercetin, catechin, gallic acid, ergosterol | Antihyperglycemic and antioxidant, anticancer activity | Ohtsuka et al. (1973) and Liu et al. (2012) |
| 3 | <i>Stropharia rugosoannulata</i> | Strophariaceae | Quercetin | Antihyperglycemic and antioxidant | Liu et al. (2012) |
| 4 | <i>Craterellus comucopioides</i> | Cantharellaceae | Quercetin, caffeic acid, gallic acid, ergosterol | Antihyperglycemic and antioxidant | Beluhan and Ramogajec (2011) |
| 5 | <i>Laccaria amethystina</i> | Hydnangiaceae | Quercetin, catechin | Antihyperglycemic and antioxidant, α -amylase inhibitory activity | Liu et al. (2012) |
| 6 | <i>Russula delicata</i> | Russulaceae | Catechin, ascorbic acid, β -carotene | Antioxidant, antimicrobial activity | Yalirak et al. (2009) and Kalač (2009) |
| 7 | <i>Russula griseocarnosa</i> | Russulaceae | Phenolics, ergosterol, β -carotene | Antioxidant | Chen et al. (2010) |
| 8 | <i>Boletus edulis</i> | Boletaceae | Ergothioneine, ergosterol, fungistero | Antioxidant | Ey et al. (2007) |
| 9 | <i>Ganoderma annulare</i> | Ganodermataceae | Applanoxidic acid A | Antioxidant, antifungal activity | Smamia et al. (2003) |
| 10 | <i>Lentinula edodes</i> (Berk.) | Marasmiaceae | Oxalic acid, homogeneous polysaccharide | Antioxidant, antifungal activity, antitumor activity | Bender et al. (2003) and Ya (2017) |
| 11 | <i>Podaxis pistillaris</i> (L.: Pers.) Morse | Agaricaceae | Pipolythiopiperazine-2,5-diones | Antioxidant, antifungal activity | al Fatimi (2001) |
| 12 | <i>Ganoderma lucidum</i> | Ganodermataceae | Triterpenes, heteropolysaccharide | Antiviral agents, antitumor, antioxidative, antidiabetic activity | Lindequist et al. (2005), Zhang et al. (2007) and Xiao et al. (2017) |
| 13 | <i>Ganoderma pfeifferi</i> | Ganodermataceae | Ganodermediol, lucidadiol, applanoxidic acid G | In vitro antiviral activity | Lindequist et al. (2005) |
| 14 | <i>Inonotus hispidus</i> | Hymenochaetaceae | Phenolic compounds, ergosterol | Antiviral agents | Ali et al. (2003) |

| | | | | | |
|----|---|------------------|---|--|---|
| 15 | <i>Collybia maculata</i> | Lyophyllaceae | Purine derivatives | Antiviral agents | Lindequist et al. (2005) |
| 16 | <i>Piptoporus betulinus</i> (Bull.: Fr.) P. Karst. | Fomitopsidaceae | Phenolics, flavonoids | Antiparasitic and antimicrobial agent, in vitro anticancer activity | Lemieszek et al. (2009) |
| 17 | <i>Schizophyllum</i> sp. | Schizophyllaceae | Schizophyllan (β -D-glucan) | Immunomodulation | c Ooi and Liu (2000) |
| 18 | <i>Sparassis</i> sp. | Sparassidaceae | Bioactive β -D-glucan, phenyl derivatives, chalcones, sesquiterpenoids | Enhancement of the hematopoietic response and induction of cytokine production | Jiang et al. (2009) and Kimura (2013) |
| 19 | <i>Polyporus</i> sp. | Polyporaceae | Nucleotides, nucleosides, proteins, amino acids, vitamins, polysaccharides, triterpenoids | Very effective against urethral blockages | Wang et al. (2004) |
| 20 | <i>Pleurotus tuber-regium</i> | Pleurotaceae | b-D-glucan | Antioxidant, hepatoprotective, anti-breast cancer | Zhang et al. (2007) and Rahi and Malik (2016) |
| 21 | <i>Auricularia auricular</i> | Auriculariaceae | Glucan | Hyperglycemia, immunomodulating, antitumor | Rahi and Malik (2016) |
| 22 | <i>Schizophyllum commune</i> | Schizophyllaceae | Glucan, schizophyllan | Antitumor | Rahi and Malik (2016) |
| 23 | <i>Herictium erinaceus</i> | Hericiaceae | Heteroglycan, heteroglycan peptide | Hyperglycemia, immunomodulating, antitumor | Rahi and Malik (2016) |
| 24 | <i>Lentinus edodes</i> | Omphalotaceae | Mannogluca, polysaccharide-protein complex, glucan, lentinan | Immunomodulating, antitumor, antiviral | Rahi and Malik (2016) |
| 25 | <i>Sclerotinia sclerotiorum</i> | Sclerotiniaceae | Glucan, scleroglucon (SSG) | Antitumor | Rahi and Malik (2016) |
| 26 | <i>Polystictus versicolor</i> | Polyporaceae | Heteroglycan, glycopeptide, krestin (PSK) | Immunomodulating, antitumor, antiradiative, hyperglycemia, anti-inflammatory | Rahi and Malik (2016) |

(continued)

Table 9.1 (continued)

| Sl. no. | Fungal species | Family | Bioactive compounds | Nutraceutical activity | References |
|---------|--------------------------------|----------------------|---|--|--|
| 27 | <i>Grifola frondosa</i> | Meripilaceae | Proteoglycan, glucan, galatomanna, heteroglycan, grifolan | Immunomodulating, antitumor, antiviral, hepatoprotective | Rahi and Malik (2016) |
| 28 | <i>Inonotus obliquus</i> | Hymenochaetaeaceae | Glucan | Antitumor, immunomodulating | Rahi and Malik (2016) |
| 29 | <i>Agaricus blazei</i> | Agaricaceae | Glucan, heteroglycan, glucan protein | Antitumor | Rahi and Malik (2016) |
| 30 | <i>Flammulina velutipes</i> | Physalaciaceae | Glucan-protein complex, heteropolysaccharide | Antitumor, anti-inflammatory, immunomodulating activity | Rahi and Malik (2016) and Feng et al. (2016) |
| 31 | <i>Polyporus umbellatus</i> | Polyporaceae | Glucan | Antitumor, immunomodulating | Rahi and Malik (2016) |
| 32 | <i>Clitopilus caespitosus</i> | Entolomataceae | Glucan | Antitumor | Rahi and Malik (2016) |
| 33 | <i>Trametes coriolus</i> | Polyporaceae | Polysaccharide, PSP-a glycopeptides | Antitumor and immunostimulant | Rahi and Malik (2016) |
| 34 | <i>Tricholomopsis rutilans</i> | Tricholomataceae | Polysaccharides | Anticarcinogenic activity, antioxidative, anti-inflammatory | Rahi and Malik (2016) |
| 35 | <i>Volvariella</i> sp. | Pluteaceae | Polysaccharide | Cardiac tonic | Rahi and Malik (2016) |
| 36 | <i>Tremella fuciformis</i> | Tremellaceae | Polysaccharide | Hypocholesteric | Rahi and Malik (2016) |
| 37 | <i>Marasmius androsaceus</i> | Marasmiaceae | Polysaccharide | Analgesic/sedative effect | Rahi and Malik (2016) |
| 38 | <i>Coralyceps</i> sp. | Ophiocordycipitaceae | Glucan, heteroglycan | Antitumor, immunomodulating, antitumor, hyperglycemia | Rahi and Malik (2016) |
| 39 | <i>Coriolus versicolor</i> | Polyporaceae | Polysaccharides PSK and PSP | Anticancerous and antiviral effect on HIV and cytomegalovirus in vitro | Rahi and Malik (2016) |
| 40 | <i>Dictyophora indusiata</i> | Phallaceae | Heteroglycan, mannan, glucan | Antitumor, hyperlipidemia | Rahi and Malik (2016) |
| 41 | <i>Agaricus bisporus</i> | Agaricaceae | Heteropolysaccharide | Immunostimulatory and antitumor activities | Zhang et al. (2014) |
| 42 | <i>Pleurotus ostreatus</i> | Pleurotaceae | Homogeneous polysaccharide | Antitumor activity | Cao et al. (2015) |

| | | | | | |
|----|------------------------------|-----------------|--------------------------|--------------------------------------|-----------------------------|
| 43 | <i>Pleurotus eryngii</i> | Apiaceae | Heteropolysaccharide | Antitumor activity | Ren et al. (2016) |
| 44 | <i>Calocybe indica</i> | Lyophyllaceae | Heteropolysaccharide | Antioxidant and antiaging activities | Govindan et al. (2016) |
| 45 | <i>Dictyophora indusiata</i> | Phallaceae | Vitamin C (55) | Antioxidant | Liu et al. (2019) |
| 46 | <i>Pleurotus ostreatus</i> | Pleurotaceae | Resveratrol | Antioxidant | Koutrotsios et al. (2017) |
| 47 | <i>Aspergillus oryzae</i> | Trichocomaceae | Agmatine (28) | Neurological benefits | Akasaka and Fujiwara (2020) |
| 48 | <i>Mucor circinelloides</i> | Mucoraceae | γ -Linolenic acid | Anti-inflammatory | Chan et al. (2018) |
| 49 | <i>Rhizopus oryzae</i> | Mucoraceae | Lipase (Lipopan F) | Decreases in glycemic response | Huang et al. (2020) |
| 50 | <i>Mortierella alpina</i> | Mortierellaceae | Arachidonic acid | Enhancement of immunity | Mamani et al. (2019) |

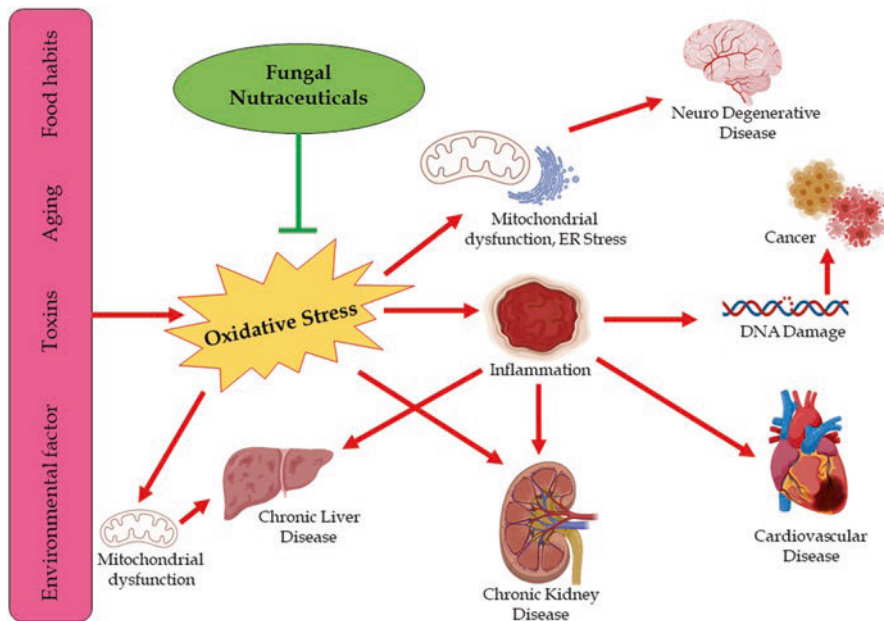


Fig. 9.2 Fungi as a nutraceutical and its function on curing different diseases

used remedies referring to fungi and also can reintroduce the metabolites as potential health supplements in modern pharmaceuticals or nutraceutical industries as potential disease-controlling agents (Fig. 9.2).

9.3 Nutraceutical Production: Using Fungi as Elicitors

Some issues regarding nutraceutical production are the time and cost for its multiple extraction process can be alleviated through advanced techniques of metabolic and some genetic approaches (Liu et al., 2017). Nutraceuticals can also be found in many plants and that co-culturing with their native endophytic fungus can increase yields of nutraceutical production. Therefore, a fungus acts as an elicitor that can influence some metabolic changes to produce the target product and also enhance its production. For example, cell cultures of *Taxus chinensis* were co-cultured with *Aspergillus niger* to increase production of paclitaxel (Wang et al., 2001). There are some other fungal elicitors for compounds like glycyrrhizin (Karwasara et al., 2010) and ajmalicine (Namdeo et al., 2002) that were also studied. Some pathogenic fungi were used as elicitors for nutraceutical production, for example, *Alternaria panax* Whetz, elicitor for *Panax ginseng* (Hao et al., 2020). Recent studies show that *Penicillium* sp. YJM-2013 activates the transcription factor expression, which promotes the increment of ginsenoside production (Wang et al., 2020). Recent studies

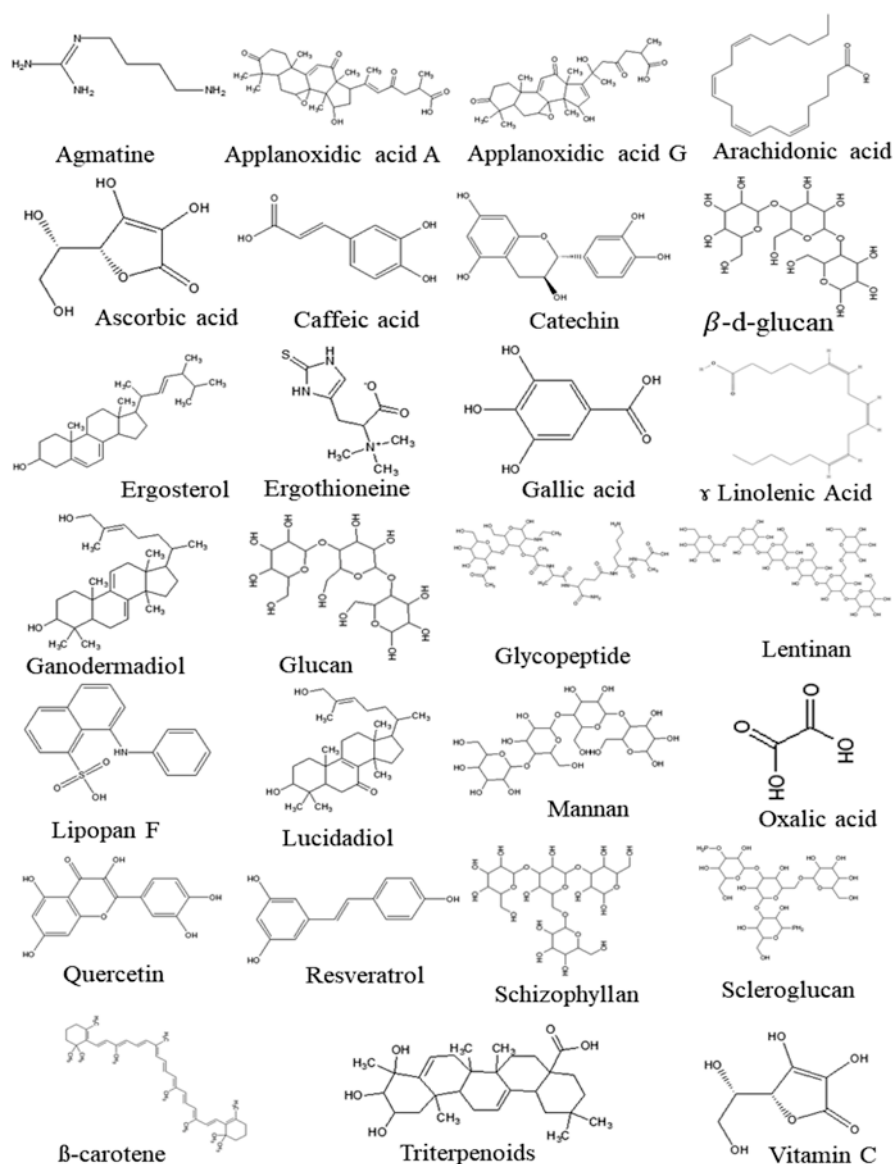


Fig. 9.3 Bioactive compounds from fungal origin

have found that the role of fungi is not only as nutraceuticals but also as a mediator in eliciting nutraceutical production from some other organisms also.

9.4 Nutraceutical Discovery: Prospects of Fungi

So many species of fungi are found on earth that can have advantages to be used as novel nutraceuticals and also as potential food supplements. Few species of fungi have been discovered till now, where it is reported that only 7% were screened for discovering some active bioactive compounds (Hawksworth, 2004). A study found 1526 out of 10,000 fungi in the screening of bioactivity level, and only a few compounds from their extract have certain bioactivity (Hoeksma et al., 2019). Studies were also carried out to discover some novel nutraceuticals from fungi species that were from marine environment and the Antarctic (Nichols et al., 2002). The *Parmeliaceae* family produces some metabolites and proteins that have promising nutraceutical activity found in the Arctic and Antarctic regions (González-Burgos et al., 2019). This will lead to the discovery of some novel nutraceutical compounds and bioactive phytochemicals of fungal species that live in extreme and remote ecological areas. Future prospects of nutraceutical research originating from fungi will highlight the traditional medicinal values of fungi, along with modern approaches in the pharmaceutical/nutraceutical industry. The aim of these types of researches is to introduce some new health supplements using those metabolites (Zhang et al., 2020).

9.5 Entrepreneurship Aspects of Fungal Nutraceutical Industry

The branches of fungal nutraceuticals, the mushrooms, are potential elements and also have global demands. The global demands of various potential mushrooms having pharmaceutical and nutraceutical benefits will lead to establishment of industries and development of various new nutraceuticals. The application of biotechnological and advanced nutrigenomics increases the relevance of nutraceutical industries to produce quality food supplements and increases the range of scope of natural nutraceuticals in potential remedies for human health. Some nutraceutical products like vitamin complexes and inorganic nutrients and also some food ingredients are taken as supplements in many developed countries, for example, there were 47% men and 50% women taking those supplements in the United States (Bas, 2017). The nutraceutical industry is nowadays leading as a functional food supplement producer in developed countries, and hence it opens new wings of entrepreneurship possibilities in natural nutraceutical development. Companies that are involved in this field are also concerned about the intellectual property rights, along with entrepreneurship knowledge (BAS, 2016). Nutraceuticals act as necessary food and also help develop the global market of fungal nutraceuticals. The Western medicinal science is looking for the most significant medicinal mushrooms that have the preventive as well as adjuvant therapeutic properties. The global market of mushrooms is quite huge for the production of nutraceuticals from mushrooms, and

Asia alone has nearly about \$2 billion-valued industries for the production of nutraceuticals from medicinal mushrooms. The publicized market for only *Agaricus* became a potential one with nearly \$450 million in retail in Japan and also for the mushrooms having a rich traditional medicine value in Japan and China craving the attention of scientific researchers to develop them. Mushroom farming is an easy and very profitable agri-business for entrepreneurs with a limited investment and less infrastructure to start. Recent data regarding mushroom cultivation and production market show a value of \$35 billion in 2015, and between 2016 and 2021 the value of the market will be about \$60 billion in 2021. Some countries like the United States, Spain, the Netherlands, Poland, China, Italy, Canada, and the United Kingdom are leading in the field of mushroom producers (Raman et al., 2018). In India, cultivation of different mushrooms is growing and has become an alternative income source for many entrepreneurs. In the recent times, mushroom have become the potential source of nutraceuticals that can help treat chronic diseases.

The global market of fungal nutraceuticals is mainly drives by the increasing demands of mushrooms, and hence mushroom cultivation and production in a large scale may be a suitable entrepreneur opportunity that leads to successful entrepreneurs in nutraceutical industries. Global mushroom demands and production rates estimated a value of US\$ 16.7 billion in 2020, which will reach a value of US\$ 20.4 billion by 2025 (Report: Mushroom Cultivation Market 2020). The innovation and development of nutraceutical products is important to emphasize on the entrepreneurship knowledge, academics, and also intellectual property rights. It is also seen that some researchers do the entrepreneurship practices along with their scientific research that creates major posts in industries and also in journals or scientific community. Nowadays, the nutraceutical industries are leading in healthy food supplement production and bio-pharmaceutical applications that can help the entrepreneurs to grow and establish new opportunities in the field of fungal nutraceuticals (Fig. 9.4). Entrepreneurship can also have an impact on the social and economic growth of the country.

9.6 Conclusion and Future Prospective of Nutraceuticals

Issues associated with nutraceutical development regarding the efficiency have been raised. Nutraceuticals are commonly used as supplements and sold with optimizing dosage. The delivery system and optimizing dosage are very important parameters for nutraceutical efficacy. There are some studies regarding the specificity of long-term effect and also for the dose optimization for positive health impact. It is quite difficult to find the exact bioactive nutrient responsible for acclaimed benefits due to their complex chemical constituents. The rising health challenges due to increasing population rate have made nutraceuticals a potential health-related food supplement obtained from different fungal sources. The higher incidences of different diseases in aging population increase the need of nutraceuticals. A study has shown some that invertebrate models can promote the beneficial areas of using nutraceuticals in

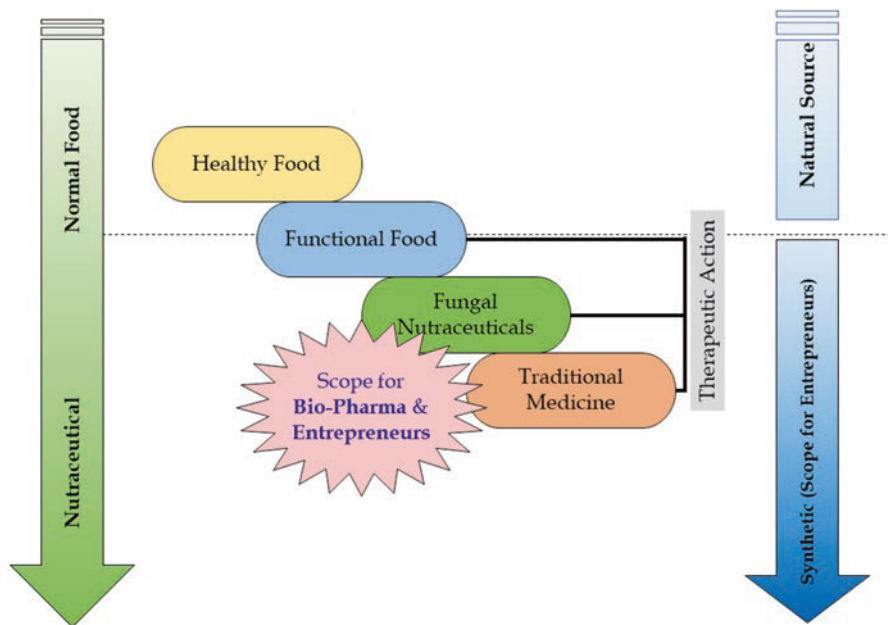


Fig. 9.4 Scopes for entrepreneurs

healthy aging, reducing risk of cancer and inflammation (Dong et al., 2012). The advanced prospective of nutraceutical development can apply in many fields of supplement manufacturing industries. The production time and cost can greatly reduce through the advanced technology of enzyme engineering, coupled with gene/protein discovery. Synthesis of nutraceuticals, the time, and cost can be reduced with the help of advanced DNA-sequencing and gene-editing methods that also open a new avenue for nutraceutical development (Goss et al., 2012). Nutraceutical developmental studies also include research on increasing the bio-availability as its bioavailability profile differs due to diverse chemical nature and structure (Santini et al., 2017). Nutraceutical research for developing a better delivery system for its potential activity also opens a new era of nutraceutical application as a health supplement. Some specialized nano-delivery methods are also used to cure cancer by nutraceuticals (Nair et al., 2010). Modern researches have shown the confidence and trust on nutraceuticals as naturally occurring food/supplements for the treatment of different chronic diseases. Some mushrooms had proved to be very active and effective nutraceutical sources having potential and numerous serious ailments to treat different diseases. Fungal nutraceuticals nowadays attract much attention in the global market for producing potential health supplements and also for their medicinal properties.

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Chapter 10

Potential Application of Edible Mushrooms in Nutrition-Medical Sector and Baking Industries



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10.1 Introduction

Mushroom belongs to the assembly of a macro-fungus with a characteristic fruiting body and morphology. It can be either epigeous or hypogeous. It belongs to Basidiomycetes and is fleshy and edible by nature (Chang, 2008). Mushrooms and their cultivation have a long history in human health and dietary food. It has been valued for its edible and medicinal properties for humankind. Humans used mushrooms as a food source even before they understood the use of other microorganisms that have been discovered later. It is utilized as a food source and medicine source since a very long and ancient time (Gupta et al., 2019). In another sense, the word mushroom is also known as the fruiting body or basidiocarp or sporophores. It has desirable flavors with high nutritive value and is mostly consumed by several countries all over the world. It is a good source of proteins, carbohydrates, vitamins, as well as minerals, dietary fiber, amino acids, etc. It is also reported for its diverse range of secondary metabolites compound, which is the biologically active agent and used as therapeutic purposes (Abdelshafy et al., 2021). Various researches reported these secondary metabolites having medicinal value and also good candidate for utilization as functional foods and used as an indirect cure of diseases.

However, wild mushrooms, edible mushrooms, and medicinal mushrooms are the three main categories in the mushroom industry that are supported by their respective international bodies or forums that have helped to bring each category to the forefront of international attention and show their positive contribution to human welfare. For many years, mushrooms have been used as a food because of their flavor and nutritional properties; of approximately 1.5 million species, there were 69,000 fungal species described and only 14,000 are recognized as mushroom species. Currently, there are still many unknown species (approximately 126,000) yet to be discovered. The growth of mushroom can be distinguished into two phases: the reproductive phase (fruiting body) and the vegetative phase (mycelia) (Ukwuru et al., 2018). There are at least 12,000 species of fungi identified as mushrooms, of which 2000 species are edible (Chang, 1999). In the overall cultivated mushrooms, about 35 species have been commercially cultivated among which 20 are cultivated on an industrial level. According to the order of cultivation patterns, the following are the mushroom species: *Agaricus bisporus*, *Lentinus edodes*, *Pleurotus* species, *Auricula auricula*, *Flammulina velutipes*, *Volvariella volvacea*, etc. (Chang, 2008). Mushrooms and their extracts have long been used in folk medicine and food due to their low calorific value and pleasant taste and are also reported to have beneficial biological activities, thus now found applications in nutraceutical and pharmaceutical products (Giavasis, 2014). The higher market growth rate of mushrooms is due to their increasing acceptability for nutraceutical and medicinal purposes. The nutraceutical sector growth has increased extremely in the past several years due to the functional characterization of a wide variety of bioactive compounds (Valverde et al., 2015). Among others, mushrooms in the Asian continent have been consumed largely in China, India, and Japan due to their traditional food and nutritional value (Boels et al., 2014). Besides, the current scientific research has also confirmed their

beneficial effects mainly as dietary supplements, antioxidants, antitumor and immunomodulator agents, and are also used as stamina and vitality supplement, etc. (Pan et al., 2018). Moreover, Europe is considered to own the largest market for mushrooms and its demand has also increased in the United States in the past years (Zong et al., 2012).

Recently, vast studies and research have been conducted worldwide to increase the interest of mushrooms as a processed food both directly and indirectly (Lin et al., 2008). Mushrooms are added in the products directly as an ingredient or used indirectly as a source of fermentation such as wine making. According to Singh et al. (2010), 55% of the produced mushrooms are used in processed form or canned form because of their short shelf life, and only 45% are consumed in their fresh form. Mushroom is also known as a functional source by containing bioactive compounds, which can be consumed directly in fresh form and processed form, and by isolating particulate compound forms. In functional food industries, mushroom powder is frequently used for making food products or fortifying the food products by using mushroom supplementation. Food supplementation by mushrooms gives a new way to generate the opportunity of employment as well as cure various diseases easily and sustainably. Functional food by mushroom also offers a rich source of nutrient for nutrient-deficient or malnourished population all over the world. This chapter covers the important bioactive compounds of edible mushrooms, their nutraceutical properties, and the application of edible mushrooms in food sectors such as bakery industries.

10.2 Edible Mushrooms and Their Bioactive Compounds

Major bioactive compounds in fungal groups are known as mycochemicals, which can be found in the form of their cell wall components like proteins and polysaccharides or as secondary metabolites like phenolic compounds, terpenes, steroids, etc. (Patel & Goyal, 2012; Gupta et al., 2019). These compounds are naturally found in the mycelium as well as fruiting bodies of mushroom, and their concentration and efficacy depend on the type of mushroom, the substrate for growth, substrate composition, growth conditions, developmental stage, culture or postharvest conditions, storage and cooking procedures, etc. (Enshasy & Hatti-Kaul, 2013; Guillamon et al., 2010). Besides their pharmacological features, mushrooms contain functional metabolites and are used as functional food also beneficial in our diet (Khatun et al., 2012). Mushroom bioactive compounds are categorized into two categories: primary and secondary metabolites. Mushroom secondary metabolites include vitamins, polyphenols, alkaloids, terpenoids, sesquiterpenes, lactones, sterols, nucleotide analogs, glycoproteins, and polysaccharides (β -glucans). Still, new proteins are being discovered from a mushroom that can further be used for the development of new drugs. Some of the identified bioactive compounds are laccase, lectins, lignin-degrading enzymes, proteases, protease inhibitors, ribosome-inactivating proteins, hydrophobins, etc. (Erjavec et al., 2012). Mushrooms contain

all essential amino acids required for the human body; also, they are an excellent resource of vitamins and minerals (Barros et al., 2008).

Numerous scientists investigated the preparation methods and structural characterizations of the molecules obtained from mushrooms bioactive compounds. These molecules mainly belong to polysaccharides, proteins, terpenes, phenolic compounds, unsaturated fatty acids, etc. These mushroom bioactive compounds are responsible for the development of efficient foods resources (Valverde et al., 2015).

10.2.1 Polysaccharides

Polysaccharides are polymeric carbohydrates that are widely distributed in animals, plants, and microorganisms. Various studies reported that mushroom polysaccharides are used as potential pharmacological agents with different bioactivities (Rodrigues Barbosa et al., 2019; Bai et al., 2019; Cheng et al., 2018; Rathore et al., 2019; Ruiz-Herrera & Ortiz-Castellanos, 2019). The polysaccharides found in the genus *Pleurotus* species have specific characteristics related to the chemical structures of glucans, which are varying degrees of bonds and conformations (Singdevsachan et al., 2016). Polysaccharides are formed in numerous types in the mushrooms cells and either can be simpler ones (monosaccharides) or they can be of complex form (oligo- and polysaccharides). Some edible mushroom polysaccharides are reported for the developed as functional food substances such as the schizophyllan from *Schizophyllum commune*, lentinan from *L. edodes*, pleuran from *Pleurotus* species, calocyban from *Calocybe* species, all of which are proved and considered as excellent representatives of D-glucans. Simple polysaccharides such as glucose, fructose, galactose, xylose, mannose, fucose, rhamnose, arabinose, trehalose, and mannitol are the most common polysaccharides found in mushrooms. It has been reported that antitumor, anti-inflammatory, and immunomodulatory activities of mushroom polysaccharides are most important health benefits for human body. Mushroom polysaccharide can activate dendritic cells, monocytes, natural killer cells, neutrophil, cytotoxic macrophages, and different cytokines (Wasser, 2011; Chandrawanshi et al., 2017). Bhandari et al. (2020) reported another group of peptide called bioactive peptides (BAPs) is small fragments (contain 2–20 amino acids) of proteins that provide some health benefits. Most of the BAPs contain >20 AAs (Ryan et al., 2011). Depending on their amino acid composition and confirmation, they perform various biological activities (Bechaux et al., 2019). Because of their size, these BAPs could be easily absorbed by the intestine and directly produce local effects in the digestive tract, or they also enter the circulatory system and show their physiological effects (Erdmann et al., 2008). According to Daliri et al. (2017) and Montesano et al. (2020), the biological function of the bioactive proteins has been credited to their encoded BAPs and can be released without losing the bioactivities. Peptide shows improved bioactivity as compared to their parent proteins (Udenigwe & Aluko, 2012). Because mushroom protein contains it, it also can be an ideal source for the discovery of BAPs. β -Glucans are biologically active

molecules or glucose polymers found in yeast, grain, alga, and fungus. β -Glucans are polysaccharides composed of multiple linear and branched D-glucose units linked by β -1-3 and 1-6 bonds. The bioactivity of polysaccharides is significantly influenced by physicochemical modifications such as the degree of branching and addition of substituent groups (sulfates, selenates) (Li et al., 2015). *Pleurotus* species mushrooms are an excellent source of crude fiber (10.2%) and particularly for β -glucans content (25.9%), which can be widely associated with prebiotic and anti-cholesterol, anticancer, and immunomodulator properties (Aida et al., 2009; Zhu et al., 2015). *P. citrinopileatus* and *P. ostreatus* have been reported for their highest crude fiber and β -glucan content, which are 20.7 and 50%, respectively. It can be used as drugs and is also known as a biological response modifier. β -Glucans, being pleuran, are more studied and recognized in *Pleurotus* genus mushrooms having bioactivity in humans and natural immunostimulant (Imunoglukan P4H[®]) (Bergendiova et al., 2011; Jesenak et al., 2013). Sulfated polysaccharide shows increased solubility, antiviral, anticancer, anticoagulant, and antioxidant activities, etc. It has anti-infective, antitumor, antiviral, and immunomodulatory activities as also wound-healing properties (Pretus et al., 1991; Williams et al., 1996; Majtan Jesenak, 2018).

10.2.2 Proteins and Peptides

Edible mushrooms are also an excellent source of bioactive proteins and peptides, which possess different biological activities. Numerous proteins and peptides are identified and characterized as well as their mechanisms of action are well established by researchers. But there are many other unidentified proteins, so fungi represent a relatively unexplored source of novel proteins. Similarly, it is also an endless source of novel peptides. The studies revealed that dry mushrooms contain 228–249 g (g/kg of mushroom) protein, which is much higher than other protein sources (Petrovska, 2020; Ma et al., 2018).

10.2.3 Terpenes

Chemically terpenes are the cluster of volatile unsaturated hydrocarbons and are classified as monoterpenoids, diterpenoids, triterpenoids, sesquiterpenoids, etc. Researchers have found and isolated various types of sesquiterpenoids from mushrooms, including drimane, bisabolane, aristolane, spiro, cuparene, lactarane, nardosinone, fomannosane, sterpurane, etc. Most of the triterpenoid compounds are lanostane type, which is isolated from mushrooms. Moreover, the mushroom terpenes have been proved to have numerous health benefits, such as antioxidant, antiviral, and anticancer activities (Klaus et al., 2017).

10.2.4 Phenols

Phenols, also called phenolic compounds, have one aromatic ring (C₆) at least and one or more (OH) groups. It is found in the form of small simple molecules to complicated polymers (Michalak, 2006). Different kinds of phenolic compounds are identified in edible mushroom extract such as phenolic acids, flavonoids, fannins, tocopherols, etc. (Abdelshafy et al., 2021). Studies revealed that the antioxidant activity is the most significant bioactivity shown by phenolic compounds. It is well known as free radical inhibitors, metal inactivators, peroxide decomposers, and oxygen scavengers (Heleno et al., 2012). It is well known that phenolic compounds are active biomolecular and protect against several degenerative disorders such as brain dysfunction, cardiovascular diseases, and aging (Finimundy et al., 2013).

10.3 Application of Bioactive Compounds from the Edible Mushroom in the Nutraceutical Sector

Bioactive compounds provide medicinal or health benefits like prevention and treatment of human disease (Rathee et al., 2012). These bioactive compounds can act as immunomodulatory, anticarcinogenic, antiviral, antioxidant, and anti-inflammatory agents (Badalyan, 2014). Nutraceuticals are defined as biologically active molecules that have both nutritional as well as pharmacological potential. Nutraceuticals are present in various foods and are natural bioactive or chemical compounds. They provide nutrient source for the human body by also having health-promoting, disease-curing, or disease-prevention properties. Nutraceuticals included lipids, vitamins, carbohydrates, proteins, minerals, secondary metabolites, etc. (Sachdeva et al., 2020). Nutraceutical industries are rising rapidly, but are still in their initial phase. It gives new opportunities for employment as well as is good for human welfare. Approximately \$379.061 billion in the year 2017 has accounted for the global nutraceuticals market and will be expected to grow up to \$734.601 billion by 2026 (Sharma et al., 2013). Currently, some of the nutraceuticals are available in the market, and some are popular worldwide, such as RoxR, an energy drink (Rox America), ProteinexR, a protein supplement (Pfizer Ltd.), Snapple-a-day™, a meal replacement beverage (Snapple Beverage Group), etc.

For effective and successful nutraceuticals industry development, research is needed and continues to rise. In recent times, there are more than hundreds of mushroom species for which a variety of bioactive properties are detected. Mushrooms are consumed as concentrate powder as well as in extracted (in hot water) form. Currently, in the food markets and pharmaceutical industries, a large number of mushroom products and its isolate compounds exist and are represented as “nutraceuticals” or “functional food or additives.” The extracts can be used as a drink, freeze-dried or spray-dried to form granular powders, which allow for easier handling, packaging, transportation, and consumption (Mizuno et al., 1995). These

liquid concentrates and dried-powdered extracts or fruiting body powder can be placed in a capsule that is considered and utilized as dietary supplements or nutraceuticals. Researchers also reported that isolated bioactive substances can be used for fortification of common food products (Shamtsyan, 2016). Bioactive compounds isolated from edible mushrooms and their biological activities are shown in Table 10.1.

10.3.1 Antioxidants

Antioxidants are substances or compounds that play a very vital role in protecting human body from the free radicals generated by oxidative damages. Therefore, it becomes very essential for food and food supplements in the diet with compounds that are rich in antioxidants. It has been well reported that mushrooms extract alone and isolated polysaccharides have antioxidant activity and can decrease the production of oxygen-free radicals (Wasser & Weis, 1999). A variety of edible mushrooms are reported to possess antioxidant activity. It is generally accepted that extracts of fungi contain many components, each of which has its specific biological effects (Kozarski et al., 2011; Chandrawanshi et al., 2018). Flavonoids, glycosides, phenolics, tocopherols, polysaccharides, carotenoids, ergothioneine, ascorbic acid, etc., are some of the antioxidant compounds found in fruiting bodies and mycelium of mushrooms (Chen et al., 2012). Water-soluble glucans, and other glucans from *A. bisporus* as well as glucans from *Auricularia auricula*, *Flammulina velutipes*, *Ganoderma lucidum*, and *L. edodes*, had notable antioxidant capacity and free radical scavenging potential, thus reported as novel natural antioxidant source (Boh & Berivic, 2007).

10.3.2 Cholesterol-Lowering Effect

Cholesterol has been divided into two major categories: low-density lipoprotein and very-low-density lipoprotein, which is formally called “bad” cholesterol, and high-density lipoprotein, called “good” cholesterol. Hypercholesterolemia is the condition that increases the risk of cardiovascular diseases that cause risk for human health.

High levels of circulating cholesterol in the blood cause deposits inside blood vessels, and this deposit generates another disease called arteriosclerosis. A major rate-limiting step in the biosynthetic pathway for cholesterol formation is at the level of the microsomal enzyme 3-hydroxy-3-methylglutaryl-coenzymeA reductase (HMG-CoA reductase) that catalyzes the reductions of HMG-CoA into mevalonate (Bobek et al., 1991). It was reported that some mushrooms produce compounds that inhibit cholesterol syntheses by inhibiting specific enzyme in biosynthesis pathway. Some species of the genus *Pleurotus* are reported for the production of mevinolin

Table 10.1 Identified mushroom bioactive compounds and their bioactivity

| Bioactive compounds | Name of mushroom | Compound name | Part | Identified bioactivity | References |
|------------------------------|------------------------------|---|---------------|--|--|
| Proteins and peptides | <i>Pleurotus ostreatus</i> | Lectin | Fruiting body | Immunomodulatory activity | He et al. (2017) |
| | <i>Pleurotus ostreatus</i> | Lectin | Mycelium | Immunomodulatory | Devi et al. (2013) |
| | Mushrooms | Fungal immunomodulatory proteins (FIPs) | Fruiting body | Useful in tumor immunotherapy, suppress tumor invasion and metastasis | Li et al. (2019) |
| | <i>Hypsizygus marmoratus</i> | Ribosome-inactivating proteins (RIPs) | Fruiting body | Used in antiviral and cancer therapies, antifungal | Wong et al. (2008) and Lam and Ng (2001) |
| | <i>Agaricus bisporus</i> | Tyrosinases | Fruiting body | Involved in the synthesis of melanin, used as biomarker for vitiligo | Zehtab et al. (2000) |
| | <i>Amanita hemibapha</i> | Ribonucleases (RNases) | Fruiting body | Antiproliferative activity/antitumor. | Sekete et al. (2012) |
| | <i>Agaricus bisporus</i> | Lectin | Fruiting body | Immunomodulatory activity, reducing the innate and adaptive responses | Ditamo et al. (2016) |
| | <i>Pleurotus eryngii</i> | Laccases | Fruiting body | Antiviral activity, inhibit the activity of HIV-1 reverse transcriptase | Wang and Ng (2006) |
| | <i>Pleurotus cornucopiae</i> | Laccases | Fruiting body | Antiviral and antitumor activities, inhibited the activity of HIV-1 reverse transcriptase, and proliferation of the hepatoma cells HepG2 and the breast cancer cells | Wu et al. (2014) |

| Polysaccharides | | | | | |
|-------------------------------|---|--|---|----------------------------|--|
| <i>Pleurotus eryngii</i> | Heteropolysaccharide | Fruiting body | Hypolipidemic and hypoglycemic activities, increase the level of cholesterol and liver glycogen of high-density lipoprotein | Chen et al. (2016) | |
| <i>Pleurotus tuber-regium</i> | β -d-glucan | Mycelium | Hepatoprotective, antibrust cancer | Zhang et al. (2007) | |
| <i>Auricularia auricular</i> | Glucan | Fruiting body | Hyperglycemia, immunomodulating, antitumor, anti-inflammatory, antiradiative | Zhang et al. (2007) | |
| <i>Lentinus edodes</i> | Mannoglucan | Culture broth, fruiting body | Polysaccharide-protein complex, glucan, lentinan immunomodulating, antitumor, antiviral | Zhang et al. (2007) | |
| <i>Pleurotus ostreatus</i> | Glycoprotein | Fruiting body | Antitumor, hyperglycemia, antioxidant | Zhang et al. (2007) | |
| <i>Lentinus edodes</i> | Lentinan | Fruiting body | Antigastric cancer | Oba et al. (2009) | |
| <i>Agaricus bisporus</i> | α -Glucans | Juice from fruiting body | Anti-hypercholesterolemia | Volman et al. (2010) | |
| <i>Agaricus bisporus</i> | Heteropolysaccharide | Mushroom wastewater | Hepatoprotective, against CCl4-induced hepatic injury | Huang et al. (2016) | |
| <i>Agaricus bisporus</i> | Homogeneous polysaccharide | Powder from fruiting body | Antitumor activity, inducing apoptosis by the mitochondrial death pathway | Pires et al. (2017) | |
| <i>Pleurotus ostreatus</i> | Heteropolysaccharide | Powder from fruiting body | Antidiabetic activity, activate GSK3 phosphorylation and GLUT4 translocation | Zhang et al. (2016a, b) | |
| <i>Lentinula edodes</i> | Heteropolysaccharide | Dried fruiting body | Antitumor activity | Wang et al. (2015) | |
| <i>Pleurotus ostreatus</i> | Heteropolysaccharide | Ultrasonic-assisted extracted of fruiting body | Alleviate cognitive impairment activity | Zhang et al. (2016a, b) | |
| <i>Pleurotus eryngii</i> | Heteropolysaccharide | Hot water extracted of fruiting body | Antitumor activity and stimulate the production of intracellular ROS of HepG-2 cells | Ren et al. (2016) | |
| <i>Pleurotus sajor-caju</i> | Lovastatin polysaccharide homopolysaccharides | Fruiting body | Lower cholesterol, prevents cardiovascular disorders | Lakshampal and Rana (2005) | |
| <i>Cordyceps sinensis</i> | Heteropolysaccharides, cordycepin | Fruiting body | Hypoglycemic activity, anti depressant activity, cellular health properties | Lakshampal and Rana (2005) | |

(continued)

Table 10.1 (continued)

| Bioactive compounds | Name of mushroom | Compound name | Part | Identified bioactivity | References |
|-----------------------|--------------------------------|---|---------------|--|-----------------------|
| Phenolic acids | <i> Amanita crocea </i> | Cinnamic, p-hydroxybenzoic, p-coumaric | Fruiting body | Antioxidant, enzyme inhibition, and antimutagenic properties | Alkan et al. (2020) |
| | <i> Agaricus brasiliensis </i> | Gallic, ferulic, benzoic, p-hydroxybenzoic, gentisic, trans-cinnamic, p-coumaric, fumaric acids, catechol | Fruiting body | Antioxidant and antimicrobial activities | Bach et al. (2019) |
| | <i> Agaricus subrufescens </i> | p-Coumaric | Fruiting body | Antioxidant activity | Ferrari et al. (2021) |
| | <i> Agrocybe aegerita </i> | Sinapic, chlorogenic, Gallic, protocatechuic, ferulic acids | Fruiting body | Antiangiogenic | Lin et al. (2017) |
| | <i> Calocybe indica </i> | Gallic, vanillin, protocatechuic acid, naringin, naringenin, homogentisic acid, hesperetin | Fruiting body | Antioxidant and antityrosinase activities | Alam et al. (2019) |
| | <i> Pleurotus pulmonarius </i> | Caffeic, gallic, and ferulic acids | Mycelium | Antioxidant activity | Contato et al. (2020) |

| | | | | | |
|--------------------|--|---|--|--|--------------------------------|
| Flavonoids | <i>Laccaria amethystea</i> and <i>Laccaria ventricosum</i> | Quercetin and catechin | Powder from fruiting body | Antihyperglycemic and antioxidant activity | Liu et al. (2012) |
| | <i>Lactarius indigo</i> | Myricetin, procyanidin, quercetin-hexoside, isoquercetin-hexoside, and kaempferol | Fruiting body | Antioxidants | Yahia et al. (2017) |
| | <i>Lentinus edodes</i> | Flavonoids rutin and quercetin | Fruiting body | Antioxidant activity | Xiaokang et al. (2020) |
| | <i>Pleurotus citrinopileatus</i> | Rutin and quercetin | Ultrasound and ethanolic extracts from fruiting body | Antioxidants | Gogoi et al. (2019) |
| | <i>Pleurotus tuber-regium</i> | – | Powder from fruiting body | Antioxidant activity | Akindahunsi and Oyetayo (2006) |
| Tannins | <i>Astraeus hygrometricus</i> | Catechin | Mushroom flours | Antioxidants | Pavithra et al. (2016) |
| | <i>Pleurotus ostreatus</i> | α -Tocopherol | Ethanollic extracts of from fruiting body | Antioxidants | Jayakumar et al. (2009) |
| Tocopherols | <i>Morchella esculenta</i> | Catechol, p-coumaric acid, rutin, hyperoside, quercetin, tocopherol, and ellagic acid | Dried fruiting bodies | Antioxidants | Wagay et al. (2019) |
| | <i>Russula delica</i> , <i>Boletus badius</i> , and <i>Agaricus bisporus</i> | α -Tocopherol | Methanolic extracts of from fruiting body | Antioxidants | Elmastas et al. (2007) |
| | <i>Pleurotus ostreatus</i> and <i>Pleurotus eryngii</i> | Tocopherol | Mushroom mycelia | Sources of lipophilic antioxidants | Bouzgarrou et al. (2018) |
| | | | | | |

(lovastatin), which is known as the inhibitor of the HMG-CoA reductase and used for the treatment of hypocholesteremia.

Another compound, eritadenine, a compound extracted from *L. edodes* mushroom, is able to lower blood serum cholesterol level. It was reported that the addition of mushrooms' dried fruit bodies or submerge mycelium in a high cholesterol diet can effectively reduce cholesterol accumulation in the blood serum and liver. It was observed that mushroom diet in the experimental rats redistributed cholesterol in favor of HDL, reduced VLDL and LDL production, and reduced HMG-CoA reductase activity and cholesterol absorption in the liver and blood serum (Popov et al., 2008).

10.3.3 Antiviral Activity

According to Pradeep et al. (2019), antiviral is an agent that kills a virus or it can suppress the replication and therefore inhibits its propagation. Antiviral compounds interfere with viral replication and its multiplication as a host. These are isolated from either natural sources like plants, bacteria, and fungi or can be chemical synthesis (Kulkarni & Sanghai, 2014). Many researchers reported the antiviral effects of *Pleurotus* mushroom from the whole extracts as well as isolated specific compounds. It is reported (Fan et al., 2006) that the extract from both mycelia and fruiting bodies showed various bioactivities such as antiviral, antibacterial, hematological, antitumor, hepatoprotective and hypotensive effects, etc. The isolated active compounds (small molecules) from mushroom played a direct role such as acting as inhibitors of viral enzymes and nucleic acids synthesis, and inhibit the absorption and uptake of viruses into mammalian cells. Polysaccharides and other complex molecules played indirect antiviral effects (Brandt & Piraino, 2000). These bioactive compounds acted either directly (exhibited especially by smaller molecules) or indirectly (the immune-stimulating polysaccharides or other complex molecules). β -Glucans had been potentially used by many viruses such as; infectious hematopoietic necrosis virus, spring viremia of carp virus, dengue virus, herpes simplex virus type 1 (HSV-1), human immunodeficiency virus, influenza virus, tobacco mosaic virus, etc. But only a few studies reported its human and animal clinical trials, and has been conducted for the treatment or prevention of viral infections (Adotey et al., 2011). β -Glucans can work directly through the inhibition and/or disruption of virus particles or indirectly by enhancing the immune system. β -Glucan's action is mediated by its several receptors such as the dectin-1 receptor, toll-like receptors (TLR 2, 4, and 6), complement receptor 3 (CR3), scavenger receptor, and lactosylceramide, among which dectin-1 receptor is the most important, which is highly expressed in many immunocompetent cells like neutrophils, eosinophils, macrophages, dendritic cells, monocytes, several T lymphocytes, etc. (Legentil et al., 2015).

10.3.4 *Antiobesity*

Obesity is a group of disorders in which body mass index is more than 30 kg/m². This causes enhancement of body fat deposited in the adipose tissue, which can cause harmful health effects. There are various complications associated with obesity such as diabetes, cardiovascular diseases (CVDs), pulmonary diseases, cancer, osteoarthritis, etc. (Ganesan & Xu, 2018). Obesity is caused by several factors or reasons like high food intake, sedentary lifestyles, lack of physical activity, and a genetic predisposition. Excess stored fat and high lipid content in the plasma in obese persons was observed. Adipose tissue is an organ that helps to maintain energy balance in the body, and in the case of obesity high fat accumulated within causes unusual progress of white adipose tissue (Kim, 2011). A balanced diet, exercise, and surgical intervention are some primary managements of obesity. Several therapeutic drugs are also available for the treatment of obesity, but these are associated with other health complications (Kennett & Clifton, 2010). Obesity also causes and generates other health complications. Obesity is a metabolic disorder, so there is a need for treatment that is associated with daily routine or that can say in the form of food. Edible mushrooms have been well documented for their numerous bioactive compounds and can be used as a functional food. It can directly or indirectly be used for the cure of obesity. Its hypocholesterolemic effects are connected to CVD (diminishes CVD)-related lipid metabolism, anti-inflammatory properties, and the prevention of oxidative stress with platelet agglutination (Sun et al., 2007). It is observed that the cholesterol-lowering effect might be due to a decrease in VLDL and decrease in the catalytic functions of HMG-CoA reductase and amplification of the rate of cholesterol catabolism (Bobek et al., 1991). It is reported that the polysaccharide from *Tremella fuciformis* prevented the variation of 3T3-L1 adipocytes by decreasing the expression of mRNA and suggesting that the possible role of the polysaccharide act as an antiobesity prebiotic. Cheskin et al. (2008) and Poddar et al. (2013) reported long-term and short-term clinical studies with obese or diabetic participants. They evaluated the impact of substituting 20% of high-energy beef with 20% to low-energy white button mushrooms in their diet, and the results showed that the mushroom regime consumers had lesser BMI, decreased belly circumference, and increased satiety without diminishing palatability. This study concluded that the consumption of *A. bisporus* has shown potential as an antidiabetic, antiobesity, etc.

10.4 Mushroom-Based Bakery Products and Their Benefits as a Functional Food

Fortification or enrichment is defined as the addition or enhancement of desired nutrients elements in developing foodstuff products. Fortification is an alternative and effective approach to prevent deficiency of one or more nutrients in certain

specific groups and population (Codex Alimentarius, 1987). A fortification strategy is used for making functional foods in which food is fortified with bioactive metabolites. The demand and research related to “functional foods” have recently attracted increasing attention worldwide. By containing various metabolites, mushrooms are considered as a popular functional food. According to the Institute of Medicine’s Food and Nutrition Board, “functional foods” are defined as foods or dietary components that provide health benefits beyond basic nutrition (Ferreira et al., 2017). Bioactive compound type and quantity in mushrooms depend upon the mushroom species, environment, and substrate in which they grow. Similarly, its functional activity depends upon the type and concentration of bioactive compounds (Ferreira et al., 2017). Till now there are hundreds of mushroom species that have been identified and extensively studied for their biological properties and potential for the prevention and treatment of several human diseases. Recently, there is a gradual increase in interest in the field of functional food from mushrooms by both direct as well as indirect use (Lin et al., 2008). It has been reported that 55% of the total mushroom produced is used in a processed form and 45% are consumed in fresh form (Moon & Lo, 2013; Singh et al., 2010). At present, advanced lifestyle demands have grown interest in the formulation and production of healthy and tasty food items that are rich in nutrients. Edible mushrooms can be used an alternative and healthy substitute for animal protein because they contain high protein as well as all the essential amino acids and vitamin D (Abigail et al., 2020). The approximate compositions of some edible mushrooms are shown in Table 10.2.

They have reported high content of sulfur amino acids and glutamic acids that provides meaty test or umami taste to mushrooms (Kim, 2011). Mushrooms are well known for their unique taste and texture flavor that differ from other food sources (Alemu, 2014). Their global and economic value is now increasing rapidly due to food as well as their medicinal and nutritional values. There are several

Table 10.2 Protein, carbohydrate, fat, ash, and moister content of some edible mushrooms

| Mushroom | Moister content | Carbohydrate content | Protein | Fat | Ash | Reference |
|------------------------------|-----------------|----------------------|--------------|-------------|--------------|-------------------------|
| <i>Agaricus bisporus</i> | 91.27 ± 0.45 | 73.80 ± 6.53 | 14.09 ± 0.23 | 2.18 ± 0.34 | 9.74 ± 1.95 | Reis et al. (2012) |
| <i>Pleurotus ostreatus</i> | 89.17 ± 2.12 | 85.87 ± 19.21 | 7.02 ± 0.55 | 1.39 ± 0.18 | 5.72 ± 0.74 | Reis et al. (2012) |
| <i>Pleurotus eryngii</i> | 82.59 ± 0.36 | 78.60 ± 0.75 | 2.09 ± 0.01 | 4.36 ± 0.14 | 14.95 ± 0.91 | Reis et al. (2014) |
| <i>Pleurotus sajor-caju</i> | 88.00 ± 0.00 | 69.05 ± 0.00 | 12.37 ± 0.00 | 1.80 ± 0.00 | 5.78 ± 0.00 | Mahamud et al. (2012) |
| <i>Pleurotus pulmonarius</i> | 89.17 ± 0.00 | 76.11 ± 0.00 | 30.30 ± 0.00 | 1.10 ± 0.00 | 13.21 ± 0.00 | Rana et al. (2015) |
| <i>Lentinus edodes</i> | 90 ± 91.80 | 67.5 ± 78.0 | 13.4 ± 17.50 | 4.9 ± 8.00 | 3.7 ± 7.00 | Crisan and Sands (1998) |

mushroom species that have been identified and cultivated worldwide. As a form of food source, mushroom is used in various forms and has generated various industrial opportunities (Waktola & Temesgen, 2018). Bakery products are one of the widely consumed food products and have a big market and income-generating industries. Bakery products are well known for their varied tastes, long shelf-life, and low cost, all these properties making it very popular in the food industry. Bakery products such as biscuits, bread, cookies, cakes, muffins, and snacks are popular products and consumed all over the world. New techniques and methods have evolved with the progress of bakery industries. Recently, bakery industry has also started making products that provide additional health benefits like fortified with specific nutrient(s) by using food sources. Mushrooms are one of the best nutrient-rich sources and can be a good alternative option for bakery products' enrichment with nutrients (Salehi, 2019), as displayed in Table 10.3. For enrichment or making bakery products edible, mushrooms are used in powder form. It provides additional protein, vitamins, mineral polyphenols, and crude fiber in bakery products. Recently, various researchers have tried to formulate and produce functional bakery products with mushrooms without the defect of their quality characteristics and sensory attributes (Lin et al., 2008). It has been reported that properties like color, physico-chemical, and texture of bakery products were affected by the replacement of wheat flour with mushroom powder, and it was observed that addition of 4–10% mushroom powder in bakery products showed the best sensory attribute. Rai and Arumuganathan (2008) prepared and analyzed various mushroom powder value-added products like biscuits, cakes, nuggets, soup powder, noodles, pickles, candies, ketchup, etc. There are several studies already present in literature that show mushrooms are potentially used for making bakery products, as shown in Table 10.3, such as bread, biscuits, and cakes. By adding mushroom powder in bakery products, it provides valuable products that are highly nutritious and can be a good source of functional food that can be consumed by a wide group of consumers.

There are some steps in the production of mushroom-based bakery products including the processing of harvested mushrooms, and their preservation and addition in the bakery product. Because of the perishable and short shelf-life of harvested mushrooms, it is processed after harvesting. Therefore, preservation and drying methods are applied. Dried mushrooms are easy and more valuable for mixing or making bakery products. Mushroom contains various metabolites; therefore, a correct drying method is applied for the drying process. Different drying methods are reported by various scientists (Giri & Prasad, 2007; Ghanbarian et al., 2016; Salehi et al., 2016; Salehi, 2019), such as sun drying, hot air, freeze drying, infrared vacuum, and dehydration being some drying methods applied according to mushroom species and physiochemical condition. For the selection of a particular drying method, it is also considered that it does not give unwanted aroma or color to the final product. Before making bakery products, mushroom must be prepared in powder form by using an efficient method (Ibrahium & Hegazy, 2014; Salehi et al., 2016). The work is based on different drying methods on shiitake mushrooms and achieves good color, improves rehydration and better aroma, etc. (Wang et al., 2015). Salehi et al. (2017) studied the drying kinetics of *A. bisporus* mushroom,

Table 10.3 Mushroom species used in bakery products and their supplementation percentage

| Bakery product | Mushroom | Quality of the product | Supplementation (%) | References |
|----------------------------|---|---|---------------------|-----------------------------|
| Bread | <i>Pleurotus ostreatus</i> | Increased protein, fiber and ash | 25 | Azeez et al. (2018) |
| | <i>Pleurotus ostreatus</i> and <i>Calocybe indica</i> | Increased protein, fiber and ash | – | Oyetayo and Oyedeji (2017) |
| | <i>Pleurotus sajor-caju</i> | Increased nutrients | 5 | Mahamud et al. (2012) |
| | <i>Lentinula edodes</i> | Increased nutrients and ash | 5 | Lin et al. (2008) |
| | <i>Pleurotus pulmonarius</i> | Increased protein, fiber, and ash | 10 | Okafor et al. (2012) |
| | <i>Lentinula edodes</i> | Less microbial counts and increased nutrients | 5 | Yen et al. (2011) |
| | <i>Agaricus blazei</i> , <i>Anthrobia camphorata</i> , <i>Hericium erinaceus</i> , <i>Phellinus linteus</i> | Increased nutrients and ash, etc. | 5 | Ulziijargal et al. (2013) |
| | Oyster mushroom | Increased protein content | 6 | Majeed et al. (2017) |
| | <i>Shiitake</i> , porcini, and white button mushroom | Increased antioxidant capacity | – | Lu et al. (2021) |
| | <i>Lentinus tuber-regium</i> | Increased nutrients and ash | – | Lee et al. (2004) |
| Cookie and biscuits | <i>Pleurotus eryngii</i> | Improve the quality and antioxidant potential | 3 | Kim et al. (2010) |
| | <i>Lentinus edodes</i> | Increase nutrients content | – | Singh et al. (2016) |
| | <i>Pleurotus sajor-caju</i> | Nutritious and glycemic index | 8 | Ng et al. (2017) |
| | <i>Agaricus bisporus</i> | Increase nutrients content | 20 | Kumar & Barmanray (2007) |
| | <i>Pleurotus sajor-caju</i> | Increased nutrients | 10 | Bello et al. (2017) |
| | <i>Pleurotus pulmonarius</i> | Increased protein, fiber, and minerals | 20 | Ibrahim and Hegazy (2014) |
| | <i>Calocybe indica</i> | Increased protein and minerals | 10 | Rathore et al. (2010) |
| | <i>Pleurotus ostreatus</i> | Higher protein | 25 | Cornelia and Chandra (2019) |
| | <i>Tricholoma matsutake</i> | Antifatigue effect | – | Ning et al. (2020) |

(continued)

Table 10.3 (continued)

| Bakery product | Mushroom | Quality of the product | Supplementation (%) | References |
|-----------------------|-----------------------------|---|---------------------|-----------------------------|
| Muffin | <i>Lentinus edodes</i> | Increased protein and fiber | – | Kim and Joo (2012) |
| Cake | <i>Agaricus bisporus</i> | Increased the protein and ash | 10 | Salehi et al. (2016) |
| | <i>Pleurotus sajor-caju</i> | Enhance essential nutritional components | 4 | Aishah and Wan Rosli (2013) |
| | <i>Pleurotus ostreatus</i> | Increased the protein and ash | 15 | Singh and Thakur (2016) |
| | <i>Pleurotus ostreatus</i> | Higher protein and Fat | 15 | Sheikh et al. (2010) |
| 3D snacks | <i>Agaricus bisporus</i> | Useful for personalized nutrition | 20 | Keerthana et al. (2020) |
| Cheese spreads | <i>Pleurotus ostreatus</i> | Increased nutrients composition and mineral | 2 | Khider et al. (2017) |

with combined IR vacuum drying. After drying, mushrooms are ground and made powder. Various scientists reported the nutritional and chemical composition of edible mushrooms in both fresh as well as dried forms.

10.5 Bakery Product Development from Mushroom as New Ways for Entrepreneurship

10.5.1 Bread

Bread is traditionally made by the process of fermentation of protein-rich sources. Salehi et al. (2016) reported that *A. bisporus* contains high-quality protein and can be a good source for making bread. Okamura-matsui et al. (2003) mixed mushroom powder with wheat flour and baked the mixture to make bread and determined the effect of mushroom supplementation on the functional properties of bread. It has been recorded that 10% addition of mushrooms (*Grifola frondosa*, *Hypsizygus marmoratus*, and *Pholiota nameko*) decreased the loaf volume of bread. Similarly, Lin et al. (2008) reported the substitution of *L. edodes* stipe for 2–7% of wheat flour for bread making and observed that the 5% shiitake stripe bread showed a higher fiber content with no interference with the bread's specific volume. Likewise, when a specific compound is isolated from mushroom like 5% of fungal chitin from shiitake mushrooms is mixed with wheat flour, it shows all similar to normal wheat

bread sensory results. It reported slow moisture loss and fewer microbial counts during storage as compared to wheat bread (Yen et al., 2011). It is also reported that increasing mushroom content significantly increases nutrient content in bakery products. Aishah and Wan Rosli (2013) formulated paratha flat bread, rice porridge, and cake with the dried oyster powder and analyzed the increased percentage of moisture, ash, and protein with respect to oyster powder content. Mahamud et al. (2012) studied wheat bread fortification by oyster mushroom at three levels – 5%, 10%, and 15% – and concluded that 5% supplementation gave better texture and acceptability. It is also observed the increased nutritional composition of bread compared to control bread with mushroom powder. Ulziijargal et al. (2013) also reported that mushroom mycelia (5%) are used for the supplementation of bread and observed that mycelium-supplemented bread showed higher umami taste intensity than control white bread. Lu et al. (2021) worked with three different species of mushrooms powder (shiitake, porcini, and white button mushroom) for wheat bread supplementation and investigated starch (gelatinization and digestibility) and its antioxidant capacities. This study reported that total starch contents decrease and increase phenolic contents of the bread with increased mushroom powder supplementation. Mushroom supplementation enhanced antioxidants capacity that was observed in the DPPH radical scavenging assay and oxygen radical absorbance ability of bread.

10.5.2 Cake

A natural fiber from natural products gives good texture and firmness to the cake or can say that quality and quantity of fiber make cakes better. Quality makes a good sensorial attribute and obtained positive acceptance by consumers. It has been observed by Jeong and Shim (2004) and Salehi et al. (2016) that the physicochemical properties of sponge cake are changed by the addition of mushroom powder. In cake making, the addition of 4% powdered oyster mushroom gives higher scores for softness and flavor attributes. It is also a cost-effective way for protein and nutrient enrichment in normal wheat bread. Fibers in cakes decreased the volume and increased the firmness, or all the changes observed in cake textures were dependent upon the fiber content in the used mushroom powder. Salehi et al. (2016) supplemented cake with *A. bisporus* powder and studied the rheological, textural, physicochemical, and sensory attributes prepared of cake batters as well as cake. In this report, it was observed that increasing the percentage of *A. bisporus* powder significantly increased the protein and ash content in cake. This study observed that the viscosity and volume of cake batters, cohesiveness, and springiness of baked cakes increased with addition of mushroom powder, but it was also observed that it reversed value in some properties like consistency, hardness, gumminess, chewiness, density, and crumb color indexes. The result indicates that 10% mushroom powder is most accepted in the sensorial evaluation test. Another study by Sheikh

et al. (2010), designated to the effect of mushroom powder addition on the quality of cake, found that 15% of mushroom addition gave the best acceptability.

10.5.3 Biscuits

Mushroom is renowned for its unique flavor and low-fat content. It is successfully utilized for making biscuits and cookies (Aishah & Wan Rosli, 2013). Mushrooms such as *L. edodes* and *P. eryngii* powder have been used for muffins and cookies making by Kim and Joo (2012) and Kim et al. (2010). Although the optimum amount of mushroom additive for product quality differs with the type of baked products and the selection of mushrooms used. The addition of mushrooms is a promising way to produce a healthy product with compatible quality characteristics to wheat flour bakery products (Moon & Lo, 2013). They formed biscuits by the substitution of wheat flour at three levels (10, 20, and 30%) with *P. pulmonarius* mushroom powder along with sweet potato flour (Ibrahium & Hegazy, 2014). This study reported that supplemented biscuits with 10 or 20% of mushroom powder and sweet potato flour exhibited a good sensory attribute. It determined the supplementation effect of oyster mushroom powder at three levels (0, 4, 8 and 12%) on the nutritional values, thermal and pasting properties, microstructure, in vitro starch digestibility, and sensorial properties of biscuits (Ng et al., 2017). They found significant increasing in the level of protein, total dietary fiber, ash, and β -glucan, with the 12% mushroom powder supplementation, etc. Another study by Singh and Thakur (2016) observed improved nutrient, physicochemical, and sensorial properties of biscuits by the supplementation of shiitake mushroom powder. This experiment also observed that the crude protein content of biscuits increases with the addition of shiitake mushroom powder. Biscuit is not much healthier for daily use because of its high carbohydrates, calories, and fat contents, but also due to its low vitamin, fiber, and mineral element contents and also having only 6–7% protein, etc. So, by using the fortification of biscuits by using mushroom powder it makes its more protein-rich source and healthier for daily use bases (Farzana & Mohajan, 2015; Serrem et al., 2011). They fortified soy flour to wheat flour with *P. ostreatus* powder and studied the effect on nutritional properties and sensory quality of biscuits (Farzana & Mohajan, 2015). A similar experiment was performed by Bello et al., 2017, who analyzed the nutrient composition and sensory characteristics of biscuits containing oyster mushroom (OM) powder. It was found that the protein content increased with mushroom powder addition. It was also observed that minerals such as K and Na are the main mineral elements in the biscuit samples and Ca, Mg, P, Fe, Cu, Zn, and Mn composition increases with mushroom powder addition. They studied functional, rheological, and physicochemical properties of biscuits that are made by gluten-free dough supplemented with fermented and unfermented polysaccharide flour of mushroom (Sulieman et al., 2019). The result signifies that *A. bisporus* polysaccharide flours in gluten-free dough formulation, having a plentiful amount of minerals, protein, essential amino acids, and rich amount of proteins,

etc. It for the first time reported the development of mushroom-based fiber-enriched snacks by using 3D food printing technology. In this technique, they used *A. bisporus* mushroom powder at different percentages (5, 10, 15, 20, and 25% w/w) along with different printing speed. They found that the 20% mushroom powder formulation printed at 800 mm/min printing speeds using the 1.28 mm nozzle with 0.383 g/min flow rate gave the best results (Keerthana et al., 2020). These printed snacks are further processed in microwaves and found that the spice-flavored snacks have an excellent sensory acceptance.

10.6 Concluding Remarks and Future Prospects

The mushrooms are utilized since ancient times as a food source as well as a medicinal source. They have several nutrients and bioactive compounds; they are a well-known natural source that can be used in many different fields of human welfare. Continuous rise in the population and its advanced lifestyle raise the demand of food and medicine. All these consequences lead to generate such type of functional food that overcomes this problem and gives both food and medicine source in a single-step manner. Therefore, edible mushrooms are the most potential candidate that successfully utilized both these criteria. It is well-established research that the mushrooms are used as a functional food and are the source of bioactive compounds; furthermore, researches are still needed for unidentified mushroom metabolites. It is well known for its nutritional and bioactive compound and also can be preserved and utilized in food industries. Food fortification with mushroom extracts offers great potential for a future generation with very high importance of functional food.

There are various products made by mushrooms and utilized worldwide, mushroom-supplemented bakery products being one of them. Bakery products are the easy and most utilized food stuff all over the world, and its supplementation by using mushroom makes it more nutritious. These supplemented bakery products are easily consumed as a form of functional food. The supplementation of bakery products by mushrooms can help improve the nutritional requirements and status of the underdeveloped as well as developing countries. By increasing awareness of the various properties of mushroom and increasing utilization of mushroom in bakery industries, it can generate rise in mushroom demand. These mushroom demands can be fulfilled by increasing mushroom cultivation. Mushroom cultivation can give a contribution to sustainable development as well as provide a livelihood for rural populations.

Employment is a bigger challenge for the rural population. The majority of the rural population mostly depend on agriculture activity (farming), and farmers are still using traditional cultivation methods that result in low productivity. The agricultural activity produces a high number of by-products such as rice straw, wheat straw, stumps, barley straw, husk, sorghum stalks, grass seed straw, flax coil seed, corn stalks, sugarcane bagasse, etc. This agro-waste was combusted for disposal, which increases carbon emissions and harms the environment. Mushroom

cultivation is the best option to utilize this agro-waste and make value-added products. In future, mushroom cultivation can be the best alternative source of employment generation in rural and backward areas. Also, mushroom cultivation is a better technique for generating income, providing employment to the rural population, and making their livelihood improvement to rural population. In the current situation, we require a food source that provides additional nutrition, and food supplementation by mushrooms is considered as a better and quick option of functional food. Mushroom cultivation and its utilization of food sector are helpful for defending malnutrition to the local population. It is an easy and cost-effective technique that provides a new way for farming, providing functional food to the nutrient-deficient population.

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Chapter 11

Fungi in Pharmaceuticals and Production of Antibiotics



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11.1 Introduction

The contribution of microbes in the enhancement of human well-being and health is immense. The natural compounds that are obtained from the microbes not only helped in treating diseases but also helped in enhancing the average life expectancy in human. The invention of penicillin from the mold *Penicillium notatum* by Alexander Fleming in the year 1928 showed that microbial species are the storehouse of a large number of natural compounds that play important role in acting against various types of disease-causing pathogens. Fungi are one of the most important among those five kingdoms as they are essential for the good growth of the plants, different crops through the nutrient cycle by degrading the cellulose and lignin (Pointing et al., 2001). In spite of having numerous negative aspects, fungi possesses biotechnological potentials for industrial exploitation. The large use of antibiotics since its discovery has resulted in the development of various types of resistant microbes by the mechanism of evolutionary selection pressure driven by the antibiotics (Davies & Davies, 2010). Thus, classic therapeutics is being rendered ineffective in various types of infections along with resulting in morbidity due to the condition of multidrug resistance (MDR). The natural products produced by the primary and secondary metabolism of microbes have a broad range of therapeutic applications, which is further modified and then amplified through biotechnology for the betterment of society. Microorganisms are the most abundant sources of diverse bioactive metabolites. The pharmaceutical industries have extended their screening program to focus the metabolites from the microbes, effective toward different types of disease areas. Fungi are one of the groups of microbes capable of producing a diverse array of secondary metabolites that have been used in the pharmaceutical industry on a large scale. As the result of incorporation of the screening process, some most important products are immunosuppressants, antitumor drugs, hypocholesterolemia drugs, enzyme inhibitors, growth stimulants, insecticides, and herbicides. This chapter focuses on the use of the fungal metabolites in various types of drug development targeting diseases at large having a special focus on the development of antibiotics with the present-day need.

11.2 Different Roles of Fungi in Pharmaceuticals

Since the discovery of antibiotics in the 1940s, millions of lives were saved by antibiotics. Due to the excessive use of antibiotics, many of the bacteria develop their own protection against antibiotics and become antibiotic resistant. This can lead to the development of lethal infections that can be diminished by the alternative strategy. The discovery and development of the new antibiotic can accomplish the urge of the alternative strategy in order to reduce the lethal infections.

In 2018, the scientific community celebrated the ninetieth anniversary of the discovery of penicillin by Sir Alexander Fleming, from which the era of the antibiotic

chemotherapy has started. After the discovery of penicillin in 1928, numerous antibiotic compounds like cephalosporins, fusidic acid, and pleuromutilin were derived from the fungi, which are effective against several diseases.

11.2.1 *Antibacterial Antibiotics*

Antibiotics are the group of chemical substances that are produced by one set of microbial cells used for the purpose of inhibiting other microorganisms. When an organism is able to produce specific secondary metabolites, which can destroy the competing organisms present in the same habitat, then that secondary metabolite is considered as antibiotics. It is evident that most of the filamentous fungi produce antibiotic compounds (Bills et al., 2009).

In the post-antibiotic era, there is a drastic change in the number and percentages of multidrug-resistant organisms and antibiotics. The inverse relationship between the drug-resistant pathogens, which is increased with the decreased number of therapeutic agents, leads to the development of new antibiotics in order to combat the pathogenicity (Cooper & Shlaes, 2011). The loss of interest of the pharmaceutical companies in research and development on the natural product gave rise to consequences like the scarcity of the antibiotics in the treatment of the multidrug-resistant human pathogenic bacteria. After two decades of this crisis, the discovery of the new antibiotics has increased, which is basically nothing but the chemical modification of the old compounds with a known mode of action. There are few compounds under the pipeline that are advanced and yet to be used in the pharmaceutical sectors.

For example, fermentation-derived compound mutilins from the basidiomycete *Clitopilus passeckerianus* is considered as a new antibacterial drug. For skin infections, several derivatives are under clinical trials, among which a derivative retapamulin has been discovered to be used as the topical antibiotic agent. However, it is difficult to culture basidiomycete on a large scale due to its slow growth rate and low yields. It has been discovered that a transfer of the biosynthetic genes into a fast-growing heterologous host *Aspergillus* can enhance the production of secondary metabolites like pleuromutilin (Bailey et al., 2016). In the middle of twentieth century, around 22% of the known 12,000 antibiotics were produced by the filamentous fungi, among which the β -lactam antibiotics including penicillin, cephalosporin, clavulanic acid, and carbapenems are the most important antibiotic class (Berdy, 1995).

Due to the immense application of those antibiotics, the production needs to be increased to reach the optimum goal. 1,3-Diaminopropane (1,3-DAP) (secreted by *P. chrysogenum* and *Acremonium chrysogenum*) and spermidine increase transcription level of penicillin biosynthetic genes *pcbAB*, *pcbC* and *penDE* to stimulate penicillin G production (Martín, 2012). *laeA* is the global regulator that regulates expression of secondary metabolism genes via heterochromatin reorganization and gets stimulated via the mechanism. Also, the cephamycin production and spermidine activity in *Amycolatopsis lactamdrans* get stimulated by 1,3-DAP. Due to the

increase of antibiotic resistance among fungi and bacteria, novel antibiotics are in dire need. One example of the new and approved antibiotic is ceftobiprole, which is a modified version of cephalosporin, is active against methicillin-resistant *S. aureus*, and is not hydrolyzed by β -lactamases from Gram-positive (Shang, 2011). Another example of antibiotic that acts as an antifungal agent and also the first discovered inhibitor of fatty acid biosynthesis cerulenin is produced by *Acremonium caereleus*. The active site of the nucleophilic cysteine of the ketosynthase enzyme of fatty acid synthetase through epoxide ring opening gets alkylated and inactivated by this antibiotic (Vance, 1972).

In a pathogenic fungi *Candida albicans*, poly(A) polymerase gets inhibited by an antifungal product parnafungin from *Fusarium layarum*. Most of the Gram-negative bacteria are resistant to the carbapenem antibiotics, mainly imipenem and meropenem, which give rise to the major antibiotic problem due to the occurrence of New Delhi metallo- β -lactamase (NDM-1), the extended spectrum metallo- β -lactamase. NDM-1 and another metallo- β -lactamase VIM-2 are inhibited by the aspergillomarasmine (AMA) from *Aspergillus versicolor* (King, 2014). AMA is the amended version of meropenem that inhibits metalloproteinases of the bacteria carrying NDM and VIM metallo- β -lactamase (King, 2014).

11.2.2 Antimycotics and Fungicides

Apart from the multidrug-resistant pathogenic bacteria, there are a number of pathogenic fungi whose resistivity is also high. Several efficient compounds are available in the market for antimycotic chemotherapy. One of those compounds is griseofulvin, which is widely used. Echinocandins (pneumocandin B_o) is the new class of antimycotic that has been launched to the market (Denning, 2002). The PKS-NRPS hybrid gene cluster regulates the biosynthesis of highly complex lipopeptides. The significant knowledge about the molecular mechanism leads to incisive manipulation of the production process, which can result in the synthesis of the new derivative by using a diverse group of fungi for the large-scale fermentation of the subsequent biosynthesis of those gene clusters (Chen et al., 2013). It was evidenced that Dothideomycetes, Eurotiomycetes, and Leotiomycetes are different classes of the producer organism Ascomycota having the high homology of the gene cluster due to horizontal gene transfer during evolution (Yue et al., 2015). A substantial number of developmental projects for the discovery of the novel antifungal compounds is under pipeline. Enfumafungin is one of the antifungal drugs originated from the fungal endophyte *Hormonema* spp. used to treat human disease after 15 years of its first discovery (Pelaez et al., 2000). Rediscovery of the old compounds can accomplish the urge of the novel antimycotics and fungicides, which have their significant effectivity against human fungal pathogens. Very few of those compounds have toxic effect toward human, but most of them have strong antifungal effects without any cytotoxicity. For example, favolon is the rediscovered form of strobilurins, which is produced by the invasive basidiomycete *Favolaschia*

calocera (Chepkirui et al., 2016), and sporothriolodes is from xylarialean fungus *Hypoxylon monticulosum*, which is now classified under the new genus *Hypomontagnella* (*H. monticulosa*) (Surup et al., 2014; Lambert et al., 2019).

11.2.3 Biofilm Inhibitor

The discovery of the new antibiotics is mainly aimed to combat the infectious diseases caused by both the fungal and bacterial pathogens, which is possible through the inhibition of the biofilm formation. It has been reported that the fungal metabolites have the potency to inhibit the biofilm formation by interfering with a quorum sensing mechanism or by destroying preformed biofilms (Abraham & Estrela, 2016). A research showed that a small molecular derivative of edible mushroom *Coprinus comatus*, coprinuslactone, acts against the *Pseudomonas aeruginosa* biofilms (de Carvalho et al., 2016). The biofilm of the Gram-positive bacteria *Staphylococcus aureus* is inhibited by roussoellenic acid from *Roussoella* sp. (Phukhamsakda et al., 2018) and microporenic acid A from *Kenyan basidiomycete* (Chepkirui et al., 2018). A human pathogenic yeast *Candida albicans* can also be destroyed by microporenic acid A. The efficacy of an antibiotic is enhanced by the biofilm inhibitor that can increase penetration through the biofilm. Though some fungal antibiotics are available and have their efficacy toward the infections, a vast range of fungi and their applications in novel antibiotics are still underexplored.

11.2.4 Antimalarial Agent

In spite of tropical and subtropical areas, worldwide malaria is one of the major causes of illness and death. Around 500 million new cases are detected every year where 1.5 million, people especially pregnant women and children, die due to this. The two major drugs quinine from the bark of the *Cinchona* tree and artemisinin from the Chinese herb *Artemisia annua* have been used successfully against malaria. In spite of being a new drug, artemisinin is used extensively due to the side effects of the quinine like arrhythmia, thrombocytopenia, and cinchonism. Endoperoxide sesquiterpene lactone artemisinin is also effective against multidrug-resistant *Plasmodium falciparum*. Genetic engineering has been implemented as very low amount of artemisinin is obtained from the *A. annua*, 0.01–1% of the weight of the dried leaves. A genetically engineered *S. cerevisiae* strain, which is able to produce 100 mg l⁻¹ of artemisinic acid, has been developed by the Keasling group in Berkeley, California, where further artemisinic acid is chemically converted to artemisinin and artemisinic acid production increased by 1 million fold. Also, by the engineered *S. cerevisiae* the artemisinin precursor amorpha-4, 11-diene is made at 40 g/L (Liu et al., 2013).

11.2.5 Anticancer Agents

After the cardiovascular disease, the second leading cause of mortality is cancer, where nearly 9.6 million cancer-related death cases had been reported. With the loss of growth factors, the uncontrolled cell division and cell proliferation are termed as “cancer.” The presence of cancerous cell is referred to as malignancy, which can invade nearby and destroy tissues by suppressing the human immune system. Several treatment processes like chemotherapy, radiation therapy, surgery, and immunotherapy have been administered according to the developmental stage of cancer disease. A wide range of cytotoxic agents are used to suppress cancerous tumor cells. Advanced cancer treatment therapy is based on regulating the gene or protein of the specific cancer cells by the drugs. Though the most used drug albeit has originated from plant and bacteria, some important natural products can be obtained from fungi for the development of novel drugs, and in most cases, those drugs are at clinical or preclinical stages. Irofulven is a semi-synthetic derivative of the natural toxin illudin S, isolated from *Omphalotus illudens* (Chin et al., 2006). Irofulven has antitumor activity, which has been evaluated at the phase I and phase II clinical trials against brain, central nervous system, breast, blood, colon, sarcoma, prostate, lungs, and ovarian and pancreas cancers (Alexandre et al., 2004). Irofulven interacts with the DNA replication complexes and inhibits the DNA synthesis to stop the abnormal cell divisions (Walser & Heinsteins, 1973). A tetracyclic diterpene aphidicolin was isolated from *Cephalosporium aphidicola* (Bucknall et al., 1973), competes with DNA polymerase α , δ , and ϵ enzymes for specific binding sites, and has the antiviral and antimetabolic properties (Crosetto et al., 2013). There are other anticancer lead compounds derived from the fungi, among which leptosins derived from *Leptosphaeria* sp. have the antitumor activity in mouse embryos (Pejin et al., 2013). Some natural compounds like β -glucans, palmarumycin, and spiroprossione A are found on fungi cell wall and also have the anticancer potency, which is yet to be explored.

The fungal and plant secondary metabolites taxol (paclitaxel isolated from pacific yew tree, *Taxus brevifolia*) and camptothecin have showed marked result in the clinical trial of anticancer drug. Taxol is a steroidal diterpene alkaloid with tetracycline ring and N-benzoylphenyl isoserine side chain that inhibits the mammalian cancer cells from dividing by promoting tubulin polymerization and interfering microtubule breakdown during cell division. The taxol was approved against the refractory ovarian cancer in 1992, and further it was used to treat breast cancer and advanced Kaposi's sarcoma (Newman & Cragg, 2007). Certain plants (angiosperms) and fungi (endophytic), *Entrophospora infrequens*, produce a modified indole alkaloid, camptothecin, which is used as an important antitumor agent and also used to treat colon cancer, unusual activity of lungs, ovarian, and uterine cancers (Amna, 2006).

11.2.6 Antidiabetic Agents

Diabetes mellitus is a chronic metabolic disorder, where insulin is not produced or effectively used in the body. Diabetic patients have high sugar of glucose in their body to the insulin imbalance. There are two types of diabetes, type 1 (insulin dependent) and type 2 (noninsulin dependent). In case of type 1 diabetes patients, due to the malfunctioning of the pancreatic beta cells, insulin is not produced sufficiently. Insulin needs to be injected from outside of the body. Worldwide, around 5–10% people, mostly children and adolescent patients, are type 1 diabetes patients (Meier et al., 2005). Rest 90–95% patients suffer from type 2 diabetes, where their body is not able to produce sufficient insulin or cannot effectively metabolize it. Around 7% of the world's elder population are affected by this disease. In 2017, the largest number of diabetic patients, around 114 million, was recorded in China. Roughly 73 million diabetic patients were recorded in India, and 30 million were recorded in the United States.

Untreated diabetic patients can be the sufferer of blindness, kidney failure, depression, cardiovascular diseases, cancer, and even death. The most common complications that are attributed to the diabetes are retinopathy and neuropathy (De Silva et al., 2016).

For the treatment or prevention of type 2 diabetes, Basidiomycota such as *Agaricus bisporus*, *Cyclocybe aegerita*, *C. cylindracea*, and *Tremella fuciformis* are used as alternative medicine as they contain very less amount of the digestible carbohydrate, which can avert high glucose intake of patients (Poucheret et al., 2006). It has been reported that the extracts of *Inocutis levis* have the ability to increase insulin resistance and sensitivity and also control blood glucose levels by increasing glucose uptake in tissues (Ehsanifard et al., 2017). Hyperglycemia (when blood glucose level is high) and hyperinsulinemia (when blood insulin level is high) both along with type 2 diabetes can be treated by *Grifola frondosa* (Poucheret et al., 2006). Some antidiabetic medicinal products like *Ophiocordyceps sinensis* capsules, SX-Fraction (enhance insulin sensitivity), Reishi-Max capsules, and Tremella are made with medicinal mushrooms, used to decrease blood glucose level of type 2 diabetes and reduce blood pressure and body weight (Li et al., 2004).

11.2.7 Improving Nerve Function

Millions of people are affected worldwide by the most common Alzheimer's, Huntington's, and Parkinson's neurodegenerative diseases. To improve nerve functioning, the discovery and development of the neuroactive compounds from the medicinal mushroom are studied to a great extent. Earlier it has been discovered that *Antrodia camphorate*, *Ganoderma* spp., *Hericium erinaceus*, *Lignosus rhinocerotis*, and *Pleurotus giganteus* can enhance the peripheral nervous system. Most of the neurodegenerative diseases occur due to the disappearance of nerve growth factor as

it is one of the important materials for the adult brain survival, maintenance, and regeneration of specific neuronal populations. Since the past 20 years, scientists are trying to discover some drug components from the fungi that can easily penetrate the blood–brain barrier and induce production of nerve growth factor. *Hericium erinaceus*, the medical mushroom, is mainly used to discover potential neuroactive compounds for the prevention of the neurodegenerative diseases. Hericenones and erinacines, the two terpenoid group of this medical mushroom, stimulate the nerve growth factor synthesis via TrkA/Erk1/2 pathway. Apart from inducing nerve growth factor, this medicinal mushroom also provides the immune support by improving digestive function and enhancing anti-inflammatory and antioxidant activities. It was demonstrated that hericenone A and erinacines C can induce synthesis of nerve factor both in vivo and in vitro (Thongbai et al., 2015). Cyathane diterpenoids, derived from *Cyathus* species, is reported as nerve growth factor enhancer, where the similar effect was observed in the cyathanes from the mycorrhizal basidiomycetes (genus *Sarcodon*) fruit bodies (Bai et al., 2015).

11.2.8 Cardiovascular Disease Control and Cholesterol-Lowering Agent

Heart and blood vessel diseases are termed as “cardiovascular disease.” which can cause the average death of 17.3 million people per year due to the elevated level of plasma cholesterol, by which arteries get clogged due to fat deposition. Around 30% of the cholesterol comes from diet and rest 70% is synthesized in the human liver. Only healthy diet is not able to control the cholesterol level in most cases, which leads to the high blood cholesterol level causing atherosclerosis, a chronic disease due to the accumulation of atheromatous plaque within the arterial wall leading toward stenosis and ischemia. Scientists came across that plasma cholesterol level can get reduced by the inhibition of de novo synthesis of cholesterol (Miller, 2001). The reduction of cholesterol is possible by inhibiting the rate-determining step where 3-hydroxy-3-methylglutaryl-coenzyme A (HMG-CoA) is reduced to mevalonate by HMG-CoA reductase (Brown et al., 1976).

Fungal secondary metabolites, hypolipidemic drugs statins are the HMD-CoA reductase inhibitor, have two different moieties: hexahydro-naphthalene system and β -hydroxylactone system. Compactin was isolated from *Penicillium brevicompactum* and has the antibiotic as well as hypocholesterolemic activity. Statin lowers the cholesterol level by inhibiting the rate-limiting enzyme 2-hydroxymethyl glutaryl-CoA of mevalonate pathway of cholesterol biosynthesis. This inhibition mechanism of the enzyme stimulates low-density lipoprotein receptors in the liver, resulting in the augmentation of low-density lipoprotein clearing from the bloodstream in order to decrease the cholesterol level. Hence, the risk of cardiovascular disease is reduced with the reduction of total plasma cholesterol by 20–40% (Endo et al., 1976).

Mevinolin is another HMG-CoA reductase inhibitor that was isolated from *Aspergillus terreus* (Alberts et al., 1980). In medicinal mushroom, certain molecules are present that are able to modify cholesterol absorption, metabolism, and modulate gene expression of cholesterol homeostasis (Gil-Ramirez et al., 2016). Some other cholesterol-emic product like compactin (from *Penicillium citrinum*) and lovastatin (from *Monascus rubra* and *Aspergillus terreus*) had been developed at the same time, which has strong medical use with the commercial success.

11.2.9 Antiviral Agents

Virus can cause a serious global epidemic through the health and mortality-deliberating diseases due to insufficient or unavailability of the antiviral chemotherapy or vaccines. Some drug-resistant viral strains are responsible for the current virus-related pandemic due to the loss of drug efficacy. Hence, to combat this viral epidemic by controlling viral infections, a new strategy related to the discovery of the natural drug leads needs to be devised. There are several natural compounds screened for the antiviral activity but are yet to reach the market; one of those compounds is plethora, isolated from fungi. Scientists are more focused on discovering natural compounds that can exhibit potent activity against human pathogenic viruses.

11.2.9.1 Inhibitory Natural Product from Fungi Against Human Immunodeficiency Virus

The discovery of the anti-HIV drug is based on virus entry, reverse transcription, and integration. The discovery of the viral entry blocker is based on the HIV interactions with proteins. HIV-1 protease, which is a key enzyme for replication and maturation of HIV-1 virus, can be inhibited by bis-indolyl quinone, hinnuliquinone isolated from *Quercus coccifera* (Singh et al., 2004). At micromolar concentration, altertoxins I–III and V (oxidized perylenes) from *Alternaria tenuissima* inhibit HIV-1 replication (Bashyal et al., 2014). A strong anti-HIV activity was observed by the dimeric tetrahydroxanthone and penicillixanthone A, from *Aspergillus fumigatus*, as they are able to inhibit CCR5-tropic HIV-1 SF162 and CXCR4-tropic HIV NL4-3 (Tan et al., 2017). Anti-HIV-1 activity is also observed by malformin C, from marine-derived *A. niger* (Zhou et al., 2015). Phenalone and cytochalasin derivatives of endophytic *Aspergillus* sp. CPC 400735 showed anti-HIV activity (Pang et al., 2017). Moderate anti-HIV-1 replication was observed in C8166 cells by novel sesquiterpenoids from *Paraconiothyrium brasiliense*, whereas significant HIV-1 activity was observed by pupukeanane sesquiterpenoid chloropupukeannolide A from *Pestalotiopsis fici* (Liu et al., 2010). The discovery of anti-HIV drug depends on the inhibition of three consecutive functions (RNA reverse transcription to DNA, degradation of RNA template by RNase H, and duplication of the remaining DNA

strand) controlled by HIV reverse transcription. *Stachybotrys chartarum*-derived phenylspirodrimane metabolite Stachybosin D has the inhibitory effects on HIV-1 replication by targeting reverse transcriptase in wildtype HIV-1 and NNRTIs-resistant strains (Ma et al., 2013). Along with protease and reverse transcriptase, integrase is one of the important proteins encoded by HIV-1. It was reported that several compounds like equisetin, phomasetin, integracins, epiphiobolins C and K, hispidin, caffeic acid, naphtho- γ -pyrones, xanthoyiricacins, etc. have the inhibitory activity against integrase.

11.2.9.2 Inhibitory Natural Products from Fungi Against Influenza Virus

Among all the influenza viruses, H1N1 and H3N2 are the most targeted ones for the development of the drug. IC₅₀ value determines the moderate to high effectivity of drug molecules against H1N1 and H3N2 viruses. The terpenoid isolate of *Stachybotrys*, stachyflin, has the modest activity against influenza A (H1N1) virus with IC₅₀ value of 39×10^{-3} μ M (Minagawa et al., 2002). Low micromolar activity against both the influenza viruses was observed by γ -pyrone isoasteltoxin from *Aspergillus ochraceopetaliformis* (IC₅₀ = 0.23 μ M) (Wang et al., 2016). γ -pyrone is a derivative asteltoxin E from *Aspergillus* sp. active against H1N1 and H3N2 (Tian et al., 2016). H3N2 influenza A and B virus replication was inhibited by the metabolite of *Gliocladium* sp. aureonitol, with an EC₅₀ value 100 nM, via influenza hemagglutination suppression (Sacramento et al., 2015)

11.2.9.3 Inhibitory Natural Products from Fungi Against Herpes Simplex Virus (HSV)

There are no such drug molecules against HSV-1 and HSV-2 pathogenic viruses available in the fungal metabolite library till date. Very few drug molecules had been reported with anti-HSV activity (Huang et al., 2017). *Scytalidium* sp. fungus derived five lipopeptides that have moderate anti-HSV-1 and anti-HSV-2 activities, which are dose and time dependent (Rowley et al., 2003). The diphenyletherglycoside cordyol C from *Cordyceps* sp. exhibited significant anti-HSV-1 activity (Bunyapaiboonsri et al., 2011).

11.2.9.4 Inhibitory Natural Products from Fungi Against Hepatitis Virus

Vanitaracin A is a novel tricyclic polyketide fungal derivative that inhibits the viral entry process. This compound can inhibit all HBV genotypes (A–D) through interacting viral bile acid transport pathway (Kaneko et al., 2015).

Through the inhibition of the Akt activity and depletion of autophagic genes LC3 and p62, the HBV-X replication is decreased by epipolythiodiocopiperazine derivative, 11'-deoxyverticillin (Wu et al., 2015). From nematode-trapping basidiomycete

Hohenbuehelia grisea, anti-HCV agent 4-hydroxypleurogrisein has been discovered (Sandargo et al., 2018). Meroterpenoid rhodatin from rare basidiomycete *Rhodotus palmatus* has the notable anti-HCV activity (Sandargo et al., 2019). Since the viral diseases are increasing day by day, the discovery of the novel antiviral molecules is gaining importance. Most of the natural compounds with nanomolar activity are at the pipeline of the drug discovery.

11.2.10 Immunosuppressive and Immunomodulatory Agents from Fungi

Immunosuppression is a therapeutic process that accounts for preventing the immunity of various patients from defending against organ transplant and tissue transplant. The entire process takes place in the absence of any type of modern medicines, presence of drugs that are immunosuppressive, any type of modern medicine related to transplantation heart, liver, and kidney. Immunosuppressants also possess the ability to control many manifestations of some autoimmune diseases and allergic conditions. In these drugs, majority of them specially devote themselves to some specific biochemical pathways, which are very important for the proper functioning of the defense mechanism of human immunity in response to invading pathogens or microorganisms coming from outside. This is done by selective inhibition of the signal transduction cascade of pathways or the immunocompetent lymphocytes. These cascade pathways or cascade of steps undergo regulation of the cytokine transcription process. This results in the hospitalization and antibiotics treatment of the patients undergoing immunosuppressants for preventing the occurrence of infection in them. Few of the noted immunosuppressive drugs are simply some natural products that are artificially formed by the application of biotechnology that is by fermentation of fungi and bacteria. The compounds tacrolimus and sirolimus are isolated from actinobacteria. Moreover, mycophenolate and cyclosporine mofetil are secondary metabolites formed by fungi. But it is also known that penicillin is considered one of the oldest natural antibiotics available. The first antibiotic invented and extracted from fungi in its crystalline form is a meroterpenoid that is formed by *Penicillium* species, which consists of *P. brevicompactum* and *P. roquefortii* (Del-Cid et al., 2016), and then undergo their process of biosynthesis that is elucidated recently in the other species. Because of different reasons, the compound did not reach its clinical development phase as an antifungal agent and antibacterial agent. So basically, the main purpose as an immunosuppressant is to become an active component of many marketed drugs like those of Myfortic® and CellCept® and to become evident. This compound undergoes selective inhibition to form inosine monophosphate dehydrogenase (IMPDH). Along with this, it also produces an enzyme that is important for the process of biosynthesis of guanosine nucleotides into the cells of mammals. This is because this enzyme is needed more inside the B-lymphocytes and T-lymphocytes than in other cell types (Allison & Eugui, 2000).

Also, the compound and its isotypes, along with the isoform in the lymphocytes, possess more sensitivity to mycophenolic acid, the drug that has more efficient cytostatic effect on lymphocytes than on the rest of the type of cells, and by this entire process, human immune system suppression takes place.

11.2.11 Traditional Chinese Medicine

Traditional Chinese medicine (TCM), scattered in Japan and other Asian countries, has been used since thousands of years. Use of natural remedies (mainly herbal medicines) is the main objective of TCM for disease therapy. The first Chinese medicine was recorded in the eleventh-century BC (1100–1001 BC), and throughout the previous centuries combination of conceptual theories and practical experiences helped to evolve the TCM. Compared to Western medicine, TCM has more holistic ideology in the pharmaceutical field. “Protector herb” or the Chinese medicinal mushrooms are being used for enhancement and maintenance of good health. Majority of Chinese medicines are plant-derived and in very few macrofungi are used. Several bioactive compounds are extracted from the fungal sources, are considered to have the potency against various diseases like cancer, diabetes, and cardiovascular and neurodegenerative diseases, and are used in fungal TCM. Due to the unavailability of a proper description, there are thorough debates regarding the utility and therapeutic applications of TCM in the modern healthcare sectors. Western medicine is more focused to develop drug from a single substance depending upon strong selectivity and pharmacological activity for the therapy of disease. TCM is a mixture of compounds that are isolated from plants, fungi, and other organisms, possessing the ability to obstruct the TCM to fit into the modern healthcare. Still there are many evidences that claim the TCM to be approved in the healthcare system (Tang et al., 2018).

11.3 Pharmaceutical Uses of Fungal Metabolites

Fungi have an important role in concurrent drug discovery and pharmaceuticals. Some fungal metabolites have a remarkable activity against the pathogenic organisms.

11.3.1 Cephalosporin

A broad spectrum of antibiotics, cephalosporins, is usually used to treat penicillin-allergic patients or against the infections resistant to penicillin. Since the 1950s, a large number of cephalosporins have been used as pharmaceutical.

Cephalosporin was isolated from the fungus *Cephalosporium acremonium*, which has potent antibiotic activity against both the Gram-positive and Gram-negative bacteria. Though cephalosporin N and cephalosporin C are closely related compounds, cephalosporin has the greater beta-lactamase withstanding ability by possessing low toxicity (Crawford et al., 1952).

11.3.2 Coumarins

Fungal metabolite dicoumarol-derived warfarin acts as an anticoagulant and anti-thrombotic. Dicoumarol is a hemorrhagic agent from which anticoagulant is formed by fungal oxidation of coumarins to 4-hydroxycoumarins along with its subsequent coupling with formaldehyde.

11.3.3 Cyclosporin

A wide range of biological activities like antifungal, antibacterial, and immunosuppressive activities were found in the cyclic peptide cyclosporin. Use of this drug was approved in 1983, and now it has wide application as first-line drug in the prevention of organ transplantation rejection. Cyclosporin A was isolated from the fermentation broth of *Tolypocladium inflatum*, which showed low antifungal but high immunosuppressive activity (Stähelin, 1996).

11.3.4 Echinocandins

An antifungal agent echinocandin inhibits fungal b-1-3glucan synthesis, which affects fungal cell wall (Saravolatz et al., 2003), mainly used to treat invasive fungal infections from *Candida* and *Aspergillus*. Due to the undesired hemolytic activity, semisynthetic investigations were performed to overcome this shortcoming. Pneumocandin B₀ and echinocandin B are the two precursors used in pharmaceutical. The modification of hexapeptide increases the antimycotic activity of pneumocandin B₀.

11.3.5 Ergot Alkaloids

The ergot alkaloid has the dopamine receptor agonist abilities to treat Parkinson's disease. The two fungal families Clavicipitaceae (e.g., *Claviceps* and *Neotyphodium*) and Trichocomaceae (e.g., *Aspergillus* and *Penicillium*) are responsible for the

production of ergot alkaloids. The use of ergot started from 1582 according to the records. This ergot alkaloid is also used as an antimigraine drug. Ergotamine and caffeine combination proved to be best for migraine treatment. It was found that ergometrine from *Claviceps* sp. can induce uterine contraction for which it is used in front-line pharmaceutical to treat postpartum hemorrhage (Wallwey & Li, 2011).

11.3.6 Fingolimod

Fingolimod is an immunomodulatory compound of fungal metabolites that is used as oral first-line treatment for multiple sclerosis. The antibiotic activity was observed in the compound myriocin, isolated from *Myriococcum albomyces*, a thermophilic microbe (Kluepfel et al., 1972). Though this compound has antibiotic activity, a significant result was not observed in the clinical study.

11.4 Biotechnology in Sustainable Production of Antibiotics

The discovery of the highly pure form of antibacterial product from specific species is done through extraction, followed by screening of the antibacterial activity and isolation process. This whole process of natural product discovery is slow, monotonous, laborious, and inept. Biotechnology contributes to the advancement of this antibiotic discovery process by controlled miniaturization for increased screening throughput. “System Duetz” (Duetz et al., 2000) and “BioLector” (Samorski et al., 2005) are used as specialized small-scale fermentation system under the controlled biotechnological approach (Fig. 11.1).

11.4.1 Biotechnological Processes-Associated Production

The most crucial step for drug development is production of natural products, which is achieved by different biotechnological approaches. Full fermentative processes using the natural producer, semi-synthetic approaches, and heterologous production in genetically modified hosts are the best three approaches of production.

11.4.1.1 Production Using Full Fermentative Processes

Filamentous fungal biotechnology is implemented for the industrial-scale fermentation of the antibiotic products (Ng et al., 2015). Most of the antibiotic production strategies were established shake flask (Erlenmeyer flasks, EMF) level, with successful scale-up. To obtain sufficient amount of isolated compounds, large-scale

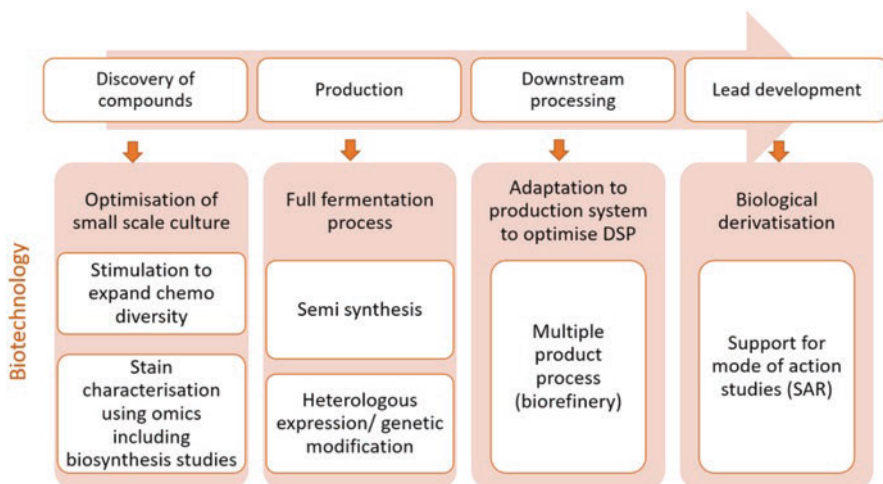


Fig. 11.1 Biotechnological approach in the production of antibiotics from fungi

fermentation is performed in multiple EMF frequently. The transfer of cultures from EMF to bioreactors under controlled conditions is required for the successful and economically expedient scale-up. Successful scale-up processes have some strategies where operating factors like dissolved oxygen, carbon dioxide, pH, temperature, aeration, and foam have influence over antibiotic production. In fungal biotechnology, stirred tank reactors (STR), moving bed, and solid-state system are most commonly used, and terrestrial fungi have been proved to be technically and economically feasible for large-scale production (Riley et al., 2000). Production of antibiotics from fungi is also dependent on the successful transfer into STR, where the controlled process can increase yield by a factor of 100. One of the examples of this process is production of tetramic acid compound ascocetin and lindgomycin by arctic fungus. Application of some specific experimental condition like high pressure for the cultivation of obligate fungi stimulates the expression of gene cluster for new antibiotic production. It was observed that 15 L of 3, 5-dihydroxydecanoic polyester exophilin A had been obtained from the 20-L culture of *Exophiala pisciphila*, member of “black yeast” isolated from the sponge *Mycale adhaerens* in a glass-bottle fermenter under controlled pH (Doshida et al., 1996), where the antibacterial activity increases from 6 days of the cultivation and reaches the highest after 10 days. Carbon and nitrogen also played an important role in compound production for terrestrial bioprocess (Xu et al., 2012). Along with pH and alternation of mycelial morphology, the adaptation of carbon to nitrogen ratio increases the production of macrocyclic polyester calcaride A from *Calcarisporium* sp. by 200-fold (Tamminen et al., 2015). Adding to the physicochemical parameters’ morphology is one of the important parameters, which influences biosynthesis of the product. From dispersed filaments to highly dense networks of mycelia, filamentous fungi have the ability to grow in various morphological appearances (Papagianni, 2004).

Morphology has a great impact on product formation through the fermentation process. Unique metabolic mechanism was observed in the presence of salt concentration in the halotolerant marine fungi, though the metabolite production rate in the presence of high salt concentration is slow. Production of ortho-quinone obioninene from marine fungi *Leptosphaeria oraemaris* has been observed in varying salt concentration. It was observed that with the increase of seawater concentration fungal growth is also increased and maximum antimicrobial activity appeared in the presence of 25–50% seawater (Masuma et al., 2001). Presence of specific salts in seawater has the impact on the osmotic effect on the cells and are sensitive toward the antibiotic production.

11.4.1.2 Production Using Semi-Synthesis

Full fermentative process is not economically feasible due to the limitation of the complex chemical synthesis; semi-synthesis is an alternative way in product development. For the stereocomplex molecules, the optically pure molecule can be obtained by stereosensitivity of the enzymatic conversions. Lactam antibiotic cephalosporin C, with increased antibacterial activity, was developed from *Acromonium chrysogenum* by semi-synthetic approach (Kück et al., 2014). Biotechnologically produced cephalosporin C is used as the precursor for the synthesis of two-thirds of commercial cephalosporin. Implementation of mutagenesis and genetic engineering improves the yield of cephalosporin C from *A. chrysogenum* in a fed-batch fermentation (Elander, 2003). The chemistry behind the full or partial fermentation process of synthesis and broad biological screening process provides the access to the synthetic analogous for structure–activity relationship (SAR) studies. One example of chemical synthesis of fungal antibiotics is corollosporine (phthalide derivative) from *Corollospora maritima* (Liberra et al., 1998). Apart from organic synthesis, total synthesis of the antibiotic was also reported, where corollosporine from *Aplidium* sp. and pestalone from *Pestalotia* sp. were subjected under total synthesis.

11.4.1.3 Genetic, Metabolic Engineering, and Heterologous System for Production

In the case of the unknown and uncultivable, microorganism's metagenomics and genome mining have been to understand the hidden chemodiversity. Transfection of the DNA from the environment to the host cell may lead to the production of the new product. Use of different molecular techniques is an alternative approach to express the silent gene cluster and manipulate the biosynthetic pathway in order to optimize production process (Xiong et al., 2013). Heterologous host is an important tool for industrial-scale bioreactor production and also helps in transient process for cultivation condition. The multiple regulatory cascade and network are responsible for the antibiotic production, where different genetic approaches to regulate genes can activate antibiotic production. Different strategies have been implemented to

increase the yield of *Saccharomyces cerevisiae* and change its product specificity along with heterologous protein production (Bhadury et al., 2006). The use of metabolic engineering has been reported for the production of novel sesterterpenoid from marine fungi *Fusarium heterosporum* and *Aspergillus versicolor* (Wang et al., 2003). Another example of using metabolic engineering is optimization in fermentation and high-yield strain improvement to obtain 30 g/L cephalosporin C from *A. chrysogenum* in fed batch fermentation (Elander, 2003). Strain improvement through mutagenesis and genetic engineering leads to the homologous cloning of the many genes involved in the biosynthetic pathway of cephalosporin synthesis, and this enables the increase of the flux of production of metabolites from primary to secondary. All the genetic and metabolic engineering are based on the significant knowledge of the omics techniques, where the RNA sequencing reveals the transcription state of a cell along with full genome sequence.

11.4.2 Role of Biotechnology in Downstream Processing

Several steps like separation followed by cell disruption, capture, concentration, extraction, purification, polishing, and formulation are involved in downstream processing (DSP), which is the second main part of the production process. DSP is the most expensive and ineffective part of the bioprocess, which is put down in biotechnological process development. It is mainly used for the commercialization purpose, where a particular batch of antibiotic is being processed before coming to the market. No standard applications of DSP are suitable for all chemical classes. A broad spectrum of downstream tools is used in the biotechnological processes, which can be applied to the marine biotechnology in antibiotic production.

11.4.3 Role of Biotechnology in Lead Development

Lead development is the late stage of discovery pipeline where biotechnology methods also can be used. In lead development, medicinal chemistry has been focused in order to generate products with improved properties like better bioavailability and metabolization with the SAR studies. Mode of action studies during lead development helps in the improvement of the therapeutic application. A small deviation in nature can affect biodiversity more than biotechnology, which was evident by the naturally occurring macrocyclic and linear polyester calcarides from a marine *Calcarisporium* sp. Though *S. epidermidis* and *X. campestris* are inhibited by all macrocyclic calcarides, neither activity was observed by closely related linear polyesters below a MIC of 100 μ M (Silber et al., 2013). The occurrence of different derivatives from marine fungi has enzymatic as well as ecological relevance. Bioconversions of biological derivatives are possible through enzyme treatment in biotechnological processes. For example, laccase-catalyzed amination generates

corollosporine derivative (Mikolasch et al., 2008) and cephalosporin C is enzymatically converted into 7-amino cephalosporanic acid (Barber et al., 2004). For the production of the deacetyl intermediate enzyme, catalyzed de-esterification method is used widely on industrial scale (Table 11.1).

Table 11.1 Different fungi, their antibacterial activity, and biotechnological approaches

| Fungal origin | Approaches in the field of biotechnology | Activity of antibiotics | Class of the compound | References |
|---|---|--|---|-----------------------|
| The fungi are usually obligate in nature living within deep sea | The cultivation is scaled 20–100 L under high pressure High-pressure cultivation scaling 20–100 L | Acts against broad groups of microbial cells | Not determined | Xu et al. (2012) |
| <i>Hypoxylon oceanicum</i> | Various optimized conditions help in the enhancement of titer values | <i>Xanthomonas campestris</i> , <i>Propionibacterium acnes</i> , <i>Staphylococcus epidermidis</i> | Lipodepsipeptide and macrocyclic polyactones | Abbanat et al. (1998) |
| <i>Halichondria panicea</i> | Helps in the increase in yield with decreased time of productivity | <i>P. acnes</i> , <i>S. aureus</i> , <i>S. epidermidis</i> , <i>Septoria tritici</i> , <i>X. campestris</i> | Tetramic acid and ascosetin | Wu et al. (2015) |
| <i>Cladosporium</i> sp. | Transfer to STR from EMF enhances the productivity | <i>Pseudoalteromonas piscicida</i> , <i>Vibrio harveyi</i> , <i>Ruegeria</i> sp., <i>Loktanella hongkongensis</i> , <i>M. luteus</i> , <i>Rhodovulum</i> sp. | Bis(2-ethylhexyl) phthalate, phthalate * | Qi et al. (2008) |
| <i>Ascochyta</i> sp. | Medium optimization is done in small scale | The two-membrane bacterial regulatory system helps in the process of regulation | Spirodioxynaphthalene Ascochytatin | Kanoh et al. (2008) |
| <i>Calcarisporium</i> sp. | Biological derivatization is performed by various types of biological optimizations The yield is dependent on C/N ratio | Macrocyclic compounds: <i>X. campestris</i> <i>S. epidermidis</i> , linear polyesters | Macrocyclic, linear polyesters and calcarides A–E | Silber et al. (2013) |

(continued)

Table 11.1 (continued)

| Fungal origin | Approaches in the field of biotechnology | Activity of antibiotics | Class of the compound | References |
|--|---|---|---|------------------------|
| Unidentified marine biofilm-producing fungi | EMF scaling | It possesses antibiofilm and antibacterial activity against <i>S. aureus</i> , <i>S. haemolyticus</i> , <i>Vibrio</i> sp., <i>Micrococcus</i> sp. | Cyclo-(Pro-Phe), diketopiperazine | Quian et al. (2006) |
| <i>Aspergillus chrysogenum</i> | The genetically engineered organisms helps in reducing the formation of the by-products | They act against broad groups of microbial species | β-Lactam cephalosporin, | Posch et al. (2013) |
| <i>Cephalosporium chrysogenum</i> | Mutagenesis-associated modification of DNA | They act against broad groups of microbial species | | Kensy et al. (2009) |
| <i>Ampelomyces</i> sp. | Scaling in EMF | <i>Vibrio</i> sp., <i>S. aureus</i> , <i>Pseudoalteromonas</i> sp., <i>Micrococcus</i> sp. | 3-Chloro-2,5-dihydroxy benzyl alcohol, benzene derivative | Samorski et al. (2005) |
| <i>Porteresia coarctata</i> , <i>Penicillium chrysogenum</i> | Enhancement of yield take place by alteration of carbon and nitrogen source | <i>Vibrio cholera</i> | Diketopiperazine Chrysogenazine | Posch et al. (2013) |
| <i>Halosarpheia</i> sp. | Reprogramming of the biosynthetic pathway | <i>S. aureus</i> , <i>Yersinia enterocolitica</i> , <i>Clostridium perfringens</i> , <i>P. aeruginosa</i> , <i>E. coli</i> , <i>Salmonella enterica</i> , <i>Shigella dysenteriae</i> | Cyclodepsipeptides, Enniatins | Duetz et al. (2000) |
| <i>Corollospora maritima</i> | Salt-dependent enzyme-associated treatment of biological derivatives | <i>S. epidermidis</i> , <i>Candida maltosa</i> , <i>Bacillus subtilis</i> , <i>Escherichia coli</i> , <i>P. aeruginosa</i> , <i>S. aureus</i> | Phthalide derivatives and Corollosporine derivatives | Bills et al. (2008) |
| <i>Phakellia fusca</i> , <i>Aspergillus terreus</i> | Agitation, inoculation, rate of aeration, pH, speed of agitation, and nutrient feeding need to be optimized | <i>B. subtilis</i> | (+)-Terrein, cyclopentenone | Wang et al. (2003) |

(continued)

Table 11.1 (continued)

| Fungal origin | Approaches in the field of biotechnology | Activity of antibiotics | Class of the compound | References |
|---|---|--|--|--------------------------|
| <i>Mycale adhaerens</i> , <i>Exophiala pisciphila</i> | Transfer from EMF to STR | <i>S. aureus</i> , <i>E. faecalis</i> , <i>E. facium</i> | 3,5-Dihydroxy-decanoic polyester, Exophilin A | Xiong et al. (2009) |
| <i>Halichondria panicea</i> | Adaptation of medium Transfer from EMF to STR enhances the yield | <i>S. epidermidis</i> , <i>P. acnes</i> , <i>S. tritici</i> , <i>S. aureus</i> , <i>X. campestris</i> | Tetramic acid lindgomycin, | Mikolasch et al. (2008) |
| <i>Aspergillus versicolor</i> , <i>Fusarium heterosporum</i> | Metabolic engineering | They act against broad groups of microbial species | Sesterterpenoid | Wang et al. (2003) |
| <i>Leptosphaeria oraemaris</i> | Salinity affects the production of antibiotics | – | Ortho-quinone, obioninene | Miller and Savard (1989) |
| <i>Arthrinium saccharicola</i> | Systematic manipulation of culture conditions | <i>Vibrio vulnificus</i> , <i>Pseudoalteromonas spongiae</i> | Not determined | Miao et al. (2006) |

11.5 Conclusion

The use of microbial species in human need has been observed for more than 100 years. Fungi proved to be one of the most beneficial microbial organisms due to a large content of biomass being associated with them. They are the storehouse of large numbers of natural compounds that have acted efficiently for curing various types of diseases. Thus, the compounds associated with fungi can be an alternative to the conventional drugs after detailed investigations.

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Chapter 12

Optimizing Physical Parameters for Amylase Production Using *Aspergillus niger* and Ammonium Molasses Medium



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12.1 Introduction

Amylases are a class of hydrolytic enzymes that converts starch to sugar like, maltose, etc. α -Amylases (EC 3.2.1.1) are starch-degrading enzymes capable of hydrolyzing internal α -1,4-glycosidic linkages in polysaccharide starch, resulting in the production of short-chain dextrin or low molecular weight carbohydrates (Sindhu et al., 2017; de Souza & de Oliveira Magalhães, 2010). Global market for industrial

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enzymes was estimated at \$4.2 billion in 2014 and expected to develop at a compound annual growth rate (CAGR) of approximately 7% over the period from 2015 to 2020 (Singh et al., 2016). α -Amylase covers about 30% shares of the total enzyme market (Paul et al., 2021). Amylase enzymes are used in diverse industrial sectors like food, pharmaceuticals, textile, and detergents. In fermentation-based industries like alcohol, etc., amylase is used in breaking starchy substrates to fermentable sugars. Growing demand for processed food, increasing demand for bio-ethanol due to rising environmental concerns, new technological innovations like dish washer, which require dish-washing detergents, and growing research and developmental activities are some of the factors driving the growth of the amylase market share.

Amylase enzyme is produced by plants, animals, and microorganisms. However, enzymes from fungal and bacterial sources have dominated applications in industrial sectors (Rana et al., 2013). Microorganisms are favored sources for industrial enzymes due to easy availability and fast growth rate. Genetic changes using recombinant DNA technology can easily be done on microbial cells for elevated enzyme production and scientific development (Illanes et al., 2012). Microbial sources of amylase are also preferred due to cost-effectiveness, consistency, and ease of process modification. Bacterial amylase is mostly obtained from *Bacillus* genus such as *B. stearothermophilus*, *B. subtilis*, *B. cereus*, *B. licheniformis*, and *B. amyloliquefaciens* that are isolated and screened for amylase production (Sivaramakrishnan et al., 2006, as cited in Elmansy et al., 2018). Among the fungi, *Aspergillus niger*, *Aspergillus oryzae*, *Thermomyces lanuginosus*, *Penicillium expansum*, and many species of *Mucor* are capable of producing amylase enzyme (El-Fallal et al., 2012).

For amylase production, different methods like solid-state fermentation (SSF) or submerged fermentation can be used. Solid-state fermentation process occurring in the absence or near-absence of free water has various advantages, such as low energy requirement, high product concentration, and little wastewater production, and is environmentally friendly compared with submerged fermentation. But a major disadvantage of SSF is thermal conduction or heat removal. Submerged fermentation (SMF) technology has the advantages of short period, low cost, high yield, easier purification of products, and easier control of the process.

Thus, the growing demand for amylase enzyme and the possibility of using cheap substrates make production of amylase a potential avenue for entrepreneurship. For economic competitiveness, it becomes important to standardize optimum parameters like media composition, temperature, pH, and time of incubation. Although there are many reports of optimum parameters for high yield, there is no consensus over the optimum parameters. So, this study attempts to determine the optimum range of pH and temperature that can help fungal amylase production entrepreneurship.

12.2 Materials and Methods

12.2.1 Isolation of the Microorganism

Aspergillus niger, used for amylase production, was isolated from soil sample by the method of serial dilution and plating on PDA media. Black fungal colonies were observed under microscope using cotton blue lactophenol stain and identified using handbook and pure cultured by streaking.

12.2.2 Screening

Screening of amylase activity by the fungi was done using starch agar medium. Starch agar medium was prepared using starch, 20 g; KNO₃, 1 g; K₂HPO₄, 0.5 g; MgSO₄·7H₂O, 0.5 g; NaCl, 0.5 g; FeSO₄·7H₂O, 0.01 g; agar, 15 g; water to 1 L and sterilized (Shirling & Gottlieb, 1966). The solidified starch agar plates were inoculated with a loop full of *Aspergillus niger* pure culture and incubated at 30 °C for 48 h. Iodine solution was poured over the plates. A colorless zone surrounding the fungal colony indicated conversion of starch to maltose by the fungal amylase.

12.2.3 Inoculation

Aspergillus niger from pure culture was streaked out and mixed in 5.0 ml of sterile deionized water. Then, from the spore suspension, 1.0 ml was inoculated to fresh sterilized potato dextrose broth medium for development of working culture and incubated at 30 ± 1 °C up to 7 days to attain about 5.0 × 10⁸ spores ml⁻¹. For the entire study, 1 × 10⁷ spores ml⁻¹ was used as inoculum to carry out the fermentation process (Sethi et al., 2013). The sterile media was allowed to cool and inoculated with spore suspension inside laminar air flow chamber and incubated according to experimental design.

12.2.4 Medium

Composition of medium used for microbial fermentation is very critical because it influences the output of the process. Various carbon sources like molasses, whey, grains, and agricultural or industrial wastes are used with the objective of cutting down cost of production, and also addresses the environmental issue of waste management. Cane molasses are often used in industrial microbiology as carbon source. In cane molasses, sugar content is diluted to about 25% sugar level by adding

double-distilled water. The molasses solution, after adding 35 ml of 1N H₂SO₄ per liter, is boiled for half an hour, cooled, neutralized with lime water (CaO), and left to stand overnight for clarification (Panda et al., 1984). Neutralized molasses solution was treated with 2% (w/v) tricalcium phosphate (TCPH followed by autoclaving at 105 °C for 5 min. The mixture is cooled and centrifuged at 3000 rpm for 15 min. The insoluble matter is discarded. TCPH treatment reduces considerably the metal loads of molasses compared to untreated molasses. (Kundu et al., 1984). Nitrogen is another important structural and functional component of all living organisms and plays a very important role in economic feasibility of any industrial fermentation process. Different organic nitrogen sources like peptone, tryptone, gelatin, yeast extract, beef extract, and inorganic nitrogen sources like ammonium nitrate, ammonium sulfate, ammonium chloride, and sodium nitrate are frequently used in microbial media. Although very good yield of organic acids, amylase, etc. has been reported by using organic nitrogen sources, because of their high cost, inorganic nitrogen sources like ammonium sulfate or ammonium nitrate are preferred for industrial production to lower the cost of production. The inorganic source ammonium nitrate (0.1%) was found suitable to produce high amount of enzyme and showed increased enzymatic activity by *Bacillus subtilis* (Deb et al., 2013). Amylase production by *Aspergillus* has been found to be higher using ammonium sulfate as an inorganic nitrogen source as compared to ammonium nitrate (Oshoma et al., 2010).

12.2.5 Extraction

Extraction of the crude enzyme was done by centrifugation of the fermented media at 2000 rpm (revolution per minute) for 5 min, supernatant collected and filtered off using Whatman No.1 filter paper. The filtrate was used as crude enzyme extract (Oyeleke et al., 2010)

12.2.6 Influence of Initial pH, Temperature, and Incubation Time

Amylase activity was checked at different initial pH values ranges 5.0–11.0 and temperature range 30–60 °C. 250 mL flask containing 50 mL of enzyme production medium was inoculated with 10 (% v/v) inoculum from 24-h-old seeds culture and was incubated in shaking incubator at 37 °C for 60 h, while, in case of temperature optimization, pH was adjusted to 7.0 and other conditions remained unchanged. Amylase activity was determined from culture broth obtained after centrifugation (12,100 for 10 min at 10 °C). To observe the optimum incubation time, the inoculated media with initial pH 7 were incubated at 37 °C for (24–96) h.

12.2.7 Enzyme Assay

The α -amylase activity was assayed by 3,5-dinitrosalicylic acid (DNS) procedure using 1% soluble starch (Bernfeld, 1955). Dinitro salicylic acid required for enzyme assay is prepared by dissolving 1 g of DNS in 50 ml of distilled water. To this solution, about 30 g of sodium potassium tartarate tetrahydrate is added in small lots, and the solution turns milky yellow in color. To this, 20 ml of 2N NaOH is added, which turns the solution to transparent orange yellow color. The final volume is made to 100 ml with the distilled water and stored in an amber-colored bottle.

To prepare a standard graph of maltose, a stock solution is prepared using 50 mg of maltose in 50 ml of water. From the stock solution, 10 ml of solution is dissolved with 40 ml of distilled water to prepare the working solution. From the working solution, different amounts, that is., 0.2, 0.4, 0.6, 0.8, and 1 ml, are taken in five test tubes, and one test tube is kept blank. The final volume of all the test tubes is made 1 ml by adding distilled water, and then 1 ml of DNS is added to each test tube. The contents in the test tubes are heated in a boiling water bath for 5 minutes. The test tubes are cooled and 10 ml of distilled water is added to each test tube. Optical density of each test tube is determined by a spectrophotometer at 540 nm (Table 12.1). A graph is plotted with the amount of maltose on X-axis vs. OD at 540 nm on Y-axis (Fig. 12.1).

For estimating enzyme activity, 1% starch solution is prepared by using 1 gram of starch in 100 ml of distilled water and 1 ml of enzyme extract maintained at different conditions under experimental design. In one of the test tube, to be treated as blank, 1 ml of distilled water is added instead of enzyme. All the test tubes are incubated for 10 minutes at 30 °C. Then, 1 ml of DNS was added and incubated in boiling water bath for 5 min. The test tubes are then cooled and 10 ml of distilled water is added and mixed well. The optical density of the sample is then determined against blank at 540 nanometers. All assays were performed in duplicate and each reading was taken three times (Chakraborty et al., 2018).

Table 12.1 Spectrophotometric absorbance shown by maltose at different concentrations

| Maltose in micrograms | Average optical density at 540 nm |
|-----------------------|-----------------------------------|
| 0 | 0 |
| 40 | 0.039 |
| 80 | 0.119 |
| 120 | 0.138 |
| 160 | 0.194 |
| 200 | 0.226 |

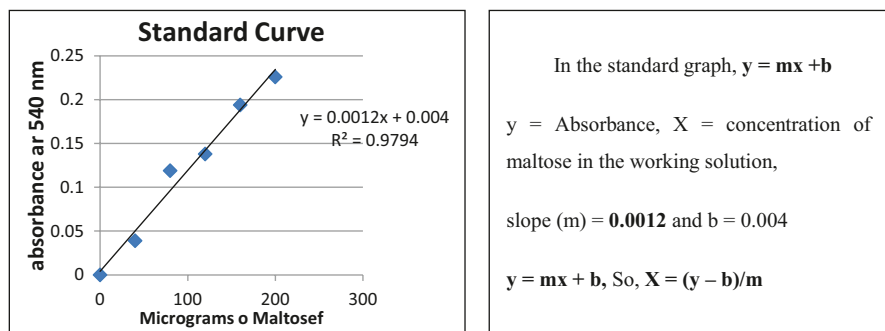


Fig. 12.1 Standard maltose curve

Table 12.2 Effect of pH on the production of amylase by *Aspergillus niger*

| Effect of initial pH | | |
|----------------------|------------|--|
| pH | Absorbance | μg of maltose units per ml of enzymes |
| 4.5 | 0.189 | 154.16 |
| 5 | 0.527 | 435.83 |
| 5.5 | 0.766 | 635.00 |
| 6 | 0.829 | 687.50 |
| 6.5 | 1.14 | 946.67 |
| 7 | 0.794 | 658.33 |
| 8 | 0.686 | 568.33 |

12.3 Results

12.3.1 Screening

Starch agar medium inoculated with *Aspergillus niger* when flooded with iodine, after incubation, showed a clear zone, indicating amylase activity on starch.

12.3.2 Effect of Initial pH

The effect of pH on amylase production is shown in Table 12.2. Of the range of pH 4.5–8, maximum enzyme activity per ml of enzyme was observed at pH 6.5. The amount of maltose produced by the enzyme was found to be 946.67 $\mu\text{g}/\text{ml}$ of enzyme extract. Enzyme activity is seen to rise gradually from pH 4.5 to 6.5 and again declines at pH 7 and 8 (Fig. 12.2).

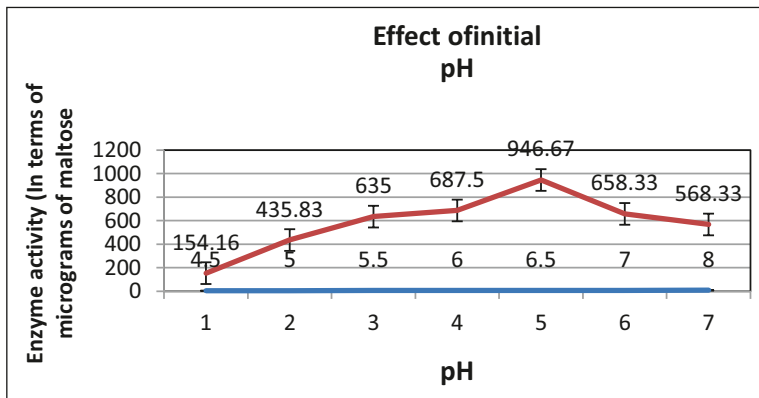


Fig. 12.2 The amount of maltose released by amylase increases from pH 4 to 6.5 and then starts declining

Table 12.3 Effect of initial temperature on the production of amylase by *Aspergillus niger*

| Effect of initial temperature | | |
|-------------------------------|------------|---------------------|
| Temperature (°C) | Absorbance | µg of maltose units |
| 25 | 0.522 | 431.67 |
| 30 | 0.732 | 606.67 |
| 35 | 1.984 | 1650 |
| 40 | 2.173 | 1807.5 |
| 45 | 1.276 | 1506250 |

12.3.3 Effect of Initial Temperature

Table 12.3 shows the effect of temperature of incubation on amylase production. Amylase production observed in terms of amylase activity on starch releasing maltose was found to be 431.67 µg/ml of enzyme extract when incubated at 25 °C and increased to maximum of 1807.5 µg/ml of enzyme extract at 40 °C. Further rise of temperature to 45 °C decreased enzyme activity 1060 µg/ml of enzyme extract (Fig. 12.3).

12.3.4 Effect of Incubation Period

The effect of incubation time on amylase production is shown in Table 12.4. It is seen that 24 h after inoculation, amylase activity in terms of maltose produced is 58.33 µg/ml of enzyme extract. The production increased to a maximum of 126.33 µg/ml of enzyme extract after 96 h of incubation and started declining with further incubation (Fig. 12.4).

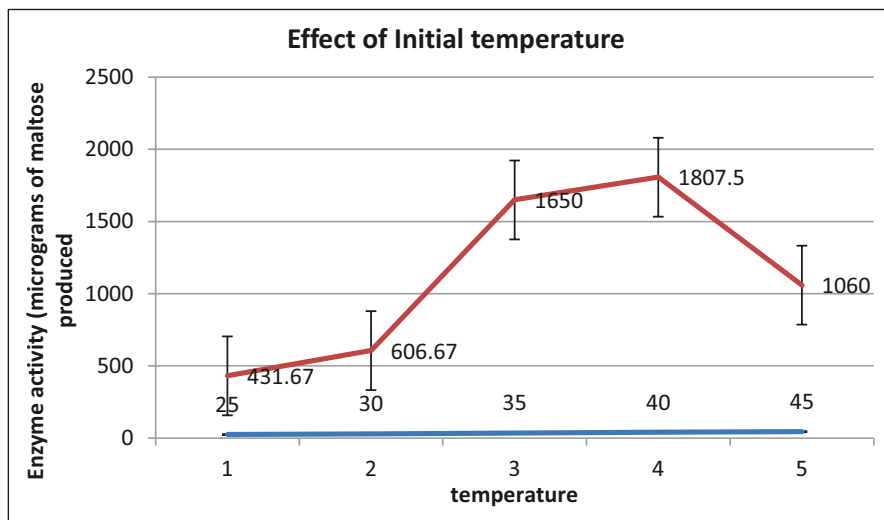


Fig. 12.3 Maltose concentration increases with temperature till 40 °C and then starts declining

Table 12.4 Effect of incubation time on the production of amylase by *Aspergillus niger*

| Incubation time (h) | Absorbance | µg of maltose units |
|---------------------|------------|---------------------|
| 24 | 0.074 | 58.33 |
| 48 | 0.092 | 73.330 |
| 72 | 0.098 | 78.33 |
| 96 | 1.52 | 126.33 |
| 120 | 1.39 | 115.5 |
| 144 | 1.08 | 89.67 |

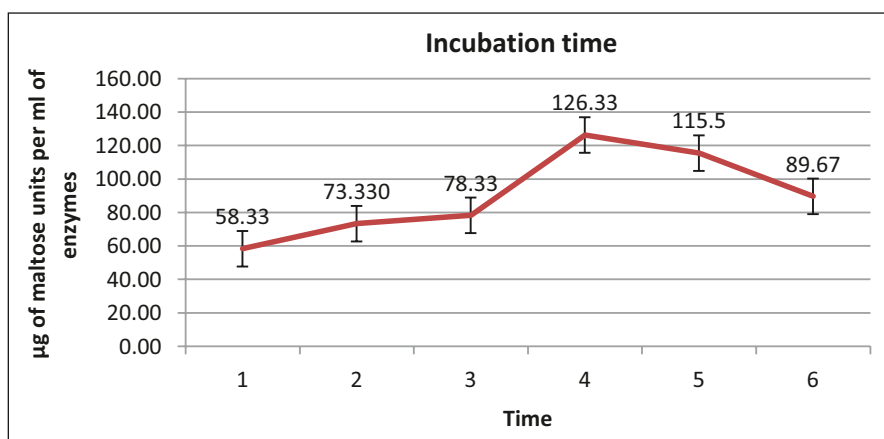


Fig. 12.4 The amount of maltose produced by maltose increases from 1-day incubation till 4 days and declines with further incubation

12.4 Discussion

Growing demands for amylase enzymes by different carbohydrate-based industries have opened up opportunities for entrepreneurship for setting small- and medium-scale industries for producing amylase enzyme at low-input cost. Startups using microbes and fermentation include MycoTechnology, which turns pea and rice protein fermented with shiitake mycelium, or fungi, into food ingredients, and Cargill produces fish food from naturally occurring microbes that use methane as an energy source (Financial Time, 2019).

To be economically competent, process optimization is very important. Effect of various physical parameters on α -amylase production was investigated in this study. pH is a very important parameter that affects any biochemical process by affecting the active sites of the enzymes involved in the process. The pH optima for production of amylase by submerged fermentation has mostly been reported as slightly acidic to neutral. In this study, the optimum pH for amylase production by *A. niger* under submerged foundation was found to be 6.5. According to other studies, the optimum pH was found to vary.

Optimum pH for amylase production was found to be 6 in various studies (Behailu & Abebe, 2018; Wang et al., 2016; Singh et al., 2014; Ahmed et al., 2020). Monga et al. (2011) reported 4.6 pH as optima for amylase production.

Temperature of incubation for amylase production has been found to affect amylase production significantly. In this study, the temperature optima for α -amylase production by *A. niger* was found to be 40 °C. In different studies, a range of optimum values of incubation temperature for amylase production has been reported. Ahmed et al. (2020) reported 30 °C, while Sethi et al. (2016) reported range of 27–36 °C and 45 °C (Behailu & Abebe, 2018).

It has been established in growing volume of studies that production of enzyme and also metabolites vary at different points of incubation. In this study, maximum amylase production observed in terms of amylase activity was observed after 96 h of incubation similar to the findings of Sethi et al., 2016. Optimum incubation time reported in other studies was 72 h (Behailu & Abebe, 2018), 144 h (Singh et al. 2014), and 120 h (Ahmed et al., 2020; Monga et al., 2011). Although most of the studies have reported the optimum parameters in a close range, but there is a need for further studies on specific conditions to optimize the process.

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Chapter 13

Plant–Fungal Interactions



K. Geetha and Vasavi Dathar

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13.1 Introduction

The most intense interactions between microbes and plants take place at the rhizosphere, which is the interface between plant roots and the soil. Soil microbes have a tremendous influence on plant health and productivity (Bloemberg & Lugtenberg, 2001). Interactions between plants and fungi are influenced by cultivars, mutants, and nonhost species. Different stages in the colonization processes may act as control points in different interactions.

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Plant–fungal interactions are of fundamental importance in crop productivity, which may be symbiotic or pathogenic. Plants produce antimicrobial compounds like phytoanticipins and phytoalexins, which act as phyto protectants. Some pathogenic fungi have evolved hydrolytic enzymes that can degrade toxic phyto protectants, thereby allowing the fungus to overcome specific chemical barriers (Tudzynski & Sharon, 2003). Fungi are versatile with respect to their nutritional requirements. Because of this versatility, they interact with plants in different ways. Thus, they play a major role in the natural ecosystems and also in the present agricultural scenario. They decompose and recycle organic materials by interacting with the plants both underground and above ground, that is, with the root and shoot system. They interact with plant roots in the rhizosphere and also above ground and decompose and recycle organic materials. The complex interactions between the fungus and plants lead to diverse outcomes. Several fungi have a combined mode of lifestyles like saprophytic, pathogenic, or symbiotic (Grigoriev, 2013). Techniques like transcriptomics and the omics era have provided an insight into the mechanism of plant–mycorrhizal interactions.

Symbiotic fungi colonize the plants and are beneficial to their hosts by promoting growth, producing secondary metabolites, and enhancing resistance to biotic and abiotic stresses. The symbiotic associations between plants and fungi include endophytic and different types of mycorrhizae. The associations between plants and endophytic fungi are governed by molecular mechanisms and deterministic factors that enhance plant growth, thus providing protection against invading pathogens. In contrast, large numbers of fungi are pathogens that constitute a major threat to crop yield and food security (Shah, 2018).

13.2 Genomics of Plant–Fungal Interactions

Understanding the molecular mechanisms by which fungal pathogens cause disease, subvert plant immune systems, and divert plant developmental and metabolic processes is crucial for development of new antifungal strategies. Identification of key virulence factors of the fungus that allow the pathogen to cause disease or to escape detection by plant's immune system is important. Also, it is important to know the mechanisms by which plants recognize and respond to fungal pathogens. Several fungal and plant genomes are being sequenced and assembled, so that the impact of functional genomics enables us to know how the plant's immune system is able to discriminate between the beneficial and harmful fungi. With the continuous progress being made toward the understanding of plant–fungal interactions, new strategies for crop protection are emerging at an increasingly rapid pace (Shah, 2018).

In nature, plants are generally resistant to many pathogenic fungi and are able to combat infections caused by the pathogenic fungi. Among the plant–fungal associations, symbiotic and neutral associations dominate, whereas parasitic associations are considered to be an exception (Staskawicz, 2001). The plant attracts fungi in the

rhizosphere by producing different secretions as a result of its metabolic activity. These secretions are determined by the genetic nature of the plant.

The outcome of the plant–fungal interaction is driven by the receptors and defense proteins expressed by the plant and the specific molecules produced by the fungi. Mutations in the genes of the plant or fungus may cause a change in these interactions where a resistant plant becomes sensitive or susceptible to the fungal infection or vice versa (Stracke et al., 2002; Giraldo & Valent, 2013). Microbes beneficial to plants have evolved strategies that suppress the defenses exhibited by the plants they infect, which allow them to colonize the plants either as epiphytes or endophytes (Zamioudis & Pieterse, 2012). Both symbiotic and pathogenic fungi use mostly similar colonization patterns like development of feeding structure and nutrient sequestration by establishing obligate relationship with the plants (Corradi & Bonfante, 2012). The outcomes of these interactions are contrasting as some plants are rewarded with symbiosis and some others suffer parasitism. In an evolutionary perspective, some fungi have evolved as parasites or pathogens, and, on the other hand, certain fungi have evolved as successful symbionts by their interaction with the host plants.

By decoding the functions of the receptors and signals of both the fungus and plant, we can understand their complex interactions and also their role in inducing immunity in the host plant. Successful colonization of the fungus in the host plant requires all of these contributing factors that also influence the resistance of the plant toward the fungal partner.

13.3 Mycorrhizal Association

Fungi are filamentous in nature and exploit diverse substrates depending on their nutritional strategy. The saprophytic fungi, called as saprobes, thrive in soil, water, and on dead and decaying plant and animal tissues. The other group of fungi, which are the parasitic and mutualistic symbionts, feed on living organisms (Carlile et al., 2001). The mycorrhizal fungi form a heterogeneous group of species that are spread over diverse fungal taxa. The word “mycorrhiza” was introduced by Frank in 1885, which means “fungus roots” (Frank, 1885; Trappe, 2005). Mycorrhiza is the symbiotic association between the roots of a higher plant and a fungus. These fungi can be divided into two major groups: aseptate endophytes such as Glomeromycota or septate Ascomycota and Basidiomycota.

More commonly, mycorrhizal classifications include two broad categories that reflect anatomical aspects (Smith & Read, 2008), referred to as ectomycorrhizae (EMs) where the fungus colonizes intercellular spaces, and endomycorrhizae, in which the fungus develops inside the cells and colonizes the roots. A third category of mycorrhizae is the ectoendomycorrhizae with mixed behavior between ectomycorrhiza and endomycorrhizae (Fig. 13.1). The endomycorrhizae are of three types: orchid, ericoid, and arbuscular mycorrhizae (AMs). The mycorrhizal fungi can be free living, which is a part of their life cycle, and also as symbionts colonizing 90%

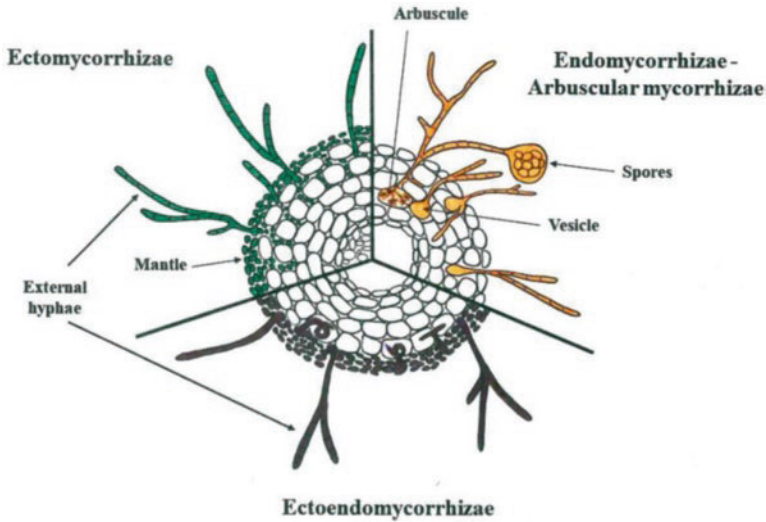


Fig. 13.1 Different types of mycorrhizae. (Courtesy: Ganugi et al., 2019)

of plant species including crop plants, forest trees, and wild grasses. Both the plant and fungi benefit from the relationship where the mycorrhizae improve the nutrient status of the host plant that influences the mineral nutrition, water absorption, growth, and disease resistance. In turn, the host plants help in the growth and reproduction of the fungus (Bonfante & Genre, 2010).

Many land plants have symbiotic association with soil-borne mycorrhizal fungi that provide nutritional benefits to both the partners and make them fit. These benefits are important in solving the food security challenge. Mycorrhizal fungi are found in the environments such as alpine and boreal zones, tropical forests, grasslands, and croplands. The belowground arbuscular mycorrhizal fungal (AMF) communities are affected by biotic factors including animals, plants, fungi, bacteria, and protists, and abiotic factors like water, soil, air, sunlight, temperature, and minerals (Larissa et al., 2019). Mycorrhizal fungi absorb and supply soil nutrients to the plant by the specific activity of the mycelia and thus play a major role in nutrient cycling, but their role in carbon flux is not much defined (Selosse & Roy, 2009).

Mycorrhizae connect plant communities by developing a wide network of the hyphae inside the soil, thus offering a horizontal transfer of nutrients between the plant and fungus (Helgason et al., 1998). Mycorrhizae form special spaces to interact with the host plant through symbiosis (Bonfante, 2001; Harrison, 2005; Parniske, 2008). The ectomycorrhizae form a thick mantle with their hyphae around the epidermal cells of the root tip developing into a Hartig net. The root tip is not colonized by arbuscular mycorrhizae. Hyphae developing from the spore produce a hyphopodium on the root epidermis. Colonization proceeds both intra- and intercellularly, leading to the formation of arbuscules inside inner cortical cells (Fig. 13.1).

13.3.1 Mechanism of Colonization

The study of plant responses illustrates how the mechanisms operate to accommodate the AM fungus inside the plant cell lumen by diverse plants and conserved during evolution. Root colonization is vital to AM fungi (Fig. 13.2); their spores feed germinating hyphae through the catabolism of storage lipids for just a few days (Bonfante & Genre, 2010). The hyphae repeatedly try to explore the soil searching for a host, and when it does not find one, it will arrest its growth and retract the cytoplasm back into the spore. When it finds a host, it re-germinates. Due to the wide host range of the mycorrhizal fungi, this situation is not very frequent in nature, but it shows their endurance because of their strict biotrophy. With the help of root organ cultures, production of nucleic acid extractions on a large scale has been made possible for growing arbuscular mycorrhizal fungi (Becard & Fortin, 1988).

13.3.2 Formation of Hyphopodium

Once a chemical acquaintance has been made between the fungus and the plant, roots and hyphae proliferate and branch in a small volume of the rhizosphere. This pre-symbiotic phase of the AM interaction culminates in a physical encounter between symbionts, when a hyphal tip touches the surface of a root (Fig. 13.2). These fungi start root penetration after carefully selecting a location, and hyphae

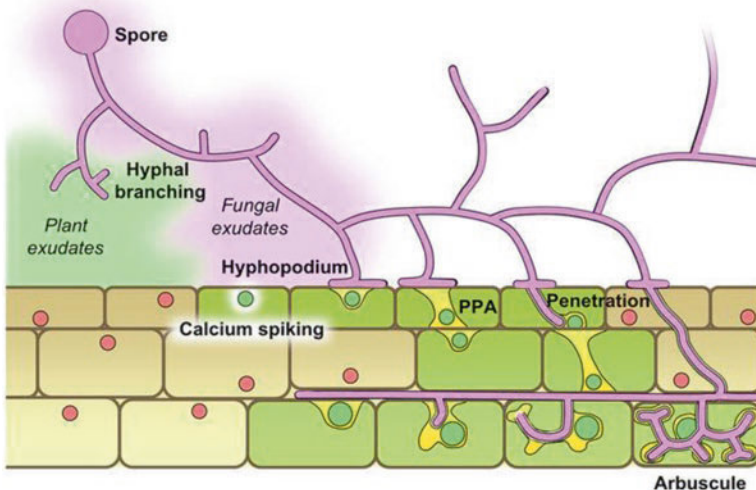


Fig. 13.2 Schematic summary of the root colonization process by AM fungi. (Courtesy: Bonfante & Genre, 2010)

wander along the root surface for several centimeters, forming a long, straight, or slightly curved structure.

This structure swells and flattens on some of the epidermal cell walls and repeatedly branch to form a structure called as hyphopodium (Genre et al., 2005). The arbuscular mycorrhizal fungi concentrate in the young lateral roots of the plants (Kosuta et al., 2003), which are the primary site of colonization of these fungi under laboratory conditions (Chabaud et al., 2002). The hyphopodial wall penetrates into the epidermal cells of the root with many protuberances. But the development of hyphopodium is followed by a pause in fungal growth for about 4–6 hours. Then, the growth of a new tip is initiated and the penetration hypha develops (Genre et al., 2005). The plant cells and tissues now prepare for colonization.

13.3.3 *Gene Expression*

The gene coding for secretion of a protein responsible for expression of symbiotic pathway, ENOD11, has been reported in the hyphopodium (Chabaud et al., 2002). Many genes of the plant are regulated before the symbiotic association and their expression also changes (Weidmann et al., 2004). During the formation of pre-penetration apparatus, the cell wall remodeling genes and the defense genes become active (Siciliano et al., 2007). The pre-penetration apparatus that is specific to arbuscular mycorrhizal fungi is required for the fungal penetration into the plant (Genre et al., 2005). The endoplasmic reticulum, Golgi bodies, and secretory vesicles are synthesized abundantly, which are concentrated in the pre-penetration apparatus (Genre et al., 2008).

After the completion of the pre-penetration apparatus, the fungus starts growing again. The hyphal tip heads through the epidermal cell wall along the track of PPA. The perifungal membrane may assemble and PPA secretory vesicles may fuse and produce an invagination of the plasma membrane. This marks the formation of a symbiotic interface, and the intracellular compartment allows the growth of arbuscular mycorrhizal fungi without disrupting the plant cell (Bonfante, 2001) (Fig. 13.2).

Fungi evolved much earlier than the plants, and plant–fungal interactions are thought to be started during the evolutionary period of the terrestrial vascular plants (Humphreys et al., 2010; Field et al., 2012). The fungal partners might have helped in the colonization of the plants on the land (Redecker et al., 2000). These associations might have started approximately 400–460 million years ago at the time of evolution of the vascular plants (Remy et al., 1994; Kemen & Jones, 2012). The stability of both the fungus and the plant is provided by beneficial fungi, whereas the pathogenic fungi cause destabilization of the host plant (Jones & Dangl, 2006).

13.3.4 Plant Growth and Development

The symbiotic association between the fungi and plants promotes the growth and development of the plant partner. Here, the fungus improves foraging of the plant, helps to acquire nutrients and water from the soil, and makes the plant in stress tolerant. The plants in turn deliver nutrition to the fungus in the form of carbohydrates (Buscot et al., 2000). All these factors contribute to a stable interaction between the two partners (Fig. 13.3).

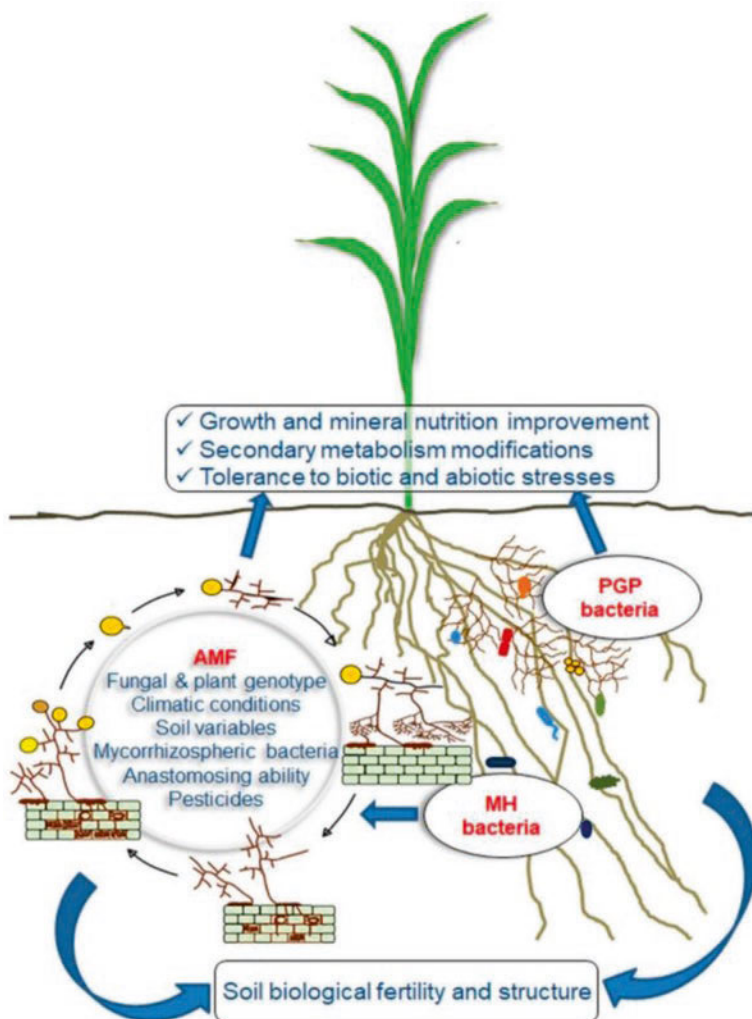


Fig. 13.3 Impact of arbuscular mycorrhizae on plant growth. (Courtesy: Giovannini et al., 2020)

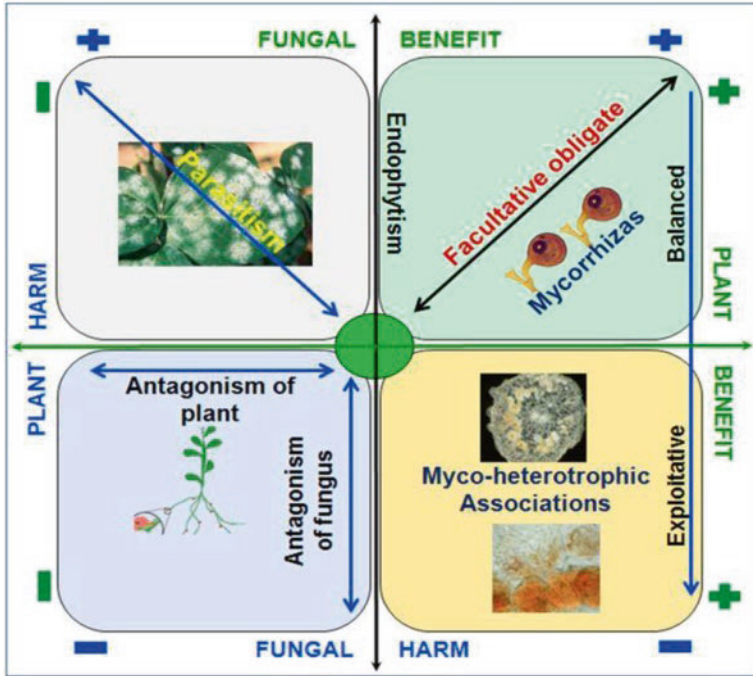


Fig. 13.4 Beneficial and harmful interaction between plant and fungus. (Courtesy: Zeilinger et al., 2016)

The endophytic fungi grow inside the living tissues of plant like the leaves, stems, and roots until the host plant attains senescence. At this stage, the endophyte may become slightly pathogenic (Brundrett, 2004). Because of the abundance of the inhabitant endophytes, most of the plants harbor more than one fungal species and become more resistant to herbivores and pathogens and also become stress tolerant (Strobel & Daisy, 2003). A comparison of different plant–fungal interactions is depicted in Fig. 13.4.

13.4 Fungi as Biofertilizers

In addition to enhancing the nutrient supply to plants, microbes also confer a degree of protection against plant diseases. Various fungi, especially of the genera AMF and *Trichoderma*, produce a range of metabolites against other phytopathogenic fungi (Walsh et al., 2001; Raaijmakers et al., 2002). With development, such microbes could become a realistic alternative to the heavy fungicide regimen used in agriculture at present. A reduction in the use of these chemicals would lead to obvious environmental benefits. The introduction of *Trichoderma* and arbuscular

mycorrhizal fungi (AMF) is known to increase the growth of many plant species, crop plants like *Vigna radiata*, *Saccharum officinarum*, *Zea mays*, *Lycopersicon esculentum*, *Cucumis sativus*, *Camellia sinensis*, *Pennisetum glaucum*, *Cicer arietinum*, *Raphanus sativus*, *Sorghum bicolor*, *Ipomoea batatas*, *Arachis hypogaea*, *Oryza sativa*, *Catharanthus roseus*, *Mentha piperita*, *Coleus forskohlii*, *Andrographis paniculata*, etc. (Selvaraj et al., 2008). Understanding the interaction between consortium of microbial inoculants and plant systems will pave the way to harness more benefits through microbial inoculants for improving plant growth and yield (Raja et al., 2006).

13.5 Disease Production

In contrast to the beneficial effects shown by the symbiotic fungi, the plant–pathogen interaction causes a sequence of events, resulting in the development and maintenance of disease cycle in the host (Daly, 1984). The disease cycle is a chain of interconnected successive events of a pathogen’s infection in a host plant. It usually coincides with the lifecycle of the pathogen with a correlation to its host and the environment (Shomrat, 2021).

A disease cycle (Fig. 13.5) in the plants consists of the following phases:

1. The fungi come into contact with the susceptible plant by wind, water, or insect bite or enter through the roots (Travadon et al., 2012).
2. Germination of fungal spores and attachment of the pathogen to the host receptors (Tucker & Talbot, 2001).
3. Penetration of the pathogen into the plant through wounds or natural openings such as stomata or direct penetration with the help of fungal appressoria (Pryce-Jones et al., 1999) or through wounds caused by insects such as *Grosmannia clavigera* on lodgepole pines (Diguistini et al., 2011) and *Ophiostoma ulmi* on Dutch elm (D’Arcy, 2000).
4. Establishment of the pathogen by invading into the plant tissues, spread from cell-to-cell causing visible symptoms.
5. Multiplication of the pathogen by reproduction in the host tissues and production of large number of spores.
6. Dispersal of the spores through wind or insects to other susceptible hosts.
7. Survival of the pathogen in a dormant stage under unfavorable conditions (Brown & Ogle, 1997).

Different species evolve together, and this adaptation results in a variety of relationships, such as mutualistic, endosymbiotic, parasitic, competitive, and pathogenic or antagonistic (Faust & Raes, 2012). Many secondary metabolites that are usually bioactive were reported to perform important functions in the fungal and interactions in the ecosystem (Braga et al., 2016) (Table 13.1).

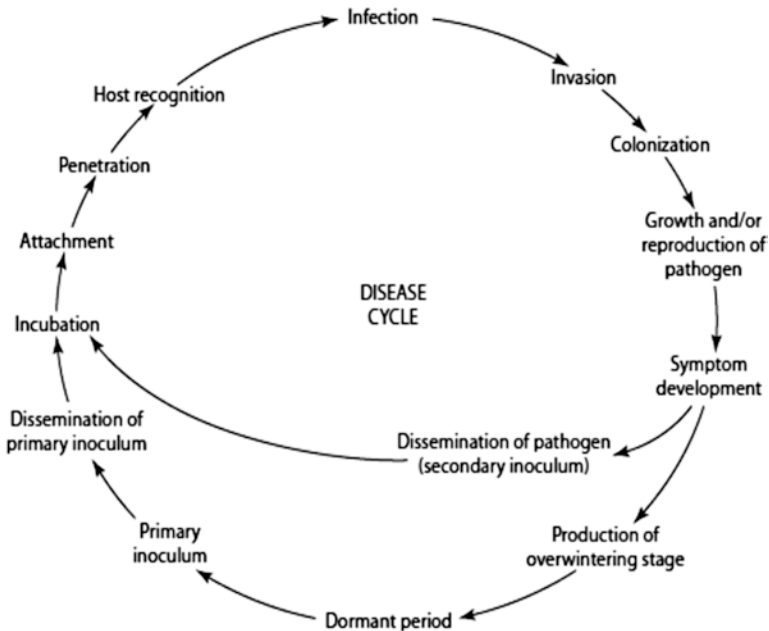


Fig. 13.5 Disease cycle. (Courtesy: Shomrat, 2021)

13.6 Conclusion

Mycorrhizal fungi are important in the fields of environmental change, ecosystem conservation, sustainable agriculture, and development of plants for future needs and food safety. These fungi mobilize P and N, and are an important C sink in the soil, having therefore an important impact on the cycling of these elements (Klironomos et al., 2005). As biofertilizers, they may counteract fertilization excess and thus promote sustainable agriculture. The selection of new crop varieties giving yields on poor soils and in low fertilization conditions should therefore be considered as new aspects, such as their responsiveness to mycorrhizal fungi, which has never consciously been taken into account during plant domestication (Sawers et al., 2008). Arbuscular mycorrhizal symbiosis can be beneficial to crops and agroecosystems in many ways including resistance to pests and improved soil structure (Thomas et al., 2017). The mycorrhizal fungi contribute to the nutritional quality of edible plant organs.

The signaling molecules released by arbuscular and ectomycorrhizal fungi are to be identified, and the potential Myc factor from arbuscular mycorrhizae is under investigation (Bucher et al., 2009), which is based on the potential similarities with the Nod factor. The study of microbes associated with mycorrhizal fungi and their role as a third component of symbiosis is promising. Hence, the mycorrhizal fungal community hidden in the soil can be exploited for the benefit of plants and humans. Newer technologies like CRISPR/Cas9 and proteomics may provide an avenue for accelerating our ability in the discovery of fungal effectors' function (Selin et al., 2016).

Table 13.1 Types and mechanisms of plant–fungal interactions

| Fungal partner | Type of interaction | Chemicals or metabolites involved | Effect | References |
|---|------------------------------|---|---|---|
| <i>Moniliophthora roreri</i> and <i>Trichoderma harzianum</i> | Phytopathogen–endophyte | Butenolide, harzianolide, sorbicillinol | Compounds produced by the phytopathogen spatially localized in the interaction zone. | Tata et al. (2015) |
| <i>Trichoderma atroviride</i> and <i>Arabidopsis</i> sp. | Endophyte–plant | Indole–acetic acid-related indoles | The fungus colonizes plant roots, thus promoting growth and enhancing systemic disease resistance of the plant. | Salas-Marina et al. (2011) |
| <i>Stachybotrys elegans</i> and <i>Rhizoctonia solani</i> | Mycoparasite–host | Trichothecenes and atranones | <i>S. elegans</i> induced alterations in the metabolism and growth of <i>R. solani</i> , thereby downregulating the production of antimicrobial compounds by <i>R. solani</i> . | Chamoun et al. (2015) |
| <i>Aspergillus nidulans</i> | Microbial community | Aromatic polyketides | Physical interaction between different microbes leads to the activation of the fungal silent genes coding for secondary metabolite. Alterations in the fungal histone are triggered by the actinomycete. | Schroeckh et al. (2009) and Nutzmann et al. (2011) |
| <i>Rhizopus</i> species and rice plant | Symbiont–phytopathogen–plant | Rhizoxin, bongkreki acid, and enacyloxins | In the absence of the endosymbiont, the fungus cannot form spores. The endosymbiont is the causal agent of Rice seedling Blight and produces the phytotoxin rhizoxin. In turn, the growth of the endosymbiont is induced by the fungus. | Partida-Martinez and Hertweck (2005), Partida-Martinez et al. (2007) and Ross et al. (2014) |

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Chapter 14

Fungal Biofertilizer: An Alternative for Sustainable Agriculture



Alka Panda

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14.1 Introduction

Why did the UN World Food Programme (WFP) receive the 2020 Nobel Peace Prize? And what does that have to do with nutrition? The answer to both questions is embedded in renewed concerns globally about food insecurity (Patrick Webb, 2020).

Since the dawn of human civilization, mankind has faced a myriad of pandemics. Presently, the COVID-19 pandemic has not only impacted food trade, food supply chains, and markets, but also people's lives, livelihoods, and nutrition. According to the latest UN estimates, as many as 132 million people went hungry in 2020 as a result of the economic recession triggered by the pandemic. At the same time, 135 million people suffer from acute food insecurity and are in need of urgent humanitarian assistance. Equally urgent is the compounding threat of the pandemic on existing crises – such as conflict, natural disasters, climate change, urban encroachment, loss of biodiversity, pests and animal diseases – that are already stressing our food systems and triggering food insecurity around the globe (HLPE, 2020).

It is in this context that the Nobel Peace Prize committee concluded that tackling hunger and malnutrition, especially in the context of COVID-19, represents a worthy cause, and that the WFP is a worthy champion of that cause (Patrick Webb, 2020).

To urgently minimize COVID-19's damaging effects on food security and nutrition while transforming global food systems to make them more resilient, sustainable, and equitable, FAO calls for immediate action in seven key priority areas, including food security (FAO, 2020).

The CFS Principles for Responsible Investment in Agriculture and Food Systems – also known as RAI – are a set of 10 principles that seek to guide all types of agricultural investment to ensure that it respects a range of environmental, social, and economic goals and supports sustainable food systems (FAO, 2014).

In the midst of such nebulous challenges, the problem of inadequate nutrition is so acute that it is beyond any single type of nutrient source to accept the challenge of appropriate nutrient supply. Integrated use of all the sources such as mineral fertilizers, organic manures, biofertilizers, etc., is the only alternative for improving soil fertility. The use of organic manures and mineral fertilizers is in practice, but

the use of biofertilizer in agriculture is not very popular. Hence, there is a need to make its use popular (Singh et al., 2014).

On one hand, it is essential that indigenous agricultural productivity should be enhanced significantly to sustain through the crucial times without degrading the environmental and agricultural ecology further by relying on sustainable agricultural practices, with minimal use of nonrenewable resources (Figueiredo et al., 2017). On the other hand, unless a situation is created for facilitating social and economic access of the rural poor to the developed sources, the organic potential of the country would not become fully available for crop production (Kumar et al., 2010). In the present scenario, exploring biofertilizers is a potent option to look into.

14.1.1 Persisting Effects of Green Revolution

Chemical fertilizers have aided farmers in time and cost-effective production of commercial crops since the 1930s. In India, the green revolution in the 1960s (Somvanshi et al., 2020) led to high productivity of crops through various adapted measures, like adoption of high-yielding variety (HYV) of seeds, which responded well to the highly increased use of inorganic fertilizers and pesticides (Singh, 2000; Brainerd & Menon, 2014). The green revolution, which was beneficial in ensuring food security, has unintended but harmful consequences on agriculture and human health (Singh, 2000; Clasen et al., 2019; Taylor, 2019). Their overuse has been found to harden the soil, cause acidification and dehydration of the soil, decrease fertility, pollute air and water, and release greenhouse gases, thereby bringing hazards to human health and the environment. Extensive use of chemical fertilizers in replenishing NPK depletes essential soil nutrients and minerals, resulting in soil degradation and the loss of equilibrium of a stable soil. This eventually led to poor crop yield, with harvest being more susceptible to pests and diseases. Besides this, chemical fertilizers can cause root burn or fertilizer burn. The nitrate salts present in the nitrogen fertilizers, being water soluble, cause ground and surface water contamination with an accumulative effect. Continuous use of chemical fertilizers is reported to negatively impact the soil inhabitants, beneficial for plant growth. In their larger threat to the environment, animals, and human health, chemical runoff of the excess fertilizer into our water bodies, ponds, streams, ground water, etc., causes eutrophication owing to oxygen depletion that eventually damages the aquatic life. Overapplication of chemical fertilizer to plants may cause the leaves to turn yellow or brown, a condition known as chemical leaf scorch. Leaf scorch can cause the leaves of the plant to wither and may cause the plant to die. Chemical fertilizers destroy soil crumbs, resulting in a highly compacted soil with reduced drainage and air circulation, referred to as soil friability effect. The use of synthetic chemicals jeopardizes soil health, which then leads to fewer or poor quality crops in the long term. Due to the green revolution, India even lost almost 1 lakh varieties of indigenous rice (Prasad, 2016). Globally, agriculture is on an unsustainable track and has a high ecological footprint now (Prasad, 2016).

On one hand, majority of Indian soils are deficient in many macro and micro-nutrients (Fertiliser Association of India, 2011); and on the other hand, with the injudicious application of chemical fertilizers alone, agricultural productivity is declining and environmental quality is deteriorating (Rakshit et al., 2015). Under such circumstances, only bio-based agro-economy is most viable for agro-industry, farmers, and consumers without compromising on soil and plant health.

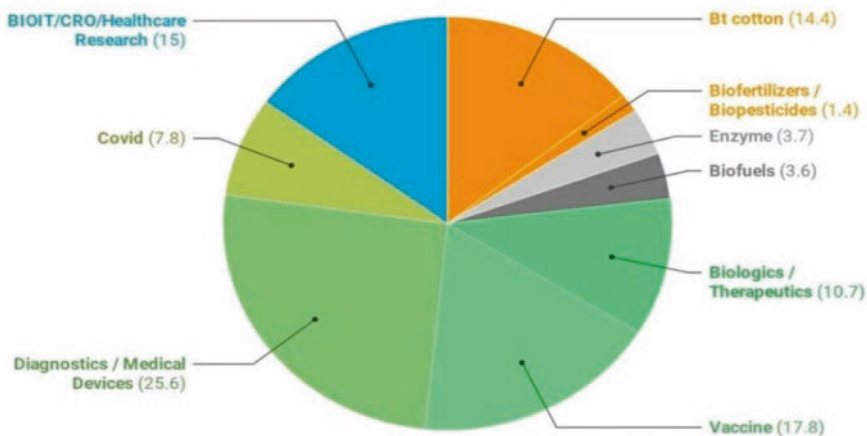
14.1.2 Biofertilizer: An Agent of Bio-based Agro-economy

The solution lies in “biofertilizer,” one of the key subsegments of Indian bio-economy (see Fig. 14.1). Bioeconomy or bio-based economy is a more recent phenomenon. The commonly accepted definition of bioeconomy is the production, utilization, and conservation of biological resources, including related knowledge, science, technology, and innovation, to provide information products, processes, and services across all economic sectors aiming toward a sustainable economy (BIO-ECONOMY INDIA.1615182060_Indian_BioEconomy_Report 2021.pdf).

The benefits of bioeconomy include increased security, economic advantages to farmers, industry, rural communities, and society, environmental benefits at the global, regional, and local levels, and other benefits to society in terms of human health and safety (Kulshreshtha et al., 2011).

[KEY SUB-SEGMENTS]

Figures are in % Share



The Total BioEconomy of India during 2019 was \$70.2 Billion. BioAgri, BioIndustrial, BioPharma, and Bio-based IT services like contract research, development, and research services are important classifications.

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Fig. 14.1 Biofertilizer is one of the key subsegments of bioeconomy. (Source: https://birac.nic.in/webcontent/1615182060_Indian_BioEconomy_Report_2021.pdf)

Bioeconomy can be hugely promoted by the extensive use of biofertilizer – a ready-to-use live formulation of soil microorganisms, which on application to seed, root, or soil, mobilizes the availability of nutrients by their biological activity. These are nothing but selected strains of beneficial soil microorganisms cultured in the laboratory and packed in a suitable carrier, which can be used for either seed treatment or soil application. Biofertilizers generate plant nutrients like nitrogen and phosphorous through their activities in the soil or rhizosphere and make them available to plants in a gradual manner. Biofertilizers are gaining momentum recently due to their availability to maintain soil health, minimize environmental pollution, and cut down the use of chemicals in agriculture. In rainfed agriculture, these inputs gain added importance in view of their low cost as most of the farmers are small and marginal and cannot afford expensive chemical fertilizers (Bisen et al., 2015). Biofertilizers are also ideal input for reducing the cost of cultivation and for practicing organic farming. The use of cheap and eco-friendly inputs like biofertilizers is especially important in India where most of the farming will continue to be in the hands of small farmers.

14.1.3 Types of Biofertilizers

Biofertilizers comprise “microbial inoculants” or assemblage of live or latent cells of efficient strains of nitrogen-fixing, phosphate-solubilizing, sulfur-oxidizing, or cellulolytic microorganisms used for application of seed, soil, or composting areas with the objective of increasing the numbers of such microorganisms and accelerate certain microbial processes to augment the extent of the availability of nutrients in a form that can be easily assimilated by plants. In large sense, the term may be used to include all organic resources (manure) for plant growth that are rendered in an available form for plant absorption through microorganisms or plant associations or microbial interactions (Rao, 1999). In addition to other biofertilizers available, fungal biofertilizers exert direct or indirect benefits on plant growth and crop yield through biological mechanisms. They can cause stimulation of plant growth by hormone action or antibiosis and by decomposition of organic residues (Pal et al, 2015; Carvajal-Muñoz & Carmona-García, 2012). The following types of biofertilizers are available to the farmers in India:

1. Nitrogen-fixing biofertilizers (N-BF)

- Symbiotic BF with all legumes – *Rhizobium*, *Bradyrhizobium*
- Nonsymbiotic BF for cereals, vegetables, horticultural crops – *Azotobacter*
- Associative BF for millets, maize, etc. – *Azospirillum* (blue-green algae, *Azolla*, *Gluconacetobacter diazotrophicus* are also N-BF, but is yet to be included in FCO) (<https://geographyandyou.com/biofertilisers-in-indian-agriculture/>)

2. Phosphorus biofertilizers (P-BF)

- Phosphorous-solubilizing biofertilizers (PSB) (*Bacillus*, *Pseudomonas*, *Aspergillus*, *Penicillium*, *Fusarium*, *Trichoderma*, *Mucor*, *Ovulariopsis*, *Tritirachium*, and *Candida*)
- Phosphate-mobilizing biofertilizers – *Mycorrhiza* (*glomus*, *gigaspora*, etc., for all crops)

3. Potash biofertilizers (K-BF) (microbes like *B. mucilaginosus* and *F. aurantia* for all crops)

4. Zinc solubilizers (Z- BF) (*Bacillus* species are capable of zinc solubilization)

5. Plant growth-promoting biofertilizers (*Pseudomonas*)

6. Enriched compost biofertilizers = cellulolytic fungal cultures (*Chaetomium bositrychodes*, *C. olivaceum*, *Humicola fuscoatra*, *Aspergillus flavus*, *A. nidulans*, *A. niger*, *A. ochraceus*, *Fusarium solani*, and *F. oxysporum*)

Generally, biofertilizers are available in a solid (using peat, lignite, charcoal, etc., as a carrier) or in a liquid base (using broth involving additives like poly vinyl pyrrolidone, gum, biosurfactants, etc., or by promoting dormant cells/spores). Biofertilizers may be prepared from either single or multiple strains, and experiments are also being conducted for formulating freeze-dried, granular, and polyacrylamide-entrapped inoculants. At present, however, the best production technology and packaging are yet to be obtained (<https://geographyandyou.com/biofertilisers-in-indian-agriculture/>). Figure 14.2 and Table 14.1 show the general

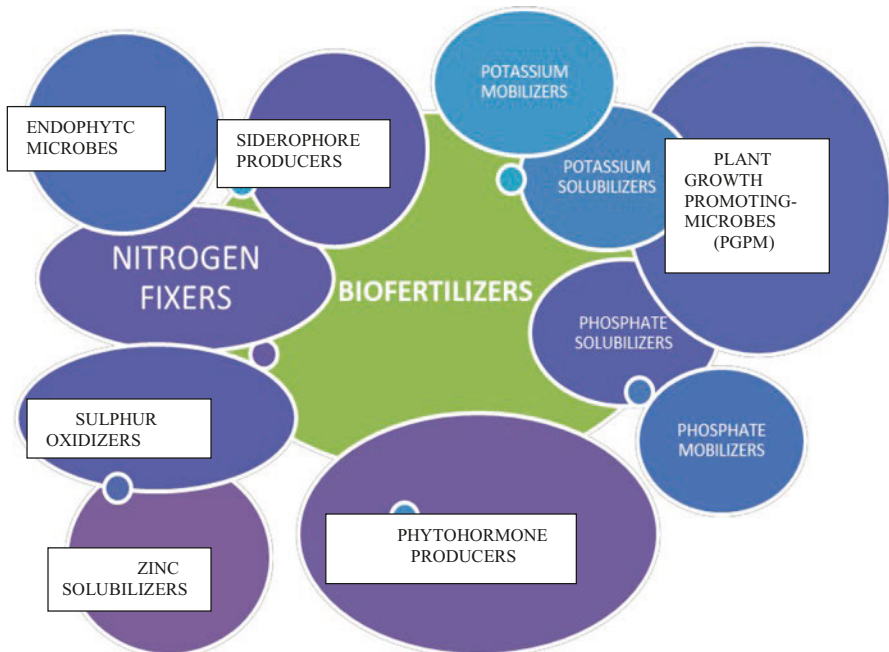


Fig. 14.2 Types of biofertilizers

Table 14.1 Classification of biofertilizers

| Types of biofertilizers | Mode of action | Examples | References |
|--|---|---|------------------------------|
| N₂-fixing biofertilizers | | | |
| 1. Free-living; nonsymbiotic | Increase soil nitrogen content by fixing atmospheric N and make it available to the plants. | <i>Azotobacter</i> , <i>Beijerinckia</i> , <i>Clostridium</i> , <i>Klebsiella</i> , <i>Anabaena</i> , <i>Bacillus polymyxa</i> , <i>Rhodospseudomonas</i> , <i>Rhodospirillum</i> , <i>Chromatium</i> , <i>Nostoc</i> | Choudhury and Kennedy (2004) |
| 2. Symbiotic | | <i>Rhizobium</i> , <i>Frankia</i> , <i>Xanthomonas</i> and <i>Mycobacterium</i> and <i>Anabaena azollae</i> | |
| 3. Associative symbiotic | | <i>Azospirillum</i> sp., <i>Enterobacter</i> , <i>Acetobacter diazotrophicus</i> , <i>Herbaspirillum</i> sp., <i>Azoarcus</i> sp., <i>Alcaligenes</i> | |
| P-solubilizing biofertilizers | | | |
| 1. Bacteria | Solubilize the insoluble forms of P in the soil into soluble forms by secreting organic acids and lowering soil pH to dissolve bound phosphates. | <i>Bacillus megaterium</i> var. <i>phosphaticum</i> , <i>Bacillus subtilis</i> , <i>Bacillus circulans</i> , and <i>Pseudomonas striata</i> | Board (2004) |
| 2. Fungi | | <i>Penicillium</i> sp., and <i>Aspergillus awamori</i> | |
| P-mobilizing (microphos) biofertilizers | | | |
| 1. Arbuscular mycorrhiza | Transfer phosphorus from the soil to the root cortex. These are broad spectrum biofertilizers. | <i>Glomus</i> sp., <i>Gigaspora</i> sp., <i>Acaulospora</i> sp., <i>Scutellospora</i> sp., <i>Sclerocystis</i> sp. | Chang and Yang (2009) |
| 2. Ectomycorrhiza | | <i>Laccaria</i> sp., <i>Pisolithus</i> sp., <i>Boletus</i> sp., <i>Amanita</i> sp. | |
| 3. Ericoid mycorrhiza | | <i>Pezizella ericae</i> | |
| 4. Orchid mycorrhiza | | <i>Rhizoctonia solani</i> | |
| K-solubilizing biofertilizer | | | |
| 1. Bacteria | Solubilize potassium (silicates) by producing organic acids that decompose silicates and help in the removal of metal ions and make it available to plants. | <i>Bacillus mucilaginosus</i> , <i>B. circulanscan</i> , <i>B. edaphicus</i> , <i>Arthrobacter</i> sp. | Etesami et al. (2017) |
| 2. Fungi | | <i>Aspergillus niger</i> | |
| K-mobilizing biofertilizer | | | |
| 1. Bacteria | They mobilize the inaccessible forms of potassium in the soil. | <i>Bacillus</i> sp. | Jha (2017) |
| 2. Fungi | | <i>Aspergillus niger</i> | |

(continued)

Table 14.1 (continued)

| Types of biofertilizers | Mode of action | Examples | References |
|---|--|--|-----------------------|
| Biofertilizers for micronutrients | | | |
| 1. Silicate and zinc solubilizers | Solubilize the zinc by proton, chelated ligands, acidification, and by oxidoreductive systems. | <i>Bacillus</i> sp., <i>Bacillus subtilis</i> , <i>Thiobacillus thiooxidans</i> , <i>Saccharomyces</i> sp., <i>Mycorrhiza</i> sp. | Kamran et al. (2017) |
| 2. Sulfur oxidizer | Oxidize sulfur to sulfates that are usable by plants. | <i>Thiobacillus</i> sp. | Itelima et al. (2018) |
| Plant growth-promoting biofertilizers | | | |
| 1. Plant growth-promoting rhizobacteria | Produce hormones that promote root growth, improve nutrient availability, control plant pathogens, and improve crop yield. | <i>Pseudomonas fluorescense</i> | Backer et al. (2018) |
| 2. Plant growth-promoting fungi | | <i>Aspergillus</i> sp., <i>Fusarium</i> sp., <i>Penicillium</i> sp., <i>Piriformospora</i> sp., <i>Phoma</i> sp., <i>Trichoderma</i> sp. | |

Table 14.2 Differences between the two production technologies of biofertilizers

| Carrier-based | Liquid-based |
|---------------------------------|---------------------------------------|
| Low shelf-life | Longer shelf-life |
| Easier to produce | Easier to produce |
| Temperature sensitive | Temperature tolerant |
| Less investment | Higher investment for production unit |
| Low cell counts | High cell counts |
| Contamination prone | Contamination free |
| Less effective | More effective |
| Product may not be 100% sterile | Product can be 100% sterile |
| Automation difficult | Automation is easier |
| Low cost | High cost |
| High dose requirement | Low dose requirement |

classification of biofertilizers, and Table 14.2 shows the differences between two different technologies of biofertilizer formulation.

14.1.4 Advantages of Biofertilizer

Biofertilizers are the natural way to get the benefits of synthetic fertilizers without risking the quality of soil health and crop products. Biofertilizers are known to play a number of vital roles in soil fertility, crop productivity, and production in agriculture as they are eco-friendly but cannot replace chemical fertilizers, which are indispensable for getting maximum crop yields. Biofertilizers can increase the crop yield

Fig. 14.3 Soil nutrients for plant growth.
(Source: FAI)

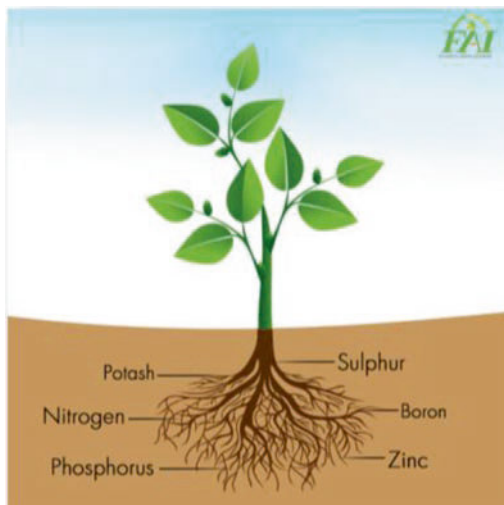


Fig. 14.4 Food security.
(Source: FAI)



by 20–30%. In addition, biofertilizers are cost-effective when compared to synthetic fertilizers. Biofertilizers unleash nutrients (Fig. 14.3) from the soil and provide us food security (Fig. 14.4).

Some of the important functions or roles of biofertilizers in agriculture are

- Phosphate-mobilizing or phosphorus-solubilizing biofertilizers/microorganisms (bacteria, fungi, mycorrhiza, etc.) convert insoluble soil phosphate into soluble forms by secreting several organic acids and under optimum conditions, can solubilize/mobilize about 30–50 kg P_2O_5 /ha, and crop yield may increase by 10–20%.

- Mycorrhiza or arbuscular mycorrhiza, when used as biofertilizer, enhance uptake of P, Zn, S, and water, leading to uniform crop growth and increased yield and also enhance resistance to root diseases and improve hardiness of transplant stock (Pal et al., 2014).
- Liberate growth-promoting substances and vitamins and help to maintain soil fertility.
- Act as antagonists and suppress the incidence of soil-borne plant pathogens and, thus, help in the biocontrol of diseases.
- Play an important role in the recycling of plant nutrients.
- Supplement chemical fertilizers for meeting the integrated nutrient demand of the crops.
- Renewable source of nutrients.
- Sustain soil health.
- Supplement chemical fertilizers.
- Replace 25–30% chemical fertilizers.
- Increase the grain yields by 10–40%.
- Decompose plant residues and stabilize C:N ratio of soil.
- Improve texture, structure, and water-holding capacity of soil.
- No adverse effect on plant growth and soil fertility.
- Stimulate plant growth by secreting growth hormones.
- Secrete fungistatic and antibiotic-like substances.
- Solubilize and mobilize nutrients.
- Eco-friendly, non-pollut, and cost-effective.

The microorganisms present in biofertilizers are available in nature. Initially, these organisms are isolated from different sources such as the root nodule for rhizobium, soil for other microbes, etc., and developed in specific media for mass culturing. Hence, the understanding of the biological attributes of beneficial microbes and their mutual interactions in conjunction with the modern agricultural practices would help to develop the most potential biofertilizer for sustainable agro-practice. Potential roles played by biofertilizers for plant growth are illustrated in Fig. 14.5.

14.2 Soil Microbiome: A Repository of Nutrients for Plant–Microbe Holobiont

Years of extensive research have revealed the significance of microorganisms in plant health. The difficulty of culturing transplants of different species in the absence of bacteria and fungi is widely known (Hardoim et al., 2008), which strongly implies the importance of such microorganisms in plant growth. It is readily acknowledged that cooperative microbial symbionts play an important role in their host's life and fitness (Kiers & van der Heijden, 2006). Thus, a plant can be regarded as a holobiont comprising the host plant and its microbiota (Zilber-Rosenberg & Rosenberg,

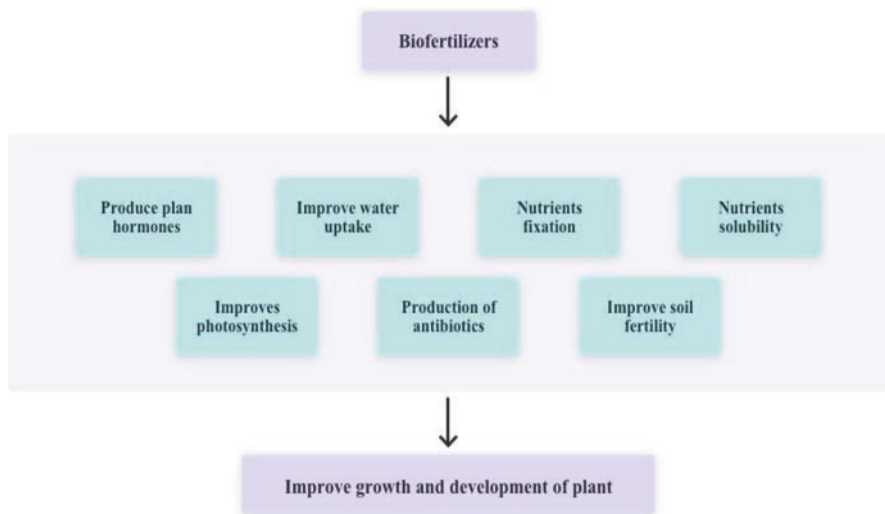


Fig. 14.5 Potential role played by biofertilizers in the growth and development of plants

2008). On the other hand, rhizospheric or root-associated microorganisms, beneficial for plant growth, are strongly influenced by the soil microbiome in the vicinity of the roots.

It is quite evident that the plant–microbiome provides the host plant with a variety of essential functions to adapt to the local environment. However, plant–microbe holobionts are impacted by different factors, generating complex interactive systems (Tosi et al., 2020). One of the most important factors is the soil microbiome, which facilitates plant growth and enhances crop yield. Owing to their rich diversity, complexity of interactions, and numerous metabolic pathways, microbes are an amazing resource for biological activity (Emmert & Handelsman, 1999; Alabouvette et al., 2006; Tejesvi et al., 2007; Mitchell et al., 2008; Raghukumar, 2008) and constitute the most important niches in the soil ecosystem. Such microbiological interactions and activities are responsible for earth geochemical stability, climatic and biogeochemical cycles, and have been exploited for enhanced agricultural, environmental, and forestry management (Barea et al., 2005; Tringe et al., 2005; Hansel et al., 2008; Frey-Klett et al., 2011; Deveau et al., 2018; Odoh et al., 2019a, b). These interactions are also responsible for enhancing plant growth by improving nutrient uptake and developing tolerance for biotic and abiotic stress (Bhardwaj et al., 2014; Clark et al., 2009). Strategic and applied research has demonstrated that certain cooperative microbial species, immersed in a framework of interactions, can be exploited as a low-input biotechnology to help sustainable, environmentally friendly, agrotechnological practices. The global need and demand for enhanced crop production, agroecology, and ecological immunity require efficient use of such biological inoculants, composed of beneficial soil microorganisms unlike their chemical counterparts (Sivasakthivelan & Saranraj, 2013). The quantity of nutrient availability rendered by these microorganisms is shown in Table 14.3.

Table 14.3 Benefits rendered by beneficial microbes

| Name | Crops suited | Benefits usually seen | Remarks |
|---|---|---|---|
| Rhizobium strains | Legumes like pulses, groundnut, soybean | 10–35% yield increase, 50–200 kg N/ha | Fodders give better results. Leaves residual N in the soil. |
| Azotobacter | Soil treatment for non legume crops including dry land crops | 10–15% yield increase – adds 20–25 kg N/ha | Also controls certain diseases. |
| Azospirillum | Nonlegumes like maize, barley, oats, sorghum, millet, sugarcane, rice, etc. | 10–20% yield increase | Fodders give higher/enrich fodder response. Produces growth-promoting substances. It can be applied to legumes as co-inoculant. |
| Phosphate solubilizers (there are two bacterial and two fungal species in this group) | Soil application for all crops | 5–30% yield increase | Can be mixed with rock phosphate. |
| Blue-green algae and <i>Azolla</i> | Rice/wet lands | 20–30 kg N/ha, <i>Azolla</i> can give biomass up to 40–50 tons and fix 30–100 kg N/ha | Reduces soil alkalinity, can be used for fishes as feed. They have growth-promoting hormonal effects. |
| Mycorrhizae (VAM) | Many trees, some crops, and some ornamental plants | 30–50% yield increase, enhances uptake of P, Zn, S, and water | Usually inoculated to seedlings. |

Pathak et al., 2017 stated that fungal–fungal (tropical plant and their foliar endophytes) and bacterial–fungal (arbuscular mycorrhizal fungi and plant growth-promoting rhizobacteria) associations enhance the turnover of soil organic matters apart from facilitating nutrient mobilization for plant growth through nitrogen (N) fixation in leguminous crops (Franco et al., 2011; Nuti & Giovannetti, 2015). The community of mycorrhiza colonizing a plant has been extensively studied over several decades. However, it was only in 2002 that the diversity of fungi colonizing the plant root was shown to be much greater than previously believed (Vandenkoornhuysen et al., 2002), indicating a gap in our knowledge concerning microorganisms colonizing roots. This explains the growing advances in mycobiome, exploring its inherent potentials in sustainable agriculture (Berruti et al., 2016). This review hence focuses on the beneficial soil fungi, which can be developed into potential biofertilizers.

When we focus upon the soil and plant health, we cannot overlook the nuisance created by agrochemicals, including pesticides – one of the deadly human interventions in destabilizing the agroecosystem and environmental ecology. Pesticides are double-edged weapons, which, although targeted on specific targets, inhibit nontarget organisms including the soil mycoflora (Daisley et al., 2022; Lata et al., 2021; Hashimi et al., 2020; Pandey et al., 2005).

Many soil-applied pesticides are intentionally introduced into the soil environment for the control of soil-borne pests and pathogens, resulting in the accumulation of their residues and metabolites in soil at unacceptably high levels (Gamon et al., 2003; Shalaby & Abdou, 2010). On the contrary, the extensive use of pesticides has often resulted in the development and evolution of pesticide resistance in insect pests, plant pathogens, and weeds (Aktar et al., 2009; Pimentel, 2005). Moreover, in the absence of a clear obsolete pesticide management strategy, over the years, significant amounts of obsolete pesticides have been stockpiling in developing countries, which endanger the environment and health of millions of people (Farrera, 2004; Karstensen et al., 2006; Ortiz-Hernandez & Sanchez-Salinas, 2010) and should be managed effectively (Karstensen et al., 2006; Dasgupta et al., 2010; Martinez, 2004).

The most promising opportunity for maximizing benefits and minimizing risks is to invest time, money, and effort into developing a diverse toolbox of crop protection and production strategies that include safe products and practices integrating chemical approaches into an overall and ecologically based framework that will optimize sustainable production, environmental quality, and human health (Damalas, 2009).

When pesticides are used rationally and carefully, in conjunction with other technologies like biopesticides in integrated pest management systems, it is more likely that their use will be justifiable (Cooper & Dobson, 2007). Therefore, biological control agents that are used in the preparation of biopesticides should be tolerant enough to pesticides (Arshad & Aishatul, 2015).

The potentiality of pesticide-tolerant fungi as biofertilizers can pave the way for bio-entrepreneurs. The combination of these potentials when fully harnessed under agricultural scenario will help to sustain agriculture and boost food security globally (Bending et al., 2002; Verma et al., 2014; Tomer et al., 2021). In the present review, we need to understand how agricultural sustainability and economic stability of rural India can largely depend upon fungal biofertilizers.

Focusing on sustainable biofertilizer production and low-cost environmental remediation, this review seeks to identify fungal species being examined individually for their unique capabilities of nutrient mobilization, nutrient solubilization, bioremediation, nitrogen fixation, or biocontrol mechanism order. This will help us to focus on the most promising bioinoculant or assemblage of bioinoculants for promoting soil and plant health.

14.2.1 Soil Mycobiome: Pool of Promising Bioagents for Sustainable Agriculture

The myco-diversity in soil plays a pivotal role affecting soil and plant health as symbionts, decomposers, nutrient mobilizers, agents for bioremediation, antistress, and pest control (Owen et al., 2015; Ajmal et al., 2018). Intensive research on the identity, abundance, distribution, and function of soil mycobiome, and on their various roles in soils has been fundamental for a better understanding of fungal biology, interrelationships, as well as the physiological mechanisms that help in the

maintenance of soil health. It is also important to assess the common dynamics that may occur due to the coexistence of fungal populations with other microbes in most cases of host–symbiont associations.

The rapid expansion of the soil mycobiome and their interaction with plants and other microbes pave way for sustainable agriculture to bring increased awareness to the diverse effect of the soil mycobiome on plant and soil health. The objective of the study is to collate a series of reviews and research articles on the role of the soil mycobiome and other associated microbes in sustainable agriculture affecting agroecology and agroecology of the nation.

14.2.2 *Biofertilizer in Agriculture*

Biofertilizers include materials derived from living organisms and microbial sources (Rola, 2000; Chen, 2006) with various benefits, such as increased access to nutrients (Chen, 2006; Vessey, 2003; Hart & Trevors, 2005), providing growth-promoting factors for plants, mediating the process of composting, and effective recycling of solid wastes (Gaur & Adholeya, 2004; Das et al., 2007). Biofertilizers are fundamentally microbial inoculants, produced from microbial cultures, that can improve soil fertility and crop productivity such as *mycorrhizae* (Malik et al., 2005; Marin, 2006). *Mycorrhizae* are fungi that form mutualistic relationships with roots of 90% of plants (Rinaldi et al., 2008; Das et al., 2007; Gaur & Adholeya, 2004). *Mycorrhizae* promote absorption of nutrients and water, control plant diseases, and improve soil structure (Rola, 2000; Marin, 2006; Zhao et al., 2003; Chandanie et al., 2006). Plants colonized by *mycorrhizae* grow better than those without them (Yeasmin et al., 2007; Singh et al., 2008) and are beneficial in natural and agricultural systems (Marin, 2006; Adholeya et al., 2005).

Some of the common microorganisms usually used as biofertilizers or in its composition are nitrogen-fixing agents (N-fixer) (Verma et al., 2016; Rana et al., 2019a, b; Odoh et al., 2019a), plant growth-promoting rhizobacteria (PGPRs) (Verma et al., 2016; Yadav et al. 2015, 2018a), potassium (Yadav et al., 2017; Kour et al., 2020), phosphorus solubilizers (Yadav et al., 2017), cyanobacteria, and endo- and ecto-mycorrhizal fungi (López-Bucio et al., 2015; Itelima et al., 2018). Intrinsically, these organisms possess the ability to produce a wide variety of extracellular enzymes for the breakdown of organic matter (Frąc et al., 2018). On the other hand, the bioagent improves plant's vigor and tolerance to abiotic and biotic factors while serving as an eco-friendly and cost-effective alternative (Rouphael et al., 2015; Itelima et al., 2018).

Agroeconomically, biofertilizers are environmental-friendly, cost-effective, and a renewable source of plant nutrients that have gained acceptance over chemical fertilizers (Kour et al., 2020; Verma et al., 2019; Yadav et al., 2020c). These bio-based materials play an important role in maintaining soil fertility, nutritional enrichment, and sustenance of healthy soil for generations (Mishra et al., 2015; Fuentes-Ramirez & Caballero-Mellado, 2005). Some fungal inoculants and their mode of action, as seen in Table 14.4, improve crop yield by enhancing a number of

Table 14.4 Some fungi used as biofertilizers and their mode of action

| Biofertilizer agents | Mode of actions/ functions | Microbial species | Plant names |
|--------------------------|--|--|---|
| <i>Fusarium</i> spp. | Enrich compost biofertilizers degrade plant residues and maintain C:N balance in the soil | <i>Fusarium solani</i> and <i>F. Oxysporum</i> | Tomato |
| <i>Trichoderma</i> spp. | Enhance compost degradation | <i>Trichoderma harzianum</i> + <i>Aspergillus</i> + <i>Penicillium</i> | Potato, corn, tomato, peanut, beans, cotton, soybean |
| <i>Mycorrhizal fungi</i> | Reduce the impact of environmental stress Improve plant health status while enhancing optimum mineral content | <i>Glomus intraradices</i> | Vegetable crops, annuals, perennials, trees and shrubs (eg: onion, garlic, carrot, ornamental grasses, ferns, aloe, rose bush, fruit trees etc) |
| <i>Aspergillus</i> spp. | Solubilize unavailable phosphate for plant use through the secretion of organic acid Improve soil quality with subsequent uses | <i>Aspergillus niger</i> | |
| <i>Penicillium</i> spp. | Supply essential minerals (p, Mn, Zn, Fe, Co, Cu, and Mo) to the plants Inducement of abiotic stress resistance in young seedling Protect the plants from fungal pathogens | <i>P. bilaji</i> , <i>P. italicum</i> , <i>P. albidum</i> , <i>P. frequentans</i> , <i>P. simplicissimum</i> , <i>P. rubrum</i> , <i>P. expansum</i> , <i>P. oxalicum</i> , <i>P. citrinum</i> | Wheat, soybean, canola |
| <i>Chaetomium</i> spp. | Enhance compost degradation | <i>C. bostrychodes</i> , <i>C. oliveaceum</i> | Tomato, corn, rice, pepper, citrus, durian, birds of paradise, carnation |
| <i>Gliocladium</i> spp. | Reduce the incidence of damping-off issue | <i>Gliocladium catenulatum</i> | |
| <i>Tritirachium</i> spp. | Organic acids secretion for mineral dissolution | <i>T. album</i> , <i>T. egenum</i> | |

biochemical processes, leading to increase in uptake of nutrients, stimulation of plant growth hormones, antibiosis, and decomposition of organic residues (Odoh et al., 2019a, b).

14.3 Fungal Biofertilizers: Biological Mechanism and Interactions

The vital role of fungal biofertilizer in agrobiolgy and biotechnological advances has been well documented (Zeilinger-Migsich & Mukherjee, 2014). These fungi, through nutritional versatility, mechanisms, and biochemical processes, interact with other organisms (bacteria, plants, and animals) in their natural habitat. At both intra- and inter-specific interactions, fungal biofertilizers comprise microbial inocula or assemblages of living microorganisms that exert direct or indirect benefits on plant growth and crop yield through different mechanisms (Fuentes-Ramirez & Caballero-Mellado, 2005). These microorganisms are able to fix atmospheric nitrogen or solubilize phosphorus, decompose organic material, or oxidize sulfur in the soil properties (Marin, 2006) that are beneficial to agricultural production in terms of nutrient supply (Malik et al., 2005). One type of biofertilizer is the arbuscular mycorrhizal fungi, which are probably the most abundant fungi in agricultural soil (Khan, 2006; Marin, 2006). The inocula improve crop yield because of increased availability or uptake or absorption of nutrients, stimulation of plant growth by hormone action or antibiosis, and by decomposition of organic residues (Wani & Lee, 2002). Žifčáková et al. (2016) in their report maintained that fungal biofertilizers regulate the balance of carbon and other nutrients in the decomposition process. They play a vital role in maintaining the soil's primary and secondary nutrient requirements – through potassium and phosphate solubilization, nitrogen fixation, and mineralization. They (Table 14.4) also help in the biodegradation of organic matter and pollutants in the soil environment (Adesemoye & Kloepper, 2009; Sinha et al., 2014; Odoh et al., 2019b). Selected fungal species that are used as biofertilizers are mentioned below.

14.3.1 *Mycorrhizal Fungi Used as Biofertilizers: Soil–Fungal–Plant (Root) Interactions*

Plants that suffer from nutrient scarcity, especially P, N, Zn, Cu, Fe, S, and B, develop mycorrhiza, which includes plants belonging to different groups (Zhu et al., 2008). *Mycorrhizae* uniquely form mutualistic symbiotic relationships with plant roots (Fig. 14.6) of more than 80% of land plants including many important crops and forest tree species (Smith & Zhu, 2001; Gentili & Jumpponen, 2006; Rinaldi et al., 2008) including xerophytes, epiphytes, and hydrophytes (Rai et al., 2013).



Fig. 14.6 Mycorrhizal growth in plant roots

Mycorrhizas enhance uptake of phosphorus (P), nitrogen (N), zinc (Zn), copper (Cu), iron (Fe), sulfur (S), and boron (B).

There are seven types of mycorrhizae: arbutoid mycorrhiza, ectomycorrhiza, endomycorrhiza or arbuscular mycorrhiza, ect-endomycorrhiza, ericoid mycorrhiza, monotropoid mycorrhiza, and orchidoid mycorrhiza (Zhu et al., 2008; Gentili & Jumpponen, 2006; Raina et al., 2000). The two dominant types of mycorrhizae are ectomycorrhizae (ECM) and arbuscular mycorrhizae (AM), which can improve water and nutrient uptake and provide protection from pathogens, but only a few families of plants are able to form functional associations with both AM and ECM fungi (Haskins & Gehring, 2005; Siddiqui & Pichtel, 2008). However, AM fungi are most commonly found in the rhizosphere roots of a wide range of herbaceous and woody plants (Finlay, 2008; Rinaldi et al., 2008). In this review, we focus on ectomycorrhizal fungi and arbuscular mycorrhizal fungi because they are most widespread and are more economically important than others.

14.3.2 Ectomycorrhizae (ECM)

The fungi in this group are characterized by the formation of a thick mantle structure within the intercellular spaces of root cortex. They also form a sheath around the feeder root, thus acting as an interface for nutrient uptake (Bücking & Kafle, 2015). They are surrounded by living cells in the host roots, leading to the development of an extensive network called the Hartig net. This Hartig net acts as both storage and transport organ for phosphorus. Most importantly, ectomycorrhizae are common in members of the families of Pinaceae, Fagaceae, Betulaceae, Salicaceae, and Myrtaceae. Ectomycorrhizal (ECM) fungi form mutualistic symbioses with many tree species (Anderson & Cairney, 2007) without penetrating the living cells in the roots, but only surround them (Gupta et al., 2000; Finlay, 2008; Raina et al., 2000), for example, pine, spruce, larch, hemlock, willow, poplar, oak birch, and

eucalyptus (Dahm, 2006; Rinaldi et al., 2008). Most ECM fungi that are associated with forest trees are basidiomycetes, such as *Amanita* sp., *Lactarius* sp., *Pisolithus* sp., and *Rhizopogon* sp., and many of these are edible (Rinaldi et al., 2008; Le et al., 2007). Some ascomycetes also form mycorrhizae such as *Cenococcum* sp., *Elaphomyces* sp., and *Tuber* sp. (Rinaldi et al., 2008; Finlay, 2008). The importance of ECM fungi to trees is in their ability to increase tree growth due to better nutrient acquisition (Gentili & Jumpponen, 2006). ECM fungi help the growth and development of trees because the roots colonized with ectomycorrhiza are able to absorb and accumulate nitrogen, phosphorus, potassium, and calcium more rapidly and over a longer period than nonmycorrhizal roots. ECM fungi help to break down the complex minerals and organic substances in the soil and transfer nutrients to the tree. ECM fungi also appear to increase the tolerance of trees to abiotic stresses like drought, high soil temperatures, soil toxins, and extremes of soil pH. ECM fungi can also protect roots of trees from biotic stress like pathogens (Odoh et al., 2019a, b; Dahm, 2006).

The most commonly widespread ectomycorrhizal product is inoculum of *Pisolithus tinctorius* (Schwartz et al., 2006; Gentili & Jumpponen, 2006). *Pisolithus tinctorius* has a wide host range, and their inoculum can be produced and applied as vegetative mycelium in a peat vermiculite carrier. These fungus inocula are applied to nursery or forestry plantations (Gentili & Jumpponen, 2006). *Piriformospora indica* (Hymenomycetes, Basidiomycota) is another ECM fungus used as a biofertilizer. This taxon can promote plant growth and biomass production and help plant tolerance to herbivory, heat, salt, disease, drought, and increased below and aboveground biomass (Waller et al., 2005; Tejesvi et al., 2007).

14.3.3 *Endomycorrhizae*

These are a group of fungi that provide biological protection against soil-borne diseases and are associated with most agricultural crops and horticultural soils (Frąc et al., 2018). They occur in most ecosystems of the world and are found in many important crop species such as wheat, maize, rice, grape, soybean, and cotton and also in horticultural species like roses and petunias. The most common among them is arbuscular mycorrhizal fungi (AMF). Bagyaraj and Ashwin (2017), in their work, reported significant increases in crop yield following inoculation with AMF, which aid in stimulating key effects such as root development, improved soil structure, increased nutrient uptake, and mobility of ions. Besides enhancing plant tolerance to stresses, endomycorrhizae improves the general well-being of plants. *Endomycorrhizae* form mutually symbiotic relationships between fungi and plant roots (Ipsilantis & Sylvia, 2007). The plant roots provide substances for the fungi and the fungi transfer nutrients and water to the plantroots (Chen, 2006; Adholeya et al., 2005). Endomycorrhizal fungi are intercellular and penetrate the root cortical cells and form structures called arbuscular vesicles and known as vesicular arbuscular mycorrhiza (VAM), but in some cases no vesicles are formed and they are known

as arbuscular mycorrhiza (AM) (Gupta et al., 2000). The agriculturally produced crop plants that form endomycorrhizae of the vesicular–arbuscular mycorrhiza type are now called arbuscular mycorrhizal (AM) fungi (Raja, 2006). AM fungi belong to nine genera: *Acaulospora*, *Archaeospora*, *Enterophospora*, *Gerdemannia*, *Geosiphon*, *Gigaspora*, *Glomus*, *Paraglomus*, and *Scutellospora* (Finlay, 2008). AM fungi are a widespread group and are found from the arctic to tropics and are present in most agricultural and natural ecosystems. AM fungi help plants to absorb nutrients, especially the less available mineral nutrients such as copper, molybdenum, phosphorus, and zinc (Yeasmin et al., 2007). They increase seedling tolerance to drought, high temperatures, toxic-heavy metals, high or low pH, and even extreme soil acidity (Chen, 2006; Gupta et al., 2000). AM fungi can also affect plant growth indirectly by improving the soil structure, providing antagonist effects against pathogens and altered water relationships (Smith & Zhu, 2001). AM fungi can reduce the severity of soil-borne pathogens and enhance resistance in roots against root rot disease (Akhtar & Siddiqui, 2008a, b; Chen, 2006). This results because of competition for colonization sites or nutrients in the same root tissues and production of fungistatic compounds (Marin, 2006). AM fungi have been found beneficial to the host plants by increasing herbivore tolerance, pollination, soil stability, and heavy metal tolerance (Hart & Trevors, 2005). Mass production of AM fungi has been achieved with several species such as *Acaulospora laevis*, *Glomus clarum*, *G. etunicatum*, *G. intraradices*, *G. mosseae*, *Gigaspora ramisporophora*, and *Gigaspora rosea* (Schwartz et al., 2006), but *Glomus intraradices* is the most common inoculum of endomycorrhizae products (Schwartz et al., 2006; Akhtar & Siddiqui, 2008b; Adholeya et al., 2005). Effective management of AM fungi involves increasing population of propagules such as spores, colonized root fragments, and hyphae using host plants and also by adoption of soil management techniques (Smith & Zhu, 2001; Tiwari et al., 2004; Kapoor et al., 2008). AM fungi are phosphate scavengers that expedite the recruitment of soluble phosphate from soil aquifers. Mycorrhizal fungi are efficient in the uptake of specific nutrients, are resistant against soil-borne pathogens, and play a key role in plant growth promotion, plant protection, and soil quality improvements. Also, besides being widely spread in agricultural systems, they have received wide acceptance as organic farming manure, especially in regions where the quest for sustainable crop production is prioritized.

14.3.4 Bacterial–Fungal Interactions (BFIs)

Studies have revealed that bacteria and fungi coexist as dynamic coevolving groups. Interaction between these microbes plays a huge role as they are considered vital players in driving activities such as biochemical cycles and contributing significantly to plant and animal pathogens (Deveau et al., 2018). BFI by-products are exploited for enhanced agricultural, environmental, and forestry management (Frey-Klett et al., 2011; Odoh et al., 2019a, b). In characterizing BFI, understanding

of microbiomes using molecular tools, chemical and microbial ecology, genomics, and biophysics is essential (Thompson et al., 2017; Bergelson et al., 2019). Freyklett et al. (2011) in their work suggested a shift from disordered polymicrobial communities to a highly specific symbiotic association of fungal hyphae and bacterial cells, following their complexities. Arbuscular mycorrhizal fungi (AMF) and bacteria (plant growth-promoting rhizobacteria [PGPR]) association has been reported to promote crop growth (Pathak et al., 2017). According to Philippot et al. (2013), *Bacillus* sp., *Pseudomonas* sp. (PGPR), and AMF interaction proffers strong viability and causes significant improvement in field study when used singly or in combined application (Pathak et al., 2017). Although this association has positive influence on crop yield, it also enhances soil nutritional status and soil microbial biodata. Studies have shown that PGPR and AMF are key bioinoculants with potential to stop plant's dependence on agrochemicals, thus aiding in sustainable agricultural practices (Pathak et al., 2017; Franco et al., 2011). PGPR boosts plant growth directly (growth-promoting hormones) and indirectly (synthesis of antimicrobial substances Zheng et al., 2018). As aid to the process of mycorrhization, mycorrhizal-helping bacteria (MHB) and PGPR symbiotically interact with mycorrhizal fungi, and mycorrhizal roots for nutrient uptake. Studies have revealed that rhizospheric AMF and PGPR elicit systemic host immune responses for plant resistance (Singh, 2018), just as experimental evidence has proven that co-inoculants of AMF and PGPR give synergistic advantage to crop growth especially in nutrient-limited agricultural soil (Gouda et al., 2018).

14.4 Production, Formulation, and Application of Mycorrhizal Fungi as Fungal Biofertilizers

AM fungi are obligate symbiotic microorganisms since they cannot be grown without the plant host on synthetic media (Hart & Trevors, 2005). As AM fungal inocula must be produced in association with the host plant, therefore there are many constraints to large-scale commercial production. Mass production by pot culture either in the greenhouse or in growth chambers is the most commonly used production method (Marin, 2006; Gentili & Jumpponen, 2006; Bagyaraj & Ashwin, 2017; Raja, 2006; Kapoor et al., 2008). AM fungal inocula have to be prepared by multiplication of the selected fungi in roots of susceptible host plants growing in the sterilized soil or substrates, for example, perlite, vermiculite, peat, sand, or a mixture of them (Naqvi & Mukerji, 2000). The inocula of AM fungi can be applied as spores, or fragments of colonized roots. The spores and hyphae can be isolated from the soil rhizosphere and mixed with carrier substrates (Gentili & Jumpponen, 2006). Spore inocula are the most resistant and can survive unfavorable environmental conditions for a long period, but they colonize new root systems more slowly than other preparations. Therefore, both types of inocula, for example, spores and fragments of colonized roots, should be combined in commercial products (Marin, 2006).

Root-based bulk inoculum production technology utilizes mass-produced seedlings grown in sterilized soil infected with selected AM fungi using spores from fruiting bodies from cultivated plants. This technology results in seedlings with infected root systems, and the roots and adhering soil are chopped up and used as the starter inoculum for scale-up production. The inocula are produced in bulk by infecting fresh seedling of selected plants (Gentili & Jumpponen, 2006). The root inocula are kept in polythene bags and used for pelleting seeds or in the preparation of granules for seed bed inoculation (Singh & Tilak, 2002). The other methods such as soil-free aeroponic, nutrient film, and root organ culture system have been used for the production of AM, but these methods are costly and preclude commercial mass production (Gentili & Jumpponen, 2006). It may be possible to mass produce plants in tissue culture in sterile agar media and induce mycorrhizal associations using spores from fruiting bodies of selected mycorrhizal fungi. The dried root tissues and fungal mycelia could then be developed into mycorrhizal seeding products.

Some steps are essential for development of a commercial fungal biofertilizers. They include selection, large-scale production, carrier selection and preparation, mixing and curing, maintenance of appropriate numbers of inocula, and strong quality control (Malik et al., 2005). The criteria for selecting AM fungi will depend on details of the local environment, soil conditions, and host plants. The AM fungi must (1) colonize roots rapidly after inoculation, (2) absorb phosphate from the soil, (3) transfer phosphorus to the plant, (4) increase plant growth, (5) persist in soil and re-establish mycorrhizal symbiosis during the following seasons, and (6) form propagules that remain viable during and after inoculum production (Adholeya, 2006).

The success of a formulation depends on whether it (1) is economically viable to produce, (2) does not alter the viability and function of the inoculum, and (3) is easy to carry and enhance dispersal during application. The inoculum formulation may comprise one or more AM fungi and other organisms that together enhance the ability of the inoculum to form mycorrhizal associations with the target plant.

The formulations are available in the form of powder, tablets/pellets or granules, gel beads, and balls (Adholeya et al., 2005). There are many ways to apply the AM inocula (Adholeya et al., 2005; Schwartz et al., 2006), including scattering by hand, in-furrow application, seed coating, root dipping, and seedling inoculation. The efficacy of the application of AM inocula depends on the product, environmental condition, delivery method, and other variables. The success of AM fungi inoculation depends on crop species, size and effectiveness of indigenous AM fungi populations, fertility of the soil, and cultural practices (Adholeya et al., 2005). The production of commercial mycorrhizal inoculum has evolved considerably, with the availability of various types of microbial cultures and inoculants in the market today that are rapidly increasing because of the technological advances (Raja, 2006). A simple flow chart illustrating biofertilizer's production is shown in Fig. 14.7.

There are more than 30 companies worldwide marketing mycorrhiza products (Tables 14.5 and 14.6) comprising one or multiple mycorrhizal fungal inocula. These products are plant growth promoters and are to be used in horticulture, agriculture, restoration, and forestry (Schwartz et al., 2006).

Other fungi used as biofertilizers (Table 14.4) are discussed below.

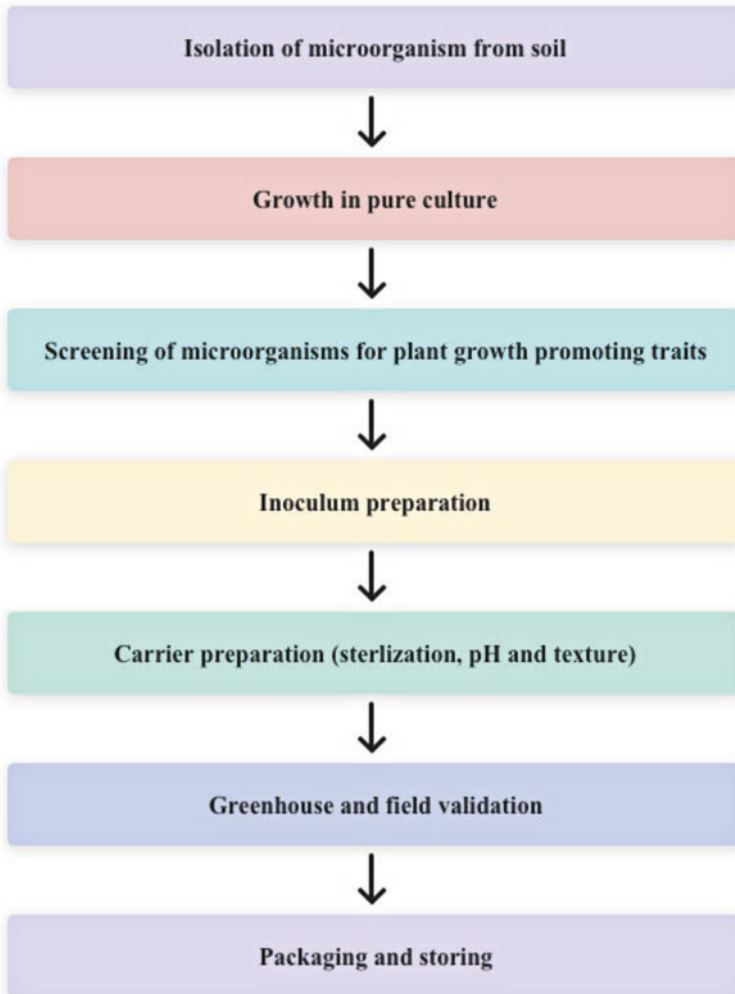


Fig. 14.7 Simplified flow chart for the production of biofertilizer

14.4.1 *Penicillium spp.*

Penicillium spp. are phosphorus-solubilizing agents that improve phosphorus absorption in plants and stimulate plant growth (Wakelin et al., 2004; Pradhan & Sukla, 2005). They act as biofertilizers by enriching compost and soil quality (Sharma et al., 2013). Some of the important species in this genus are *P. bilaji*, *P. italicum*, *P. simplicissimum*, *P. oxalicum*, *P. frequentans*, and *P. rubrum* (Yadav et al., 2018b). They also act as zinc solubilizers (Anitha et al., 2015; Ghosh et al., 2019).

Table 14.5 Contribution of arbuscular mycorrhizal fungi to plant growth promotion and soil nutrients

| Mycorrhizal fungi | Plants | Effect on plant | Effect on soil | References |
|---|---------------------------|--|---|------------------------------|
| <i>Glomus versiforme</i> <i>Glomus mosseae</i> | Tomato | Promotes growth and yield under water stress and more efficient conditions | Increases phosphorus concentration in the soil | Maaloum et al. (2020) |
| <i>Glomus etunicatum</i> | Maize | Improves chlorophyll content and nutrient uptake in maize | Increases soil quality | Xu et al. (2019) |
| <i>Acaulospora lacunosa</i> | Strawberry | Enhances nutrient uptake in strawberry | Increases soil nutrient for horticultural crops productivity | Chiomento et al. (2019) |
| <i>Rhizophagus irregularis</i> | Wheat | Improves tolerance to stress, enhances plant growth, and increases seed yield | Increases soil nutrient needed for wheat production | Tohidi et al. (2018) |
| <i>R. irregularis</i> | Maize | Enhances tolerance to salt stress and improves growth parameters | Reduces the concentration of salt in the soil for better plant development | Krishnamoorthy et al. (2016) |
| <i>G. mosseae</i> and <i>G. geosporus</i> | Strawberry | Enhances growth and improves its tolerance to water stress | Increases soil nutrient to enhance its colonization on the plant root system | Boyer et al. (2015) |
| <i>Rhizophagus irregularis</i> | Tomato | Protects plants against pathogens (<i>Sclerotinia sclerotiorum</i>) and improves nutrient uptake in plants | Increases soil micronutrient and triggers the defense of the plant against pathogens | Mora-Romero et al. (2015) |
| <i>Glomus deserticola</i> | Snapdragon | Increases the total dry matter, chlorophyll content, and improves Snapdragon tolerance to water stress | Increases soil nutrients needed for plant growth promotion | Tognon et al. (2016) |
| <i>Glomus</i> spp. and <i>Mortierella</i> spp. | Seashore mallow | Increases shoot and root weight under salt stress | Increases soil nutrient and enhances its absorption by plants | Mota et al. (2019) |
| <i>Glomus versiforme</i> | <i>Mentha arvensis</i> L. | Increases dry weight and improves nutrient uptake in salt stress conditions | Increases soil nutrient and enhances its absorption by the plant to enhance its tolerance to salinity | Bharti et al. (2016) |

Table 14.6 Fungal biofertilizers available in India

| Biofertilizer | Active ingredient | Mode of action | Dose | Manufacturer |
|---|--|--|--|---|
| A. Phosphate-solubilizing microorganisms | | | | |
| Grotop PSB Powder | Phosphate-solubilizing microorganisms (<i>Bacillus</i> sp.), powder 107–109 cfu g ⁻¹ and liquid 109 cfu ml ⁻¹ | Solubilize unavailable organic and inorganic forms of phosphorus (80%). | Seed: 5–10 g kg ⁻¹ seed Soil: 0.5–1 kg acre ⁻¹ along with 40–50 kg manure Foliar: 3 ml l ⁻¹ water | MD Biocoals Pvt. Ltd., Haryana |
| Mani Dharma's biopromoter | <i>Bacillus megaterium</i> + <i>Aspergillus niger</i> | Biopromoter facilitates root formation and plant growth. It improves soil quality with subsequent uses. It should not be mixed with antibacterial agents and inorganic fertilizers. | Foliar: 200 g/200 ml ⁻¹ of "rice kanji" or 5% jaggery water | Mani Dharma Biotech Private Limited, Tamil Nadu |
| Multiplex Nalapak | Homogenous mixture of <i>Azotobacter</i> + <i>Azospirillum</i> + phosphate solubilizer + potash mobilizer | It produces amino acids, vitamins, and growth-promoting substances like IAA, GA, and cytokines, which helps in better growth and development of crop plants. Improves physical, chemical, and biological properties of the soil. | Soil: 500 ml 5 kg ⁻¹ acre ⁻¹ along with 100 kg well decomposed FYM Foliar: 10 g l ⁻¹ water | Multiplex Bio-Tech Pvt. Ltd., Karnataka |
| Ambiphos | Phosphate-solubilizing microorganism (<i>Aspergillus niger</i>) | PSM secrete organic acids, which dissolve unavailable phosphate into soluble form and make it available to the plants. | Foliar: 3–5 ml l ⁻¹ water | Ambika Biotech & Agro Services, Madhya Pradesh, |

(continued)

Table 14.6 (continued)

| Biofertilizer | Active ingredient | Mode of action | Dose | Manufacturer |
|---------------|---|--|--|--|
| Biophos | <i>Bacillus megaterium</i> var. <i>phosphaticum</i> | PSM secrete organic acids, which dissolve unavailable phosphate into soluble form and make it available to the plants. | Seed: 5–10 ml kg ⁻¹ of seed Seedling: 125–250 ml in 25–50 liter of water as coating, sett treatment 125–250 ml in 60–80 l ha ⁻¹ for 30 min Soil: 500–625 ml ha ⁻¹ mixing with 250–375 kg FYM | Biotech International Limited, Delhi |
| BioP-P | Phosphate-solubilizing microorganism (2 × 10 ⁸ CFU g ⁻¹) | PSM secrete organic acids, which dissolve unavailable phosphate into soluble form and make it available to the plants. | Foliar: 3–5 ml l ⁻¹ water | Sundaram Overseas Cooperation, Gujarat |
| PSM | Phosphate-solubilizing microorganisms | PSM secrete organic acids, which dissolve unavailable phosphate into soluble form and make it available to the plants. | Soil: 0.5–1 kg acre ⁻¹ along with 40–50 kg FYM | Shree Biocare India, Shree Biocare Solution Pvt Ltd, Gujarat |
| PSM | Phosphate-solubilizing microorganisms | PSM secrete organic acids, which dissolve unavailable phosphate into soluble form and make it available to the plants. | Soil: 0.5–1 kg ha ⁻¹ a | KRIBHCO, UP |

(continued)

Table 14.6 (continued)

| Biofertilizer | Active ingredient | Mode of action | Dose | Manufacturer |
|--|--|--|---|--|
| B. Enriched compost | | | | |
| Multiplex Sagar (compost poly culture) | Homogenous mixture of <i>Azospirillum</i> + <i>Trichoderma</i> + <i>Pleurotus</i> | Nitrogen-fixing potential, biopesticidal activity, enhances compost degradation. | Soil: 1 kg acre ⁻¹ along with 1000 kg well-decomposed FYM Foliar: 100 ml 10 l ⁻¹ water | Multiplex Bio-Tech Pvt. Ltd., Karnataka |
| Enriched compost culture | <i>Trichoderma harzianum</i> + <i>Aspergillus</i> + <i>Penicillium</i> | Enhances compost degradation F. | Foliar: 100 ml 10 l ⁻¹ water | Organic Biotech Pvt Limited, Maharashtra |
| Biomanure culture | <i>Trichoderma harzianum</i> + <i>Aspergillus</i> | Enhances compost degradation F. | Foliar: 100 ml 10 l ⁻¹ water | Uno Natural and Greens Private Limited, Tamil Nadu |
| LignoBiocompost culture | <i>Trichoderma resei</i> , <i>Phanerochaete chrysosporium</i> , and <i>Aspergillus awamori</i> | Enhances compost degradation. | Foliar: 75–100 ml 10 l ⁻¹ water | Peak Chemical Industries Limited, West Bengal |
| C. Phosphate-mobilizing biofertilizer | | | | |
| ManiDharma VAM | Vesicular arbuscular endomycorrhiza (VAM) | Solubilize phosphate and supplies P, Zn, Mn, Fe, Cu, Co, and Mo to the plants. Increases the plant vigor by inducing drought resistance in young seedlings. Protects the plants from the fungal pathogens. | Soil: 200 g/m ² or 2–5 g seedling ⁻¹ ; 50–200 g trees ⁻¹ ; 3–5 kg acre ⁻¹ (2–3 cm depth). | Mani Dharma Biotech Pvt Ltd, Tamil Nadu |
| Ecorhiza-VAM/ Nurserrhiza-VAM | Arbuscular mycorrhiza | Improved uptake of nutrients, root development, and growth in plant. | Soil: 200 g/m ² or 2–5 g seedling ⁻¹ ; 50–200 g trees ⁻¹ ; 3–5 kg acre ⁻¹ (2–3 cm depth) | Mani Dharma Biotech Pvt Ltd, Tamil Nadu |
| Ecorhiza-VAM/ Nurserrhiza-VAM | Arbuscular mycorrhiza | Improved uptake of nutrients, root development, and growth in plant. | Soil: 3–5 kg acre ⁻¹ with the 200–250 kg FYM, one tablet plant ⁻¹ in 2–4 inches deep hole near the plant root | TERI, New Delhi |

(continued)

Table 14.6 (continued)

| Biofertilizer | Active ingredient | Mode of action | Dose | Manufacturer |
|--------------------------|---|--|--|---|
| Root care | Mycorrhiza (<i>Glomus intraradices</i>) | Improved plant health, uptake of nutrients and reduction of environmental stress. | Soil: 5 kg acre ⁻¹ | Ambica Biotech, MP |
| Mycorrhizae | 105 propagules/kg with carrier material (talc powder/ vermiculite) | Mobilize major nutrients like phosphorus and potassium and certain micronutrients like zinc, calcium, etc. | Soil: 5–10 kg of VAM ha ⁻¹ is recommended along with the 1 q of FYM; 25 g plant ⁻¹ | Dr. Rajan Laboratories, Tamil Nadu |
| JOSH Super/ JOSH Plus | Mycorrhizal (<i>Glomus intraradices</i>) | Root development in plant enhances growth. | Soil: 5 kg acre ⁻¹ for all crops, 60 infective propagules g ⁻¹ | Cadila Pharmaceuticals Limited, Gujarat |
| Shubhodaya | Vesicular arbuscular mycorrhiza (with three species: 1. isolated from desert and adoptable to harsh water-strained condition; 2. adoptable to water-lodging conditions; and 3. adaptable to general and acidic/basic soil conditions) | Improved uptake of nutrients. | Soil: 5–10 kg acre ⁻¹ | Cosme Biotech, Goa |
| TARI VAM | Vesicular arbuscular mycorrhiza | Mobilize nutrients like, phosphorus and zinc. | Soil: 5–10 kg acre ⁻¹ | TARI Biotech, Tamil Nadu |
| BioCarry | Vesicular arbuscular mycorrhiza (VAM) | Helps in efficient mobilization and uptake of fertilizers and other nutrients by plants. | Soil: 5 kg acre ⁻¹ | Sundaram Overseas Operation, Gujarat |
| Symbion VAM | Solid formulations of arbuscular mycorrhiza | Improved uptake of nutrients, root development in plant and growth S. | Soil: 5 kg acre ⁻¹ | T. Stanes and Company Limited, Tamil Nadu |

(continued)

Table 14.6 (continued)

| Biofertilizer | Active ingredient | Mode of action | Dose | Manufacturer |
|-----------------------------|--|--|--|--|
| CAMBAY's VARDHAK | Arbuscular mycorrhiza (powder and tablets) | Helps in efficient mobilization of nutrients. | Soil: 5 kg acre ⁻¹ ; one tablet plant ⁻¹ | Neesa Agritech Private Limited |
| Mycorrhiza–VAM | Arbuscular mycorrhiza | Mobilize nutrients like P, Zn, Cu, and B. | Soil: 10 kg acre ⁻¹ | KCP Sugar and Industries Corporation Ltd |
| Mycorrhiza–AM Biofertilizer | Arbuscular mycorrhiza | Helps in efficient mobilization of nutrients and improved plant growth. | Soil: 5 kg acre ⁻¹ | Majestic Agronomics Pvt. Ltd., HP |
| Colonizer | Arbuscular mycorrhiza (powder) | Colonizes living plants roots. Improves phosphorus uptake and imparts stress and disease resistance. | Soil: 2–3 kg acre ⁻¹ | Krishidhan Seeds Pvt. Ltd, Maharashtra |

Penicillium simplicissimum and *Penicillium* sp. in combination with *Aspergillus awamori* are used as phosphate-mobilizing biofertilizers. Mishra et al. (2013) in their finding suggested that a combination of *Penicillium* and *Trichoderma* increases the development of wheat and soybean under nursery condition. *Penicillium bilaiae* was formulated as a commercial product named Jumpstart® (wetttable powder) and has been found to enhance accumulation of phosphorus (P) and increase dry matter and seed yield in canola (*Brassica napus*) (Wakelin et al., 2004; Sane & Mehta, 2015; Chandrashekrappa & Basalingappa, 2018; Whitelaw et al., 1999). *Penicillium radicum*, isolated from the rhizosphere of wheat roots, has shown a good promise in plant growth promotion (Whitelaw et al., 1999), while *P. italicum* from the rhizosphere soil has the ability to solubilize tricalcium phosphate (TCP) and promote efficient soybean growth and production (Ram et al., 2015). In general, *Penicillium* spp. serves as phosphate solubilizer and general plant growth stimulator (Pal et al., 2006).

14.4.2 *Aspergillus* spp.

Aspergillus spp. helps in unleashing the soil phosphate complexes through the synthesis of an organic acid, which dissolves the soil phosphate for plant uptake (Zarafi & Dauda, 2019). A good example is *Aspergillus niger*, which facilitates plant growth through root formation (biopromoter). It improves soil quality and can be used to produce zinc-solubilizing biofertilizers. An example of the biofertilizer

made from *Aspergillus niger* includes “Ambiphos.” It secretes organic acids, which aid in dissolving unavailable phosphate into a soluble form and makes it available for plants use. In addition, “Ligno Biocom” post culture is also a good biofertilizer made from *Aspergillus awamori*, which serves as an enhancer for compost degradation (Pal et al., 2006). Most importantly, these fungi genera (*A. flavus*, *A. tubingen-sis*, *A. awamori*, *A. terreus*, *A. fumigates*, *A. niger*, and *A. melleus*) are able to solubilize inorganic phosphate through the production of citric acid, gluconic acid, glycolic acid, oxalic acids, and succinic acid (Barroso et al., 2006; Akintokun et al., 2007). *Aspergillus fumigatus*, which most often isolated from compost, has been reported to be a potassium-releasing fungus (Barroso et al., 2006; Akintokun et al., 2007; Sharma et al., 2013). *Aspergillus* spp. are also considered as microphos biofertilizers because of their ability to release phosphate from the bounded and insoluble state (Jehangir et al., 2017).

14.4.3 *Chaetomium* spp.

Chaetomium species are found mostly in soil and organic compost. It is a mycofungicide agent used mostly for protective and curative purposes. According to Dhingra et al. (2003), *Ch. bostrychodes* and *Ch. olivaceum* have been used severally as biofertilizers for compost enrichment. Most interesting is that the product of *Chaetomium* species has in recent time been used as fungal biofertilizers. A good example is seen in Ketomium, a bioproduct formulated from *Ch. globosum* and *Ch. cupreum*, which serves as plant growth stimulants (Jehangir et al., 2017). As a unique strategy, *Chaetomium* spp. have developed the ability to enable the suppression of bacterial and other fungal growth through competition, mycoparasitism, and antibiosis (Marwah et al., 2007). *Chaetomium globosum* and *Ch. cupreum* most importantly control root rot disease in black pepper, citrus, soybean stem, and strawberry with tendency of reducing sugar beet damping-off diseases (Dhingra et al., 2003; Zarafi & Dauda, 2019). Tomato, corn, rice, pepper, citrus, durian, birds of paradise, and carnation treated with Ketomium® have a greater plant growth and high yields than nontreated plants (Soytong et al., 2001).

14.4.4 *Gliocladium* spp.

Gliocladium species are common soil saprobes. They are parasites to many plant pathogens such as *Sporidesmium sclerotiorum* and *Fusarium* spp. They attack the fungal host by direct hyphal contact and forms pseudo appressoria. Engineered strain of *Gliocladium catenulatum* (strain JI446) has also been used as a wettable powder applied in soils, roots, and foliage to reduce the incidence of damping-off disease in the greenhouse (Viterbo et al., 2007; Youssef & Eissa, 2014). *Gliocladium virens* has been used as a biological control agent against a wide range of soil-borne

pathogens such as *Pythium* and *Rhizoctonia* under greenhouse and field conditions. They also produce antibiotic metabolites such as gliotoxin, which has antibacterial, antifungal, antiviral, and antitumor activities. A good example is *Gliocladium catenulatum*-JI446, a commercialized bioagent that has been used to reduce the impact of damping-off disease caused by *Pythium ultimum* and *Rhizoctonia solani* (Kaewchai et al., 2009; Nissipaul et al., 2017).

14.4.5 *Trichoderma* spp.

Trichoderma is a fungal genus commonly found in most habitats. They are predominantly found in root and soil ecosystems and considered as ubiquitous saprobes. *Trichoderma* spp. can be easily isolated from decaying wood soil and organic materials (Rosa et al., 2012). Studies have shown a number of successfully reduced plant diseases through their highly effective antagonistic and mycoparasitic activity (Kour et al., 2019; Sharma et al., 2019; Verma et al., 2017). Some *Trichoderma* strains interact with their plant's host, increasing directly their growth potential and imposing resistance against diseases and abiotic stresses (Rosa et al., 2012). Some of the fungal disease-causing agents acted upon by *Trichoderma* (*Trichoderma harzianum* T-22) include *Rhizoctonia* spp., *Pythium* spp., *Botrytis cinerea*, *Pythium*, *Rhizoctonia*, and *Fusarium* spp. As an agent with mycofungicides properties, *Trichoderma* spp. has been engineered overtime to successfully support crops such as potato, corn, tomato, peanut, beans, cotton, and soybean (Sneha et al., 2018). They are considered an excellent competitor within the rhizosphere because they exercise resistance against soil unfavorable conditions and high efficiency in harnessing soil nutrients while also being hostile to phytopathogens (Chandrashekrappa & Basalingappa, 2018).

Members of the fungal genus *Trichoderma* are found to establish beneficial interactions with plants either antagonizing phytopathogens or directly influencing morphogenesis (Benítez et al., 2004; Harman et al., 2004; Harman, 2006; Hermosa et al., 2012). Furthermore, *Trichoderma* induces plant defense responses and improves crop performance under different stress conditions (Rawat et al., 2013; Hashem et al., 2014).

Mastouri et al. (2010) demonstrated that, when tomato seeds were treated with *Trichoderma harzianum* strain T-22 (T22), a range of biotic and abiotic stresses were alleviated. This is because after the colonization of roots *Trichoderma* spp. are able to chemically communicate with the plant and, sometimes, they act as endophytic symbionts. They are thus capable of correctly altering the expression of various plant genes and, as a consequence, plant physiology (Harman et al., 2012).

Recently, the use of the T22 strain of *T. harzianum* was also investigated as a new approach for controlling viruses, considering that chemical treatments have no effect on these pathogens (Vitti et al., 2015).

Trichoderma species improve mineral uptake, release minerals from soil and organic matter, enhance plant hormone production, induce systematic resistance

mechanisms, and induce root systems in hydroponics (Ali et al., 2021). For these reasons, *Trichoderma* species are known as plant growth-promoting fungi. *Trichoderma* species have therefore successfully been used as biofungicides and biofertilizers in greenhouse and field plant production (Harman et al., 2012; Harman, 2006; Hermosa et al., 2012). These studies emphasize that *Trichoderma* can be widely applied as important sources of antibiotics, enzymes, decomposers, biopesticides, and plant growth promoters in agricultural practices. There are many *Trichoderma* products as fungal biofertilizers available in the market (some of them are shown in Table 14.6).

14.4.6 Formulation of Fungal Biofertilizers

Fungal biofertilizer is formulated either in liquid or solid state. Often, these bio-products are packaged as dry biomass (dusts, granules, and briquettes) and suspension (water-based and emulsions). Dust particle size ranges from 5 to 20 μm and contains <10% of an organism by weight. Granules are discrete masses, 5–10 mm^3 in size. Pellets are >10 mm^3 , and briquettes have large blocks up to several cubic centimeters. The granules are discrete masses contain an inert carrier like charcoal, lignite, clay minerals (vermiculite, bentonite), starch polymers, dry fertilizers, and ground plant residues. In some cases, the choice of carrier depends on absorption, hardness, bulk density, and product disintegration rate in water. During formulation, these microbial bioagents get coated with various materials to slow and control the rate of their release. In the solid-based state, however, their shelf-life is about 6 months as they are not tolerant to UV rays and temperatures greater than 30 °C. Usually, the population density of these microbes ranging from 10⁸ CFU/ml reduces with time (10⁶ CFU/mL) at fourth month, reaching almost nil at the end of sixth month. Again, the improper sterilization of carrier material and their handling such as mixing the organism with carrier and packing serve as source of contaminations. Because of this, the inoculant packet could not hold desired biofertilizer organism for longer time and has not become effective and popular among the farmers in some climes. Furthermore, the quality of the biofertilizers gets deteriorated with longer and unsafe storage conditions. These have in many ways contributed to the limitation associated with biofertilizer application and in some cases where it could not be able to give viable results in the field. Owing to these challenges, the liquid formulation of organisms with a count of 10⁹ CFU/ml (*Azospirillum* and *Phosphobacteria*) was developed to avoid drawbacks of biofertilizers and to increase the quality and shelf-life of bioinoculants. Here, preservatives are added with the microbial cultures and packed in bottles with improved shelf-life of about 2 years. According to Pal et al. (2015), the application of 1 ml of liquid biofertilizers is equivalent to the application of 1 kg of 5-month-old carrier-based biofertilizers. Increasing research has often suggested amendment of carrier materials to increase effectiveness of the biofertilizers (Itelima et al., 2018). These carriers are the delivery vehicle of live biofertilizers from the factory to the field that upon application

Table 14.7 Classification of carrier materials for the production of biofertilizer

| Categories of carrier material | Carrier materials |
|--------------------------------|--|
| Natural materials | Peat, lignite, coal, clay, and organic soil |
| Inert materials | Talc, vermiculite, perlite kaolin, bentonite, silicate, rock phosphate, calcium sulfate, and zeolite |
| Synthetic polymers | Polyacrylamide, polystyrene, and polyurethane |
| Natural polymers | Xanthan gum, carrageenan, agar agar, and agarose |
| Organic materials | Charcoal, biochar, composts, farmyard manure, sawdust, maize straw, vermicompost, cow dung, corn cob, and wheat husk |
| Agro-industry by-product | Sludge ash, jaggery |

turns into organic, inorganic, and/or synthetic materials in the environment (Mahanty et al., 2016). Bashan et al. (2014) in their work reported that carriers have tremendous importance in delivering the right number of viable cells in good physiological condition. It also provides a short-term protective niche to the biofertilizers in soil, either physically via the provision of a protective surface of pore space or nutritionally via the provision of a specific substrate (Arora et al., 2010); hence, a good carrier should be nontoxic, of high moisture absorption capability, easy to process, and free of lump-forming materials, easy to sterilize, available in adequate amounts, and cost-effective with good pH buffering capacity (Mahanty et al., 2016). Different carrier materials used for formulating biofertilizers are shown in Table 14.7.

14.4.7 Mode of Application of Biofertilizers

Given below, are several ways that biofertilizers can be applied:

- Seed inoculation with powder formulations
- Dry biofertilizers mixed with the seeds in the seed hopper
- Sprinkle method (a small amount of water mixed with seeds before peat powder is added and mixed)
- Slurry method (the biofertilizer is suspended in water then added to the seeds and mixed)
- Seed pelleting
- Biofertilizers and adhesive are applied as slurry to seeds and coated with ground material like lime
- Peat suspension in water sprayed into the furrow during sowing

14.5 Role of Fungal Biofertilizer in Sustainable Agro-Practices

The use of fungal biofertilizers in sustainable agriculture has been gaining prominence in recent times, not only because it allows for effective utilization of mineral elements such as nitrogen and phosphorus, but also enhances mineral acquisition, transportation, as well as acting as an efficient biocontrol agent. The use of fungi as agents to improve soil fertility, water uptake, nutrient availability, induce environmental stress tolerance, biocontrol, as well as in limiting the use of agrochemicals has long been identified (Odoh et al., 2019a; Figueiredo et al., 2017). In a nutshell, the potentiality of fungal biofertilizers can be attributed to its unique functional niche in the agroecosystem, which are as follows:

- The fungal mycorrhizal symbiosis is arguably the most important symbiosis because of its agricultural importance (Yadav et al., 2019). Fundamentally, these plant species form mycorrhizal associations by acting as a critical linkage between soil and plant roots. This association is usually characterized by the movement of fungal-acquired nutrients to plants and the movement of plant-produced carbon to fungus (Johri et al., 2015). The fungal mycelium that extends from the root surfaces into the soil matrix captures nutrients from soil solution while the minuscule diameter of the fungal hyphae increases the surface area for absorption, thus increasing its ability for nutrient acquisition and utilization. Interestingly, these occur when mycorrhizal fungi colonize plant root systems; just as seen in the endo (arbuscular), ecto, ectendo, arbutoid, ericoid, monotropoid, and orchidaceous mycorrhizae. Pal et al. (2014) noted that when mycorrhizae are used as biofertilizers they enhance uptake of phosphate, zinc, and water, leading to improved hardiness and uniform growth of transplanted stock. The arbuscular mycorrhizal fungi also improve soil structure and control plant diseases, thus implying that plants colonized by these microbes grow better than those without them (Tripathi et al., 2017). Arbuscular mycorrhizal fungi (AMF), a group of soil fungi that live in symbiosis with most agricultural crops, have the ability to mobilize mineral nutrients from soil for their plant hosts.
- Fungal agents act as inhibitors or antagonizers of phytopathogens (biocontrollers), promote plant growth (biostimulants), activate mineral nutrients, and fix atmospheric nitrogen (biofertilizers) as part of sustainable agricultural practices (Odoh et al., 2019a).
- Many fungal bioinoculants such as endophytic fungi (*Penicillium*, *Aspergillus*, *Piriformospora*, and *Curvularia*) solubilize and mineralize phosphorus (Rai et al., 2013). However, for phosphate solubilization, endophytes are the major contributors. Here, they colonize plants without inducing symptoms of disease, making them a more aggressive colonizer of plant root than non-endophytic microbes. These endophytes convert insoluble soil phosphate into soluble forms through the secretion of organic acids with concomitant changes in the texture, structure, and water-holding capacity of the soil.

- *Trichoderma* biofertilizers (Zhang et al., 2018) were shown to increase soil anti-fungal compounds that suppressed pathogenic fungi and allowed for improved grassland biomass. This suggests that *Trichoderma* biofertilizers could be an important tool for sustainable management of soil and plant productivity.

14.5.1 Fungal Biofertilizers as an Alternative to Chemical Fertilizers

Fungal biofertilizers are a renewable source of plant rhizospheric nutrients that could serve as a close alternative to chemical fertilizers (Kannaiyan, 2002). The use of fungal biofertilizers can help to curb the excessive use of synthetic fertilizers, which when used indiscriminately causes detrimental effects to the environment and human health (Odoh et al., 2019b; López-Bucio et al., 2015; Yadav et al., 2020c). Fungal biofertilizers include plant growth-stimulating fungi (*Trichoderma*), mycorrhizal fungi (ectomycorrhiza and arbuscular mycorrhizae), enzyme-producing fungi for compost production, and P/K-solubilizing fungi. Mycorrhiza fungal biofertilizers, also known as vesicular arbuscular mycorrhiza (VAM), help in retaining moisture around the root region and increase resistivity towards different root and soil pathogens unlike the chemical fertilizers (Sadhana, 2014).

14.5.2 Fungal Biofertilizers Enhance Mineral Acquisition and Transport

The primary goal of AMF inoculation is to increase and enhance mineral acquisition and transport. Clark (1997) in his work observed increased plant acquisition of mineral nutrients upon AMF addition. During seasonal variation (dry season), soil structure and porosity determine soil water retention ability. As a result, this agent “AMF” influences hormonal flow and changes information from plant roots to shoots, and affects stomatal responses when soil water potential is lowered (Odoh et al., 2019a, b). In the work of Jagnaseni et al. (2016), they maintained that mycorrhizal associations increase nutrient uptake during water stress through hydraulic conductivity in the roots. They occur as the hyphal sheath, Hartig net, and extraradical mycelium. These fungal mantles represent a significant apoplastic barrier creating a closed interfacial apoplast, thus enabling nutrient transfer (Bücking & Kafle, 2015). Upon application, the differential expression of plant and fungal uptake transporters in the mycorrhizal interface plays a role in the development of a strong concentration gradient across the mycorrhizal interface. High-affinity phosphorus (P) and nitrogen (N) transporters of ECM and AM fungi are expressed in the ericoid mycorrhiza (ERM), but downregulated in the intraradical mycelium (IRM). This favors the active absorption of nutrients by the ERM, but reduces nutrients

reabsorption process in IRM (Bücking & Kafle, 2015). Consequently, AM and ECM fungi store significant amounts of P as poly P where they are assumed to play an important role in the transport of P and N from the ERM to the IRM (Bücking & Kafle, 2015), unlike in hydrolysis of poly P where IRM would release P_i and Arg into the fungal cytoplasm to facilitate the efflux catabolic product of the urea cycle into the mycorrhizal interface. In general, AM and ECM fungi regulate nutrient transport in host cells via accumulation of poly P while inducing carbon supply in host plant, a process that also triggers poly P hydrolysis (Bücking & Kafle, 2015).

14.5.3 Fungal Biofertilizers Enhanced Nitrogen Fixation

Soil nitrogen is considered a valuable nutrient for plant growth. It is also one of the most essential micronutrients for food production. They exist in the form of ammonium, nitrate, and amino acids. Bücking and Kafle (2015) in their research reported that the ERM of AM and ECM fungi takes up inorganic N sources (ammonium NH_4^+ or nitrate NO_3^- from the soil). Through protein transporter named AMT1, nitrogen is taken up by plants as ammonium. During glutamine synthetase formation, this ammonium in combination with glutamate forms glutamine in the extraradical mycelium. Interestingly, NH_4^+ has been described as the preferred N source of mycorrhizal fungi because its uptake is energetically more efficient than the uptake of NO_3^- . In *Hebeloma cylindrosporum*, AMT1 and AMT2 are expressed as NH_4^+ transporters and regulated by exogenous NH_4^+ supply. The expression of both transporters is upregulated under low NH_4^+ supply conditions, but downregulated in response to an exogenous supply of NH_4^+ . More importantly, low-affinity NH_4^+ transporter (AMT3) also enables the fungus to maintain a basal level of N uptake and assimilation at high exogenous supply conditions (Bücking & Kafle, 2015).

14.5.4 Fungal Biofertilizers Enhanced Phosphorus Content

Phosphorus (P) is an extremely immobile element present in the soil. It is very vital for plant growth. One of the major roles of vascular–arbuscular (VA) fungi is the supply of phosphorus to plant roots via phosphate transporters in the hyphal membrane (Rastegari et al., 2020). This enables the fungi to provide phosphorus as poly P pool to the plants. The mycorrhizal hyphae present outside the root of the host plant known as the networks of filamentous, extraradical hyphae of AM fungi help in the uptake of freely available phosphates. Generally, the fungal hyphae extend beyond the host root to enable greater soil volume contact for phosphate acquisition. This fungus (AM fungi) hydrolyzes organic phosphates to soluble forms for plant use. This uptake is done using the phosphate transporter of the Pht1 family. According to Vergara et al. (2019), this fungus has two separate uptake systems for phosphorus: (a) a high-affinity system that works against an electrochemical

potential gradient. This takes up the Pi from the soil via proton co-transport and (b) a low-affinity system that facilitates the diffusion of Pi across the fungal plasma membrane. The expression of these transporters is often regulated by responses to externally available P concentration and P demand of the fungus (Johri et al., 2015).

14.5.5 Fungal Biofertilizers as an Efficient Biocontrol and Bioremedial Agent

Biological control agents are referred to as organisms that suppress plant diseases. In other words, they are considered as the utilization of introduced or resident living organisms such as fungi to reduce pressure and effects exerted by plant pathogens or suppressing reproduction of their kinds (Yadav et al., 2020a). Various fungal biofertilizers possess this ability, for example, *Trichoderma harzianum* – a species with biocontrol potential against *Botrytis cinerea*, *Fusarium*, *Pythium*, and *Rhizoctonia* (Redda et al., 2018); *Chaetomium globosum* and *C. cupreum*, having biocontrol activity against root rot disease caused by *Fusarium*, *Phytophthora*, and *Pythium* (Aggarwal, 2015). Besides inducing biocontrol, mycorrhizal associations also protect plants against heavy metal toxicity (copper, zinc, iron, manganese, cadmium, nickel, etc.). Ectomycorrhizal fungi protect trees by accumulating and immobilizing these toxic metals in the mycorrhizal mantle. Also, this association enhances detoxification mechanisms through extracellular heavy metal chelation (e.g., glycoprotein glomalin), binding of heavy metals to rhizodermal cell walls, and distorting heavy metal uptake mechanism. In addition, fungal biofertilizers also act as biocontrol agents (BCA) by competing for nutrients and space or by producing metabolites that impede spore germination (fungistasis), kill the cells (antibiosis), or modify the rhizosphere (acidifying the soil) so as to prevent pathogens multiplication (Khokhar et al., 2012). In mycoparasitism, biocontrol may result from a direct interaction between the pathogen itself and the BCA as they rely on the recognition, binding, and enzymatic disruption of the host fungus cell wall (Abdallah et al., 2018).

14.6 Fungal Biofertilizers as an Agent of Pesticide Tolerance

The most concerning agroecological issue is the unintended detrimental effects of pesticides due to its extensive use for decades for crop protection and yield. The versatile biological mechanism and interaction of fungi with soil and plant microbiome assessed under pesticide application can pave the way to isolate and develop fungal strains with high tolerance for pesticides, making it most compatible in the integrated agricultural management. Since the application of pesticides is unavoidable in conventional agricultural practice, to develop effective biofertilizers, with both pesticide tolerance and plant growth-promoting (PGP) traits are of paramount significance. A few such examples are discussed below.

14.6.1 *Compatibility of Aspergillus niger Isolates with Pesticides*

One such example is *Aspergillus niger* Van Tieghem, which is a good phosphate solubilizer (Khan & Anwer, 2007), siderophore producer (Khan & Anwer, 2007), antagonistic to many pathogens (Khan & Anwer, 2007, 2008, 2018; Khan et al. 2006a, b), and does not produce mycotoxins and ribotoxin (Campbell, 1994). Interestingly, the fungus is reported to decrease aflatoxin contamination. It is xerophilic and grows at a wide range of temperatures (10–50 °C), pH (2.0–11.0), and osmolarity (from nearly pure water up to 34% salt), and can also improve its thermostability (Zhang et al., 2007). The fungus is extremely resistant to herbicides, fungicides, and pesticides at very high concentrations (Braud et al., 2006). In view of the effectiveness of *A. niger* for their metal tolerance (Ahmad et al., 2006), bio-sorption potential, and pesticides tolerance, it was considered desirable to identify more efficient soil isolates among the diverse *A. niger* aggregate collected from different crop fields of several districts in Uttar Pradesh (North India). The compatibility test of *A. niger* with different groups of pesticides varied with isolates (Geetha et al., 2016). Some strains of *A. niger* have shown to degrade endosulfan pesticides (Bhalerao & Puranik, 2007) and some of them have produced organophosphate-degrading enzymes (organophosphorus hydrolase) that split P-S bonds of organophosphate (Liu et al., 2001, 2004). *A. niger* has also been reported to degrade carbamates by hydrolyzing several N-methylcarbamate insecticides (Qing et al., 2006) and pyrethroids pesticides through a novel pyrethroid hydrolase from their cell extracts (Liang et al., 2005).

14.6.2 *Compatibility of Trichoderma Isolates with Pesticides*

Similarly, the compatibility of fungal biocontrol agents (*Trichoderma harzianum*, *Trichoderma virens*, and *Pochonia chlamydosporia*) was assessed with different concentration of six pesticides, viz., carbendazim, mancozeb, metalaxyl, captan, thiram, and nemacur commonly used by farmers in India for the control of soil-borne plant pathogens (Mohiddin & Khan, 2013). Their efficacy in pesticide tolerance broadened the use of these strains as biopesticides under integrated disease management for managing soil-borne plant pathogens.

Currently, *Trichoderma* spp. share almost 70% of fungal biocontrol agents (BCAs) market. *Trichoderma harzianum* is extensively used for its dual advantages of controlling several soil and seed-borne phytopathogens in addition to its capacity to act as an effective biofertilizer (Keswani et al., 2013). Competence of *Trichoderma* spp. to other fungal biocontrol agents owes to its relatively faster metabolic rates; secretion of antimicrobial secondary metabolites and physiological conformation are some of the key factors that contribute to its vigorous antagonistic effect in field conditions. On the other hand, *Trichoderma* spp. has also been exploited for

commercial production of enzymes like cellulases, hemicellulases, proteases, and α -1, 3-glucanase. This article (Keswani et al., 2013) advertises the commercial potential of *Trichoderma* spp. discussing its status in various aspects, viz., pest control, plant growth promotion, bioremediation, and production processes.

14.6.3 Compatibility of ECM with Pesticides: A Novel Path

Endeavors pertaining to application of intriguing ectomycorrhizal association for on-site pesticide remediation are acquiring huge consideration from global scientific community (Assad et al., 2020). These ectomycorrhizal (ECM) fungi represent a functional group of enormous importance, that interconnects plants through below-ground colossal hyphal networks and serve as potential indicators of changes in environmental quality. Owing to their specialized lifestyle and mycorrhizal remediation potential, ECM fungi remediate pesticides through enzymatic conversion of noxious pesticides into innocuous residues. As an innovative emerging discipline, ectomycorrhizal biotechnology engrosses application of subsist ECM fungal strains or their prominent fungal enzymes, via contemporary biotechnological approaches, for sustainable remediation and recuperation of gravely disturbed environs. This technology has persuasive potential to arise as an effectual substitute of conventional remediation system. Ectomycorrhizal ecology, mycorrhizoremediation of pesticides, fungal enzymes entailed in remediation, role of ectomycorrhizal biotechnology in pesticide remediation, and plant–pollutant–mycorrhiza interactions with reference to climate change are the novel paths to be explored.

14.6.4 Plant Growth-Promoting Microorganism (PGPM): Prospects and Constraints

Bioremediation and microbial degradation are being used for the destruction of chemicals in soil and ground water. These pollutants are removed from the soil by indigenous microorganisms, including bacteria for their extraordinary characteristic to use a wide range of xenobiotics as sole energy and carbon source (Siddique et al., 2003).

In other words, indigenous microorganism is also known as plant growth-promoting microorganism (PGPM), as represented by the microbial cells of bacteria, actinomycetes, and fungi. PGPM inoculants are presently commercialized as an innovative and efficient solution for plant growth enhancement through direct and indirect mechanisms.

Presently, the PGPM available in the market as the only single or dual microbial consortium is not effective for all the crops and exhibit the inability to degrade or tolerate pesticides. They have been sourced from other agroclimatic region, and that

is why they lose their potency to function and survive in the indigenous soil, with different edaphic and climatic conditions. This poses a huge problem with PGPM as biofertilizers, currently available in the market for farmer's application. In order to overcome these problems, the need of alternative practices like the effective pesticide-degrading plant growth-promoting microbial consortia are required for improving plant growth of vegetables and degrade or minimize the pesticides in soils. These consortia can degrade and utilize the pesticides for co-metabolism or as a sole source of carbon and enhance plant growth. Being indigenous in nature, they may easily grow and help in pesticide bioremediation in crop fields. Therefore, the future prospects lie in screening indigenous fungal strains for the development of efficient indigenous pesticide-tolerant plant growth-promoting microbial consortia from rhizospheric and non-rhizospheric soil.

Similar studies were undertaken with an intention to isolate and characterize the plant growth-promoting properties and pesticide (chlorpyrifos) tolerance ability of microbial strains for sustainable vegetable productions (Verma et al., 2016a, b). Fifty microbial isolates from vegetable field were successively tested on the basis of morphological, biochemical, and plant growth-promoting properties. Ultimately, three strains were found as effective pesticide-tolerant plant growth-promoting microbial consortia for vegetable production under sustainable agriculture.

According to results of microbial populations, fungal populations were recorded comparatively higher than actinomycetes and bacteria. Therefore, fungi are found to have higher resistance to pesticides (Ambethgar, 2009).

A similar study was conducted to screen for pesticide-resistant isolates of plant growth-promoting microorganisms such as *Azotobacter* spp., *Azospirillum* spp., *Rhizobium* spp., and phosphate solubilizers, the activity of which would not be affected by the presence of pesticides owing to their pesticide resistance, and to check their efficacy as biofertilizers. Screening was carried out using rhizospheric soil samples from fields treated with pesticides (Prabhu & Raut, 2011). Plant studies and viability studies proved that the isolates, mainly the consortium, not only proved to be effective in promoting plant growth in the presence of 1000 ppm of six commonly used pesticides, but also proliferated in pesticide-contaminated soils and improved the soil quality. These studies indicate that the isolates can be used as potential biofertilizers in pesticide-contaminated fields, thus bringing about enhanced plant growth even in the presence of pesticides.

Mishra and Sundari (2015) investigated the effects of direct inoculation of selected consortia on plants to study its effects on supporting plant growth in the presence of root pathogen *Sclerotium rolfsii* and organophosphate pesticides Malathion (ML) and Methyl Parathion (MP). Candidate plant growth-promoting microbial (PGPM) isolates chosen for the study are two bacterial isolates (PGPM2, a diazotrophic bacterium, PGPM9, a fluorescent Pseudomonad) and one fungal species (T103, a biocontrol fungus), originating from native agricultural fields of western U.P., India. Host plants inoculated with individual species showed a distributed growth enhancement pattern, while isolate T103 improved root biomass, isolate PGPM9 enriched photosynthetic pigment content and isolate PGPM2 expanded root and shoot lengths.

The results affirm the hypothesis that synergistic action of carefully selected PGPMS can escalate the benefits of plant growth promotion even in presence of pathogen and pesticide; hence, this consortium may be a valuable option for sustained plant growth in modern agriculture systems.

14.6.5 Developing Nation Through Developing Effective Biofertilizers

Shen et al. (2019) screened rice endophytic biofertilizers for fungicide tolerance and plant growth-promoting characteristics. Out of 17 different strains obtained from rice seedling roots, except for one, all strains could tolerate two or more fungicides. Rice roots inoculated with the endophytic bacteria conferred rice growth-promoting ability and should be considered a potential bacterial biofertilizer.

However, to develop effective biofertilizers, PGP traits (Mahanty et al., 2017) along with pesticide tolerance should be confirmed before their application because applying pesticides is inevitable in most agriculture practices (Ahemad & Khan, 2012). The hypothesis of such study should be based upon the application of biofertilizers with pesticide-tolerant ability to the soil, where they would be likely to retain their functions and would be capable of promoting the growth of the crops treated with pesticides.

Conclusively, all such studies suggest that it is important to examine the impact of pesticides during the process of developing biofertilizers with pesticide tolerance and plant growth-promoting characteristics. The rising demand for environmentally friendly, organic, and sustainable agricultural practices is driving the application on the use of beneficial biological products. Sustainability has become an integral component of the agriculture system. The use of fungi in agriculture sector is potentially useful for improved plant health and growth, water uptake, nutrient availability, stress tolerance, and biocontrol. Fungal species served as very important biological tools in sustainable agricultural ecosystem with the process of mycoremediation, mycocontrol–mycoherbicides, mycoinsecticides, and as mycorrhiza fungi (Thakur, 2020) and have potential role for attaining sustainable agricultural systems. Examples of fungi used as mycoremediators are *Pleurotus ostreatus*, *Rhizopus arrhizus*, *Phanerochaete chrysosporium*, *P. sordid*, *Trametes hirsute*, *T. versicolor*, *Lentinus edodes*, and *L. tigrinus*.

14.7 Biofertilizer Production: Indian Scenario

- Since the time the first commercial production of biofertilizer was initiated in 1956 at the Indian Agricultural Research Institute, New Delhi, and Agricultural College and Research Institute, Coimbatore, India has shown slow yet a steady growth in its production.

- The Ministry of Agriculture under the Ninth Plan initiated the real effort to popularize and promote the input with the setting up of the National Project on Development and Use of Biofertilizers (NPDB) during 1983–84.
- During 1965–1990, around 30 biofertilizer production laboratories were set up in the country to meet the demand, and a lot of schemes formulated to popularize their use in different legume crops. The 1990s saw a dramatic surge in biofertilizer industries with adding of new biofertilizers such as *Azotobacter*, *Azospirillum*, PSBs, and VAM (Adholeya, 2012) added to the list and total production jump was observed ten times in between 1989 to 2000 (Bhattacharya & Dwivedi, 2004).
- Starting from a few tons, production and consumption increased gradually and reached a moderate figure of 28,000 MT by 2009–10 (Fertiliser Association of India, 2011).
- Research in the field of microbial fertilizers is in progress in different research institutes of India. Several institutes of ICAR are producing commercial microbial fertilizers. However, further researches and encouragement from the government are needed to isolate and characterize more salt-tolerant microbial strains using modern biotechnological tools.
- The government should campaign for the utilization of these efficient and economic fertilizers because instead of considerable efforts by scientists, majority of the farmers in India are unaware of the use and fruitfulness of these microbial fertilizers.
- The share of bioinputs in agriculture is abysmally low. The market share is not even 1% in India. Relatively speaking there is more rigor in estimation of biofertilizer market in India because of the presence of some large producers in production of biofertilizers and comparatively this is more organized than other green inputs market (Rakshit & Bhadoria, 2002).
- Based on the gross cropped area in India and recommended doses of biofertilizers, potential demand is segregated into different categories of biofertilizer, such as *Rhizobium*, *Azotobacters*, *Azospirillum*, BGA, and phosphate solubilizer, where demands differ widely.
- Contrary to the world biofertilizer scenario, Indian biofertilizer industry is not restricted to *Rhizobium*. Presently phosphorous-solubilizing bacteria accounts for about 55% compared to other nitrogenous biofertilizers, which account for 45% (NCOF Annual Report, 2007–08).
- Although actual production and distribution of biofertilizers are below the targeted level, but there has been a positive trend in India, with respect to production.
- The growth rate in installed biofertilizer capacity is comparatively more stable than the growth rate in total production, consumption, and distribution.
- This is an indication that there is not only a need but also a role for market development for green agriculture inputs in India.
- In spite of the impressive growth rate of more than 200% in production capacity and around 300% growth rate experienced in production and consumption of biofertilizers in India, in the recent past yet it is only around 1.5% of the estimated demand potential for biofertilizers in the country.

- The region-wise distribution of biofertilizers is more dispersed in relation to chemical fertilizers across the country. As per the latest data available on biofertilizers, South Indian state Andhra Pradesh has overtaken another South Indian state Tamil Nadu in biofertilizer production to reach the top. The other major producers of biofertilizers are Karnataka, Kerala, Maharashtra, and Madhya Pradesh. The highest number of units is located in the state of Maharashtra.
- A good degree of large industrial chemical fertilizer units like IFFCO, KRIBHCO, TCL, MFL, NFL, and RCF are also involved in biofertilizer production as their corporate social responsibility along with chemicals-based fertilizers, insecticides, and pesticides. Among the PSUs, KRIBHCO produced highest amount followed by IFFCO, MFL, NFL, and RCF.
- Due to intensive and continued assistance from Government of India in this regard, overall production of biofertilizers in the country is continuing with a positive growth rate. However, the market for biofertilizers is still not well developed and the biofertilizer industry has not anticipated the growth.
- The current strategy of marketing biofertilizer products in India is through niche markets. The scope for a biofertilizer is often perceived to be limited.
- Biofertilizers are often perceived to be more expensive than the chemical fertilizers. This is more so since the farmers and small holders received fertilizers heavily subsidized by the government, enabling a few farmers, who appreciate the benefits of use of certain biofertilizers.
- Another perception on biofertilizer is its slow effect on the crops as compared to chemical fertilizers.
- Special care (storage, mixing with powders, etc.) is also needed to handle microbial inocula, so that these remain effective for extended use. These inoculants too favors certain environment. Concerning microbial inoculants, some users realized their potential.
- There was difference of opinion on the effectiveness of microbial inoculants available in the market. Some felt that the performance of these products is often disappointing, unreliable, and not as claimed by the manufacturer. Some products, however, do give good results. All these perceptions contribute to influencing the user on the use of microbial inoculants and biofertilizers.
- The way to forward the produce is that it may satisfy the users in terms of versatility, ease of use, and cost. The use of biofertilizers has still not spread uniformly, although there has been a steady rise in their use by certain group of farmers.

14.7.1 Quality Control

- Quality of biofertilizer is one of the most important factors resulting in their success or failure and acceptance or rejection by end user, the farmers. Efforts to promote the use of these substances in the past have been hampered by poor and uneven quality. A survey of farmers shows that poor quality of biofertilizers is largely responsible for the poor acceptance by users.

- Basically, the quality means the number of selected microorganisms in the active form per gram or milliliter of biofertilizer. Specifications of biofertilizer differ from country to country and may contain parameters like the microbial density at the time of its manufacturing, microbial density at the time of expiry, expiry period, permissible contamination, pH, moisture, microbial strain, and the carrier.
- Quality has to be controlled at various stages of production (during mother culture stage, carrier selection, broth culture stage, mixing of broth and culture, packing and storage). Inoculant shall be packed in 50–75 microns 1000 density polythene packets and should be marked prominently the following:
 1. Name of the product
 2. Name and address of manufacturer
 3. Name of the carrier
 4. Batch number
 5. Date of manufacturing
 6. Date of expiry
 7. Net mass
 8. Crop for which intended
 9. Storage instruction ($15\text{--}30 \pm 0.2$ °C)

Countries like India have regulations for inoculant quality, but neither it is regulated properly nor are the existing regulations well enforced. Bureau of India Standards (BIS) has published necessary specifications/standards for different biofertilizers. But these specifications are purely voluntary in nature and are being regulated on firms and producers, who have opted for BIS certification and putting for BIS certification and putting ISI mark on their products. Gaps exist between India and some of the leading countries in all areas of biofertilizer production and application technology. These gaps are important, particularly in the field of strains, techniques used for sterilization, fermentation, and carriers. The effects of these are aggravated further by poor quality control maintained by most of the producers. Under such circumstances, a significant increase in the level of penetration and demand of biofertilizers are possible only if these gaps are reduced and strict quality control is maintained. While promotional efforts are important, but the success of such efforts will depend on the availability of biofertilizers of high and consistent quality. A system by which the quality is monitored by the central and state-level authorities must be devised and enforced.

14.7.2 Challenges and Prospects of Fungal Biofertilizer Usage

Despite significant improvement and progress in biofertilizer technology over the years, the progress in the field of its production technology is below satisfaction. Further, there do exist obstacles for biofertilizer use. These are not only technical problems, but also socio-economic and human resource obstacles. The technical

problems can be addressed through a comprehensive program of basic and applied research up to a certain extent. Overcoming the socioeconomic and human resource obstacles would require an emphasis on education, training, and promotion of private enterprise. The difficulties to expand the use of biofertilizer by farmers in India are as follows:

1. Difficult handling of biofertilizer
2. Problems with distribution
3. Low quality of biofertilizer
4. Cannot be stored for longer period
5. Lack of demonstration and low visual effect of biofertilizer
6. Low in public relation and technology transfer
7. Low knowledge of farmer on sustainable agriculture and environmental effect of biofertilizer

One of the most important limitations of biofertilizers is seen in their nutrient content when compared to inorganic fertilizers as a number of works have reported deficiency symptoms associated with plants grown in biofertilizer-treated soils (Itelima et al., 2018). However, this challenge is curbed by the addition of substances such as bone meal (rich in phosphorus), wood ash (rich in potassium), or other substances of natural origin such as phosphate rock to enrich the fertilizer (Itelima et al., 2018). Also, the use of nutrient-rich organic materials such as palm wastes (rich in potassium) and wood ash (rich in potassium also) in making biofertilizers serves as a good remedy. Mahimairaja et al. (2008) stated that the addition of phosphorus to wastes makes the biofertilizers more balanced and reduces nitrogen losses. Notably also are the limitations associated with the storage conditions of biofertilizers; this goes a long way in affecting its efficacy. Biofertilizers should be stored in cold storage conditions away from heat and the polythene used in packaging should be of low-density grade and thickness of about 50–75 microns (Mishra & Dadhich, 2010). As good as they are (biofertilizers), sometimes they do bring about the expected outcome owing to exposure to high temperatures and other harsh environmental conditions.

Other constraints limiting the use of biofertilizers technology may include human resource, unavailability of suitable strains, and unavailability of suitable carrier. Chen (2006) has identified some bottlenecks for effective biofertilizer inoculation:

- Unavailability of suitable strains: This is one of the major constraints in the production of biofertilizers because only suitable strains have the ability to survive both in the broth and in the inoculant carrier.
- Unavailability of suitable carrier: If suitable carrier material is not available, it becomes difficult to maintain the shelf-life of biofertilizer. In terms of suitability of carrier, the order is peat, lignite, charcoal, farmyard manure, soil, and rice bran.
- Lack of awareness among farmers: Farmers are mostly not aware of biofertilizers and their usefulness in increasing crop yields. They are unaware of the damages caused on the ecosystem by continuous application of inorganic fertilizer and increasing bioaccumulation of toxic materials in plants.

- Skill and technical knowhow: It is another problem. This is because the unskilled and inadequate staff farmers are not given proper instructions about the application. This however affects in general terms the success of biofertilizers.
- Environmental constraints: Soil characteristics like salinity, acidity, drought, water logging, and a number of climatic conditions affect the use of biofertilizers. Table 14.8 throws light on various constraints in the production and commercialization of biofertilizers.

Table 14.8 Constraints/limitations in the production and commercialization of biofertilizers

| Constraints/limitations | | Recommendations |
|-------------------------|---|---|
| Technical constraints | Unavailability of suitable carrier material | Identification and selection of appropriate economic carrier to maintain shelf-life and effectiveness of biofertilizers Skilled staff should be hired and manpower should be trained via proper training |
| | Lack of skilled staff in production | Frequent monitoring of the biofertilizer production units for quality assurance |
| Marketing constraints | Lack of regulation and units standards for biofertilizer | Necessary legislation for monitoring biofertilizers should be done by the government |
| | Limited transportation and storage facilities | |
| | Poor and incomplete labeling of biofertilizer products | Proper labeling of biofertilizers should be done (giving genus name, viable count, and expiry date, etc.) |
| | Lack of promotion network and publicity among the end users. | Wide publicity of biofertilizers should be done through scientific training, fairs, exhibitions, or media |
| Biological constraints | Unavailability of appropriate strain. Tendency of strain to mutate during fermentation | Continuous efforts for identification of strains and screening for their efficiency across the type of soil, agro-climatic conditions, and farming situations are recommended |
| | Nonavailability of right inoculant at the right place at the right time | Standardization of biofertilizer dose in a particular crop and soil. Understanding of strain effectiveness should be strengthened through extensive research in this field |
| Field-level constraints | Soil conditions like acidity, presence of salts and toxic elements in the soil, and extreme climatic conditions may make the results of biofertilizers inconsistent | It is needed to strengthen the research and technologies to reduce effects and to counteract stated soil and environmental conditions |
| | Inadequate awareness among the farming community about bioinoculants | A strong training and awareness program may be initiated to aware and motivate farmers |
| Financial constraints | Nonavailability of sufficient fund and equipment from government and private bodies | Use of high-tech equipment is required for quality products |
| | | Government should provide funding and loans for development of production units |

14.8 Conclusions

- Fungi as biofertilizers in agriculture sector are one of the emerging areas for growth and enhancement of crop production for sustainable agriculture. This review envisioned the future beneficial role of plant growth-promoting fungal communities in plant growth promotion and soil fertility. These PGP fungi can be used as biofertilizers, bioinoculants, and biocontrol agents in place of chemical fertilizers and pesticides in an environmentally and eco-friendly manner.
- Use of beneficial microbes as biofertilizers and biopesticides in agriculture sector are one of the methods that can pave way for the next agricultural green revolution. Beneficial fungi, associated with plants, are playing an essential role in the development and growth of plants through various mechanisms including solubilization of different insoluble and unavailable nutrients and production of plant growth regulators. Fungi also help in alleviating biotic stresses like plant pest and pathogen and abiotic stresses like drought, salinity, temperature, heavy metals, etc. Fungi are playing significant role in bioremediation of heavy metals and other pesticides also.
- There are many commercial fungal biofertilizers available worldwide. Using fungal biofertilizers offers more environmentally friendly alternatives than chemical fertilizers. However, there are some limitations in using these products. The success can be affected by environmental conditions, while application difficulties, limited shelf-life, and slow action as compared to chemical products may discourage farmers to use the biofertilizers. A lot of research done in the past few decades has enabled these fungi to emerge as a potential biofertilizer, a cheap and environment-friendly alternative to expensive and harmful chemical fertilizers. This aspect of a conventional to alternative route to more food grain production in a sustainable manner especially gains significance for developing countries. Among the fungal biofertilizers, mycorrhizal biofertilizer products greatly appeal to the agro-industry mainly due to its versatility and use of environment-friendly technology. The AM biofertilizer technology can be called poor man's technology.
- Microbial mixed inoculum (biofertilizers and biopesticides) or consortium, used as an alternate to synthetic fertilizers and pesticides to increase the soil fertility and control disease and pests in agriculture is gaining prominence. Biofertilizers and biopesticides are environment-friendly products and can be used in integrated nutrient management (INM) and integrated pest management (IPM) techniques.
- White biotechnology refers to the use of living cells and/or their enzymes to create industrial products that are more easily degradable, require less energy, and create less waste during production. The fungal white biotechnology includes biodiversity of fungi from different habitats, including extreme environments (high temperature, low temperature, salinity, and pH) and associated with plants (epiphytic, endophytic, and rhizospheric) and their industrial applications in diverse sectors. Endophytes have generally greater beneficial effects (FAO,

2014) and are less likely to encounter an extreme change of the soil environment because of the insulation of plants. Thus, to achieve better yield and protection of crop, the use of endophytic biofertilizers can be preferred over other microbial species. Endophytes can be explored as effective biofertilizers as they have an advantage of living within a plant's tissues and therefore more readily exert a direct beneficial effect (Vandenkoornhuyse et al., 2002).

- Future research in rhizosphere biology will rely on the development of molecular and biotechnological approaches to increase our knowledge of rhizosphere biology and to achieve an integrated management of soil microbial populations.
- In the agricultural sector, vegetables form the most vital part of Indian agriculture and nutritional security due to their short life cycle, high yield, nutritional richness, economic viability, and also provide employment opportunities to rural nonfarming and farming sector. India has diverse climatic conditions with different weather phenomena, which help in growing a wide range of vegetable and fruit production. India ranks second in worldwide vegetable and fruit production. In our continual combat with pests along with other agro-ecological and environmental challenges, while meeting the nation's demand to feed the ever-increasing human population, the use of pesticide is unavoidable. The inefficient farm management practices and improper application of agrochemicals and pesticides have caused degradation of soil fertility and health apart from affecting different biotic and abiotic factors (Setboonsarng & Gilman, 2006).
- The need of the hour is to develop effective biofertilizers, with both pesticide tolerance and plant growth-promoting (PGP) traits, isolated indigenously from the crop field soil. Sustainable agricultural practices are essential for providing better food and fiber for a healthy nation. Only effective microbes are suitable alternatives to chemical fertilizers and pesticides for enhancing plant production under sustainable agriculture. These microbes also have the ability to survive under pesticide threat and adverse environmental conditions because indigenous microbes can change their physiological and genetic character according to environmental conditions.
- The interest in biofertilizers is increasing and so is the potential for their use in sustainable agriculture. However, many of the products that are currently available worldwide are often of very poor quality, resulting in the loss of confidence from farmers. The formulation of an inoculant is a crucial multistep process that should result in one or several strains of microorganisms included in a suitable carrier, providing a safe environment to protect them from the often harsh conditions during storage and ensuring survival and establishment after introduction into soils. One of the key issues in formulation, development, and production is the quality control of the products at each stage of the process. Quality control of products is of great importance. This is always a challenge with microbial products because the mass production of living organisms tends to select those best suited to mass production rather than those found more effective in the field. Research in the field, supported by advanced technology, will enhance biofertilizer use in the country as well as profitability for the small and marginal farmers.

- Potentiality of pesticide-tolerant fungi as biofertilizers can pave the way for bio-entrepreneurship. Agricultural sustainability and economic stability of rural India depend upon scientific capacity building amongst the farmers. The combination of the biological potentials of plant–holobiont and soil mycobiome, when fully harnessed under agricultural scenario, will help to boost food security globally. The biofertilizers are important to ensure a healthy future for the generations to come. Long-term use of biofertilizers is economical, eco-friendly, more efficient, productive, and accessible to marginal and small farmers over chemical fertilizers.

In a nutshell, the future is envisioned to be most promising for rural India by following the key points mentioned below:

- Shifting the focus from chemical fertilizers to biological fertilizers will lead to the proliferation of biofertilizer industries and incentivizing the ecological benefits will encourage the farming community for application of biofertilizers.
- Biofertilizers would play a key role in productivity and sustainability of soil and protect the environment as eco-friendly and cost-effective inputs for the farmers.
- Strain improvement using genomic and biotechnological tools for better nutrient fixation/solubilization and plant growth-promoting attribute will further lead to gaining confidence in biofertilizers by stakeholders.
- Selection of effective and competitive multifunctional biofertilizers for a variety of crops.
- Quality control system for the production of inoculants and their application in the field.
- Study of microbial persistence of biofertilizers in soil under stressful environments conditions.
- Establishment of "Bio-fertilizer Act" and strict regulation for quality control in markets and application.

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Chapter 15

Efficacy of Plant Products as Biofungicides for Postharvest Decay of Root, Tuber, and Bulb Crops: An Opportunity for Bioentrepreneurship



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15.1 Introduction

There is a worldwide swing to the use of eco-friendly methods for protecting the crops from pests and diseases as the use of potentially harmful chemicals is viewed with several disadvantages. Plants have been known for their medicinal and antimicrobial properties since ancient times, thereby indicating that the plant kingdom is a vast storehouse of chemicals that can check many plant pathogens. Recently, in different parts of the world, attention has been paid toward exploitation of plants as novel chemotherapeutants in the plant protection in view of their long-term effect on crop disease management, low cost, and safety to ecosystem. Biofungicide

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means fungicides of biological origin which may be microorganism such as bacteria, fungi, and animal or plant-based product like secondary metabolite.

Vegetables are an important source of human nutrition and are a rich and comparatively cheaper source of vitamins and minerals. They are important for the maintenance of good health and are also beneficial in protecting against some degenerative diseases. The vegetables grouped under root, tuber, and bulb crops are most important as compared to other vegetables being nonperishable in nature. After harvest, they can be stored and consumed throughout the year. Some important vegetables included in this group are potato (*Solanum tuberosum* L.), sweet potato [*Ipomoea batatas* (L.) Lam.], radish (*Raphanus sativus* L.), carrot (*Daucus carota* L.), beet (*Beta vulgaris* L.), colocasia [*Colocasia esculenta* (L.) Schott.], yam (*Dioscorea alata* L.), Elephant's foot yam [*Amorphophallus paeoniifolius* (Dennst.) Nicolson], onion (*Allium cepa* L.), and garlic (*Allium sativus* L.). After harvest, the produce is stored in locally prepared bins, in open store houses, or in bamboo baskets for their consumption and marketing throughout the year.

The issue of postharvest disease management of these crops has become an important concern due to inadequate agricultural storage practices for produce preservation from microbes-induced spoilages (Kana et al., 2012). These crops are drought tolerant and provide a wide harvesting window which makes it act as a famine reserve. However, postharvest deteriorations caused by microbial invasion of the tubers are the most important causes of loss in its production and contribute hugely to the unsuccessful long-term storage of the root tubers. The pathogenic organisms including fungi gain entry into tubers probably through the area where the tubers are separated from stems at harvest or the root tips which break during harvesting, or other natural openings and cracks on the tuber's surfaces sustained during harvesting, transit, or storage (Okigbo, 2009). Improper handling leaves off bruises on the produce during harvesting, transit, or storage which may become contaminated from adherent soil or air-borne propagules which later colonies such abrasions to cause rots (Okigbo & Nmeka, 2005). In Odisha, the postharvest decay of these vegetables during storage poses serious problems and sustains heavy losses particularly for the warm and humid climate. According to various workers, rotting usually starts from the field and progresses in storage (Okigbo & Ikediugwu, 2000). Unhealthy store conditions lead to the absorption of moisture by produce in storage as a result of defects in the storage facility, thus encouraging the development of hotspots and molds.

Antimicrobial chemicals such as benzimidazoles, aromatic hydrocarbons, and sterol biosynthesis inhibitors are often used in the control of plant diseases in agriculture. However, there is a series of problems against the effective use of these chemicals in areas where the fungi have developed resistance (Brent & Hollomon, 1998). In order to overcome this problem, higher concentrations of the chemicals had been used, but this increased the risk of high-level toxic residues in the products which were not beneficial for human beings and the environment. Thus, there has been a growing interest on the research of possible use of plant extracts for control of pest and diseases in agriculture which was less harmful to human health and the environment (Costa et al., 2000; Duru et al., 2003).

To avoid use of synthetic fungicides, it is imperative to find alternative sources from naturally occurring compounds that are easily biodegradable and of low mammalian toxicity for a safe control of fungal pathogens (Adongo et al., 2012). Plants are the source of natural pesticides that make excellent leads for new pesticide developments (Shanmugavalli et al., 2009). Recent studies on the use of plant extract have opened a new approach to control plant diseases. These plant extracts have been reported to be safe, nontoxic to man, but effective against plant pathogens (Shivpuri et al., 1997). Plant diversity serves humankind as renewable natural resources for a variety of biologically active chemicals. Active constituents of the medicinal and aromatic plants have been found to be less phytotoxic, more systemic, and easily biodegradable (Fawcett & Spencer, 1970). Plants are the sources of natural pesticides that make excellent leads for new pesticide development (Gangadevi et al., 2008; Satish et al., 2008; Shanmugavalli et al., 2009; Swarna Latha & Neelakanta Reddy, 2009).

In India, Dastur (1916) first studied the control measure for ripe rot of plantains. After a long period of gap, studies in this field gained rapid momentum, and within two decades, much knowledge was gained on fungi causing postharvest rot of fruits and vegetables from several parts of the country, particularly from Pune, Allahabad, Lucknow, Delhi, Kolkata, Bhagalpur, and Bangalore. Unfortunately, little work has been done so far in this field in Odisha (Das, 2012). In continuation of the previous work, the present topic was selected in view of the limited studies on postharvest fungal rots of root, tuber, and bulb vegetables. The present article is, therefore, based upon the postharvest decay of root, tuber, and bulb vegetables and their control by biofungicides. It is expected that the present investigations shall throw some light on the etiology and control of postharvest storage decay of vegetables such as potato, sweet potato, colocasia, yam, amorphophallus, beet, carrot, radish, onion, and garlic caused by various species of fungi and their management by application of botanicals to avoid serious economic losses in the markets and store houses.

15.2 Methodology

15.2.1 Pathological Studies

(i) *Collection of Rotten Samples of Root, Tubers, and Bulb Crops*

A total of 1000 samples of rotten potatoes, sweet potatoes, radish, carrot, beet, colocasia, yam, amorphophallus, onion, and garlic were collected from various markets and store houses of Odisha during 2014–2016. Rotten plant samples were collected in separate polythene bags with labels in respect of (1) date of collection, (2) place of collection, (3) extent of damage in sample lots, (4) viable symptoms (smell, color, and mycelial growth), and (5) name of collector. All the samples were brought to the laboratory and surface sterilized with 0.1% mercuric chloride solution for 2–3 min. Some specimens from each sample were examined for the

presence of fungal infections by taking a small portion of the rotten tissues on a clean glass slides, macerated with lactophenol-cotton blue, and examined under a microscope for the occurrence of fungal mycelium and fruiting bodies, if any. Specimens showing the signs of fungal infections were immediately processed for phytopathological analysis, after recording the visual symptoms of rotting.

(ii) *Isolation of Fungi from Rotten Samples*

Sterile moist chambers were prepared out of Petri dishes, 100 mm in diameter, by placing triple-layer moist filter paper over which a platform was prepared by placing two pieces of clean glass slides over a pair of equal-sized glass rods. It was sterilized at 15 PSI for 20 min in an autoclave.

Small pieces of tissues including both rotten and healthy areas were surface sterilized with 0.1% mercuric chloride solution for 2–3 min, washed thoroughly three times with sterile distilled water, and transferred aseptically into the sterile moist chamber at the rate of 5 pieces per chamber. Similar sterilized tissues were also transferred into petriplates containing potato dextrose agar (PDA) at the rate of 5 pieces per petri plate.

All the materials were incubated at room temperature ($28\text{ }^{\circ}\text{C} \pm 1\text{ }^{\circ}\text{C}$) for four to five days. At the end of incubation, the sterile chambers as well as the petriplates were examined for the presence of fungi. Fungi of all kinds appearing on the incubated materials were transferred aseptically into PDA slants. Each fungus as obtained in PDA slants was subsequently purified by single spore/hyphal tip isolation technique and maintained by periodic subculturing in PDA slants for further studies. The frequency of incidence of each fungus on different samples of rotten samples was estimated by the formula:

$$\text{Frequency of incidence} = X / Y \times 100$$

Where X = the number of pieces showing the presence of different fungi and Y = total number of pieces investigated. The frequency of incidence was expressed in terms of percentage.

(iii) *Identification of Fungi*

The colony characteristics of each fungal isolates on PDA were recorded. The details of morphology of each isolate were studied under varying magnifications ($\times 100$ to $\times 1000$) of a microscope. The characteristics of mycelium and fructifications were recorded. The dimensions of each structure were measured with the help of an Ocular micrometer. Each fungal isolate was taxonomically identified with the help of available literatures and confirmed by the microbiology Department, OUAT, Bhubaneswar, and Odisha. The fungal isolate which was found to be new to their molecular identification was carried out by sequencing of 26S rRNA gene followed by submission of sequence to NCBI GenBank. Genomic DNA was extracted from these fungal isolates (Sambrook et al., 1989) followed by amplification of 26S rRNA gene using universal primers DF (5'-ACCCGCTGAACTTAAGC-3') and DR (5'-GGTCCGTGTTTCAAGACGG-3') and sequencing at Xcelris Genomics, India.

Raw sequences were analyzed by Bio-Edit software (v7.0.5.3) and identification of the isolates by BLASTN in NCBI database (www.ncbi.nlm.nih.gov/nucleotide). The phylogenetic tree was constructed using the UPGMA distance algorithm (Wheeler & Kececioglu, 2007) using Mega software v5.1. The resultant tree topologies were evaluated by boot-strap analysis of muscle datasets based on 500 resamplings. The final sequences were submitted to NCBI GenBank to obtain the accession number of each isolate.

15.2.2 In Vitro Evaluation of the Antifungal Efficacy of Plant Extracts

(i) Selection of the Test Fungal Isolates

A total of ten fungal species were found to be associated with postharvest decay of potato, sweet potato, radish, carrot, beet, taro, yam, amorphophallus, onion, and garlic during the present investigations. All the fungal species were purified, identified, and maintained in potato dextrose agar slants by periodically subculturing to use in the present studies.

(ii) Selection of Plants and Plant Parts

On the basis of local availability and medicinal values, the leaves of the 16 plant species such as *Abutilon indicum* (L.) Sweet (Malvaceae), *Ageratum conyzoides* L. (Asteraceae), *Alstonia scholaris* (L.) R.Br. (Apocynaceae), *Artocarpus heterophyllus* Lam. (Moraceae), *Averrhoa carambola* L. (Averrhoaceae), *Cassia fistula* L. (Caesalpinaceae), *Centella asiatica* Urban. (Apiaceae), *Dillenia indica* L. (Dilleniaceae), *Eucalyptus globulus* Labill (Myrtaceae), *Haldinia cordifolia* (Roxb.) Rids (Rubiaceae), *Justicia adhatoda* L. (Acanthaceae), *Lawsoniainermis* L. (Lythraceae), *Murraya paniculata* Jack. (Rutaceae), *Pithecellobium dulce* (Roxb.) Benth. (Mimosaceae), *Pongamia pinnata* (L.) Panigrahi (Fabaceae), and *Tamarindus indica* L. (Caesalpinaceae) were selected for the present investigations. Some of these plants have been reported to exhibit antifungal properties.

(iii) Collection, Processing, and Extraction of Plant Material

Leaves of 16 angiosperm plants belonging to diverse plant families were collected from different parts of Odisha. Each material was collected in separate polythene bags and brought to the laboratory of Department of Botany, S.J. College of Education and Technology, Bhubaneswar, Odisha (India), for further studies.

Three solvent systems were used for leaf compound extraction, viz., aqueous, petroleum ether, and methanol. For the preparation of aqueous extract (cold water extract), fresh plant leaves were washed thoroughly with sterile water, chopped into pieces, and the extracts were obtained by macerating with 100 ml of sterile distilled water for each 20 g of fresh leaf material. The extracts thus obtained were separately filtered through a three-fold sterilized muslin cloth into sterilized conical flasks.

Then the extracts were refiltered with Whatman No 1 filter paper using a power-operated suction filter pump. These extracts formed the standard plant extract solution (20%) and were stored at 5 °C in the refrigerator for further use. For dried leaf extraction, reflux method of solvent extraction was used for purification of different organic constituent. Solvent series for successive separation was as follows: petroleum ether and methanol. First, the soxhelt extraction unit was cleaned and rinsed with petroleum ether and then 30 g of air-dried and coarse-powdered plant material were kept in it. Then, this unit was kept on water bath and exhaustively extracted with petroleum ether for 30–36 h. During extraction, the round bottom flask (RBF) which contains the extracts should be with porcelain chips for well siphonation. To confirm whether the extraction is complete or not, the extract from siphonating tube of soxhelt was taken in a watch glass. After petroleum ether extraction, the extracted plant material was then air dried, repacked in soxhelt apparatus, extracted with methanol for such time hour, and maintained again. The crude extract and the fraction obtained at every step were filtered and distilled to evaporate the solvent from the extract. The liquid extracts were concentrated separately by using a rotary vacuum evaporator, and the resulting dried extract was preserved in desiccators until further use. The diluted plant extracts were prepared by dissolving purified petroleum ether and methanolic extract in dimethyl sulfoxide (DMSO) @ 20 mg plant extract in 1 ml DMSO.

(iv) *Evaluation of Antifungal Efficacy of Plant Extracts*

For the evaluation of antifungal effect of plant extracts on inhibition of mycelia growth of ten isolated fungal species, namely, *Apergillus flavus*, *Apergillus niger*, *Fusarium oxysporum*, *Geotrichum candidum*, *Junghuhnia* sp. AK15 (NCBI Gene Bank accession number KT 946990), *Mucor* sp. BOT15 (NCBI Gene Bank accession number KT 946994), *Neoscytallidium* sp. AM15 (NCBI Gene Bank accession number- KT 946992), *Penicillium* sp. KBS15 (NCBI Gene Bank accession number KT 946991), *Penicillium* sp. UU15 (NCBI Gene Bank accession number KT 946993), and *Rhizopus oryzae*, poison food technique was adopted. For this purpose, Potato Dextrose Agar (PDA) medium was prepared and sterilized at 15 PSI for 20 min. After sterilization, 1 ml of each diluted plant extract solution (20 mg extract/1 ml DMSO) was mixed with 19 ml of PDA (Potato Dextrose Agar) and poured into each petriplate @ 1 mg/ml, mixed thoroughly, and allowed to solidify. PDA medium without the plant extracts but substituted with 1 ml distilled water was served as control.

The mycelial disc of 5 mm in diameter was taken from the periphery of 2- to 3-day-old colonies of the desired fungi with the help of a sterilized cork-borer. These discs were transferred aseptically inside the laminar flow with the provision of ultraviolet light and inoculated at the center of each petriplate containing PDA with plant extracts. Three replications were maintained for each fungus × plant extract combination. A separate set of petri plates containing only PDA was inoculated by each test fungus to serve as control. All the test petriplates were incubated at room temperature (28 ± 1 °C) and examined at regular intervals. Growth of the isolates was considered as optimum for recording the observations when it reached

around 8 cm in any of the control sets. *Rhizopus oryzae* was incubated for 1 day, *Neoscytallidium* sp. AM15 for 3 days, *Junghuhnia* sp. AK15 and *Mucor* sp. BOT15 for 5 days, *Aspergillus flavus* for 7 days, *Aspergillus niger* for 8 days, *Penicillium* sp. UU15 and *Fusarium oxysporum* for 10 days *Geotrichum candidum* for 11 days, and *Penicillium* sp. KBS15 for 15 days depending upon their complete growth (8 cm) on the petriplate. At the end of incubation period, the diameter of the colony in each petriplate was measured in two planes, one at the right angle to the other. Further, the data of each test plant extract was compared with that of the control in order to estimate the inhibition (positive) or stimulation (negative) effects of the plant extract tested. The formula used for this purpose was:

$$\% \text{inhibition / Stimulation} = X - Y / X \times 100$$

Where X = colony diameter in control and Y = colony diameter in treated plated.

For the sake of convenience of description of the results, it was classified by the method of Rath and Mohanthy (1978) with slight modifications as given below:

| | |
|-----------------|----------------------|
| % of inhibition | Rating |
| 0 or negative | No inhibition |
| 1–10 | Incipient inhibition |
| 11–30 | Mild inhibition |
| 31–70 | Moderate inhibition |
| 71–100 | Severe inhibition |

While screening the data on the basis of the above-noted scale, it was noted that most of the effective inhibition of the colony diameter was in the range of 31–70%. Only in a few instances, it exceeded 70%.

(v) *Comparison of Plant Extracts and Estimation of Inhibition of Mycelial Growth by Selected Plant Extracts*

For this purpose, three plants were selected that show the mean maximum inhibition of mycelial growth of ten test fungal species, namely, *Aspergillus flavus*, *Aspergillus niger*, *Fusarium oxysporum*, *Geotrichum candidum*, *Junghuhnia* sp. AK15, *Mucor* sp. BOT15, *Neoscytallidium* sp. AM15, *Penicillium* sp. KBS15, *Penicillium* sp. UU15, and *Rhizopus oryzae*. The selected plant species were *Ageratum conyzoides*, *Alstonia scholaris*, and *Averrhoa carambola*.

(vi) *Comparison of the Efficacy of Synthetic Fungicides and Selected Plant Extracts*

A separate experiment was conducted in order to compare the relative efficacy of four synthetic fungicides, viz., Dhanustin, Blitox-50, Indofil-M-45, and Mancozeb, with the petroleum ether and methanolic leaf extracts of *Ageratum conyzoides*, *Alstonia scholaris*, and *Averrhoa carambola*. The details of concentrations are given in Table 15.1:

Table 15.1 Comparison of plants and fungicides used along with their concentration

| Treatment | Solvent system | Concentration mg/ml | Chemical constituent |
|----------------------------|------------------|---------------------|---|
| Blitox-50 | | 0.1 | Copper oxychloride |
| Dhanustin | | 0.1 | |
| Indofil-M-45 | | 0.1 | Carbamate |
| Mancozeb | | 0.1 | Coordination product of zinc and magnesium ethylene bis dithiocarbamate |
| <i>Ageratum conyzoides</i> | <i>Aqueous</i> | 4 | Natural plant product |
| | <i>Pet-ether</i> | 1 | |
| | <i>Methanol</i> | 1 | |
| <i>Alstonia scholaris</i> | <i>Aqueous</i> | 4 | Natural plant product |
| | <i>Pet-ether</i> | 1 | |
| | <i>Methanol</i> | 1 | |
| <i>Averrhoa carambola</i> | <i>Aqueous</i> | 4 | Natural plant product |
| | <i>Pet-ether</i> | 1 | |
| | <i>Methanol</i> | 1 | |
| Control | – | – | No fungicide or plant product |

Incorporation of the test fungicides/natural plant extracts was carried out as per the method described earlier. The required quantity of fungicides and plant extracts was incorporated after sterilization of the medium. The petriplates containing PDA and fungicides/plant extracts were inoculated by 10 test fungal isolates in three replicates and incubated for 1st to 15th days depending upon their respective complete growth as described earlier. At the end of incubation period, the colony diameter of each fungus-fungicide/plant extract combination was measured and transferred to percentage of mycelial inhibition as per the methods described earlier.

(vii) Formulation of Herbal Fungicides and Recommendation for Use

A combination of petroleum ether leaf extracts of *Ageratum conyzoides* (10 mg) and *Alstonia scholaris* (10 mg); *Ageratum conyzoides* (10 mg) and *Averrhoa carambola* (10 mg); and *Alstonia scholaris* (10 mg) and *Averrhoa carambola* (10 mg) was tested against ten fungi tested. The concentration of the formulated combination was 20 mg/ml DMSO (10 mg extract of each plant).

15.3 Results and Discussion

The data on the incidence and taxonomic identity of fungi in root, tuber, and bulb crops revealed that eight genera comprising of ten species of fungi were responsible for storage decay of ten root, tuber, and bulb vegetables collected from different market places and storehouses of Odisha in frequencies ranging from 9.74% to 50.3% of the cases studied. *Aspergillus*, *Geotrichum*, *Penicillium*, and *Rhizopus* were found to be the dominant fungi causing maximum loss of those vegetables

during storage due to rotting. It was also noted that these fungi are generally the soil inhabitants which come in contact with the tubers and fleshy roots of these vegetables during the crop stand as surface microflora. After harvest and during transport, transit, and storage, they infected the tubers and roots through wounds arising out of harvest, transportation, and storage handlings or by injuries caused by insects and other microorganisms when the environmental factors are favorable for their growth and disease development, incurring heavy economic losses to both growers and consumers. Experiments conducted on the pathogenicity tests indicated that fungi isolated from rotten samples of potato, colocasia, amorphophallus, sweet potato, yam, carrot, radish, beet, onion, and garlic were pathogenic to their respective hosts in an artificial inoculation test.

The data on the pathogenicity test of six fungi on potato tubers indicated that *Aspergillus niger* was the most pathogenic one and caused the complete rotting of tuber within 10 days of incubation. The pathogenicity test data of colocasia rotting caused by five fungi indicated that *Rhizopus oryzae* was the most pathogenic to corms than other isolates, and complete rotting of the corm within 10 days of inoculation was recorded. Amorphophallus rotting was caused by three fungi, and the most virulent was *Geotrichum candidum* which caused the rotting within 30 days of inoculation. Artificial rotting of this host took a long time. Among the two fungi, *Rhizopus oryzae* was the most virulent one causing the complete rotting of sweet potato within 15 days. Similarly, among the five fungi causing storage rot in yam, *Neoscytallidium* sp. AM15 was the most virulent to cause complete rotting within 20 days. Rotting in carrot, radish, and beet root was caused by three fungi, and among them, the most virulent fungus for carrot was *Rhizopus oryzae*, while for radish and beet was *Aspergillus niger*, but both the fungi cause complete rotting of their respective host within 10 days of inoculation. Rotting of onion and garlic was found to be caused by two and three fungi, respectively. Among them, *Aspergillus niger*, for onion, and *Penicillium* sp. UU15, for garlic, were found to be more virulent than others. They cause the complete rotting of the respective host within 15 days of inoculation.

Studies on the inhibition of ten fungal species causing storage rot of root, tuber, and bulb vegetables, by aqueous, petroleum ether, and methanolic leaf extract of sixteen angiospermic plants revealed that petroleum ether and methanolic leaf extracts were more efficient than aqueous extracts (Tables 15.2, 15.3, and 15.4 and Figs. 15.1, 15.2, and 15.3). Petroleum ether and methanolic extracts of all the sixteen test plants were severe to moderately efficient against the ten fungi tested, while the aqueous leaf extract was mild to incipiently efficient to inhibit the mycelial growth. Out of the sixteen plants tested, extracts of *Ageratum conyzoides*, *Alstonia scholaris*, and *Averrhoa carambola* were comparatively more efficient than extracts of other plants tested (Table 15.5 and Fig. 15.4).

Comparative efficacy of plant extracts (petroleum ether and methanolic extract of *Ageratum conyzoides*, *Alstonia scholaris*, and *Averrhoa carambola*) with four commercial fungicides (Blitox-50, Dhanustin, Indofil, and Mancozeb) revealed that plant extracts were more efficient to inhibit the mycelial growth of ten fungi tested (Table 15.6 and Fig. 15.5). Mishra and Dixit (1976) reported that the leaf extracts of

Table 15.2 Screening of aqueous, petroleum ether, and methanolic extracts of *Ageratum conyzoides* L. against ten test fungi

| Test fungi | % of Inhibition of mycelial growth | | | |
|---------------------------------|------------------------------------|-------------------------|--------------------|-------|
| | Aqueous extract | Petroleum ether extract | Methanolic extract | Mean |
| <i>Aspergillus flavus</i> | 25.13 ± 0.56 | 62.90 ± 0.17 | 56.95 ± 0.68 | 48.33 |
| <i>Aspergillus niger</i> | 21.25 ± 0.73 | 62.54 ± 0.17 | 46.51 ± 0.63 | 43.43 |
| <i>Fusarium oxysporum</i> | 15.68 ± 0.29 | 59.80 ± 0.76 | 57.41 ± 1.05 | 44.3 |
| <i>Geotrichum candidum</i> | 16.42 ± 0.63 | 89.03 ± 0.56 | 79.08 ± 0.65 | 61.51 |
| <i>Junghuhnia</i> sp. AK15 | 33.12 ± 0.51 | 98.13 ± 0.38 | 100 | 77.08 |
| <i>Mucor</i> sp. BOT15 | 42.77 ± 0.73 | 99.9 ± 0.08 | 88.48 ± 1.19 | 77.05 |
| <i>Neoscytallidium</i> sp. AM15 | 20.8 ± 0.80 | 86.86 ± 0.26 | 87.87 ± 0.67 | 65.17 |
| <i>Penicillium</i> sp. KBS15 | 26.81 ± 0.66 | 70.29 ± 0.80 | 58.72 ± 1.04 | 51.94 |
| <i>Penicillium</i> sp. UU15 | 14.79 ± 0.40 | 58.35 ± 0.36 | 59.5 ± 0.88 | 44.21 |
| <i>Rhizopus oryzae</i> | 11.78 ± 0.35 | 88.97 ± 1 | 88.88 ± 0.59 | 63.21 |
| Mean | 22.85 | 77.68 | 72.34 | |

Results expressed as mean ± SEM of three determinations

Table 15.3 Screening of aqueous, petroleum ether, and methanolic extracts of *Alstonia scholaris* (L.) R.Br. against ten test fungi

| Test fungi | % of Inhibition of mycelial growth | | | |
|---------------------------------|------------------------------------|-------------------------|--------------------|-------|
| | Aqueous extract | Petroleum ether extract | Methanolic extract | Mean |
| <i>Aspergillus flavus</i> | 16.98 ± 0.62 | 54.26 ± 0.82 | 45.66 ± 1.44 | 38.97 |
| <i>Aspergillus niger</i> | 10.88 ± 0.48 | 34.32 ± 0.36 | 54.72 ± 0.98 | 33.3 |
| <i>Fusarium oxysporum</i> | 5.58 ± 0.54 | 58.55 ± 0.22 | 57.33 ± 1.51 | 40.48 |
| <i>Geotrichum candidum</i> | 18.44 ± 1.11 | 73.41 ± 1.87 | 80.14 ± 1.03 | 57.33 |
| <i>Junghuhnia</i> sp. AK15 | 25.46 ± 0.65 | 100 | 99.46 ± 0.23 | 74.97 |
| <i>Mucor</i> sp. BOT15 | 15.97 ± 0.99 | 100 | 74.48 ± 0.83 | 63.48 |
| <i>Neoscytallidium</i> sp. AM15 | 8.99 ± 0.53 | 84.28 ± 0.75 | 87.25 ± 1.03 | 60.17 |
| <i>Penicillium</i> sp. KBS15 | 10.5 ± 0.23 | 51.66 ± 0.98 | 59.91 ± 0.53 | 40.69 |
| <i>Penicillium</i> sp. UU15 | 14.14 ± 0.65 | 63.1 ± 1.15 | 79.04 ± 1.05 | 52.09 |
| <i>Rhizopus oryzae</i> | 21.18 ± 0.93 | 86.29 ± 0.68 | 89.14 ± 0.65 | 65.54 |
| Mean | 14.81 | 70.58 | 72.71 | |

Results expressed as mean ± SEM of three determinations

Clematis gouriana were superior to Blitox-50, Dithane M-45, Dithane Z-78, and Ziram, while Renu et al. (1985) reported that essential oil from leaf of *Aegle marmelos* possessed greater fungitoxicity than that of six commercial fungicides. El-Shami et al. (1985) evaluated garlic extracts along with five different fungicides *in vitro* against *Fusarium oxysporum* sp. and found that the garlic extract inhibited spore germination and mycelia growth of the fungus in a similar manner to five different fungicides. Bandara et al. (1989) reported that the petroleum and ethyl acetate

Table 15.4 Screening of aqueous, petroleum ether, and methanolic extracts of *Averrhoa carambola* L. against ten test fungi

| Test fungi | % of Inhibition of mycelial growth | | | Mean |
|---------------------------------|------------------------------------|-------------------------|--------------------|-------|
| | Aqueous extract | Petroleum ether extract | Methanolic extract | |
| <i>Aspergillus flavus</i> | 17.49 ± 0.38 | 57.39 ± 0.46 | 54.57 ± 0.86 | 43.15 |
| <i>Aspergillus niger</i> | 20.66 ± 0.64 | 27.8 ± 0.12 | 45.47 ± 0.75 | 31.31 |
| <i>Fusarium oxysporum</i> | 14.44 ± 0.72 | 73.5 ± 0.62 | 58.57 ± 0.44 | 48.83 |
| <i>Geotrichum candidum</i> | 14.03 ± 1.22 | 77.75 ± 0.65 | 70.46 ± 0.84 | 54.08 |
| <i>Junghuhnia</i> sp. AK15 | 19.19 ± 0.45 | 98.93 ± 0.42 | 99.66 ± 0.27 | 72.59 |
| <i>Mucor</i> sp. BOT15 | 9.3 ± 0.53 | 84.81 ± 0.53 | 79.83 ± 0.47 | 57.98 |
| <i>Neoscytallidium</i> sp. AM15 | 22.86 ± 0.69 | 88.93 ± 0.67 | 90.21 ± 0.74 | 67.33 |
| <i>Penicillium</i> sp. KBS15 | 15.93 ± 0.26 | 75.20 ± 0.73 | 73.08 ± 0.73 | 54.74 |
| <i>Penicillium</i> sp. UU15 | 26.08 ± 0.58 | 67.38 ± 1.01 | 67.5 ± 0.7 | 53.65 |
| <i>Rhizopus oryzae</i> | 22.63 ± 0.65 | 87.94 ± 0.62 | 89.7 ± 1.56 | 66.75 |
| Mean | 18.26 | 73.96 | 72.9 | |

Results expressed as mean ± SEM of three determinations

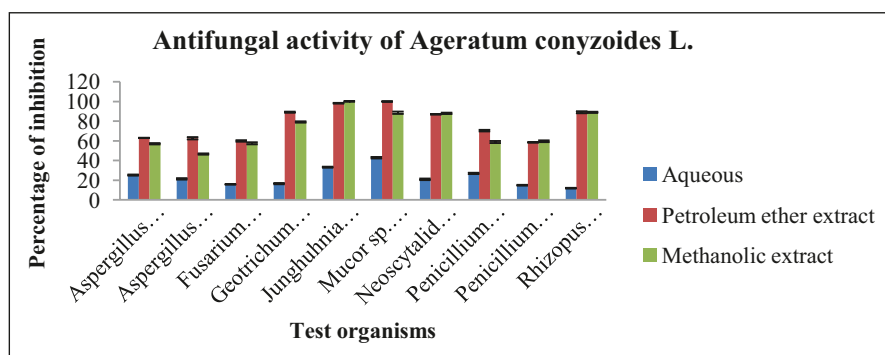


Fig. 15.1 Antifungal activity of *Ageratum conyzoides* L. against ten tested fungi

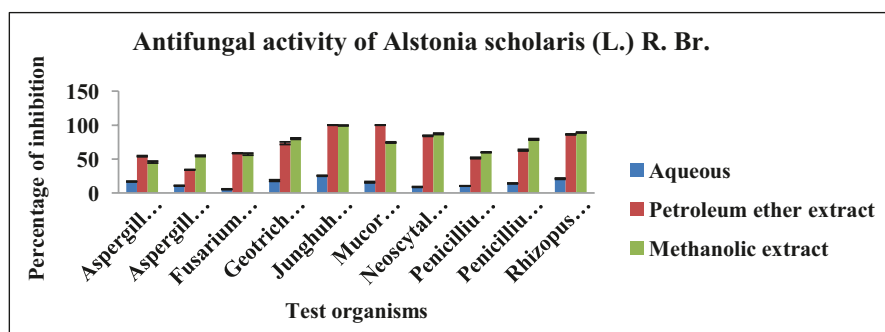


Fig. 15.2 Antifungal activity of *Alstonia scholaris* (L.) R.Br. against ten test fungi

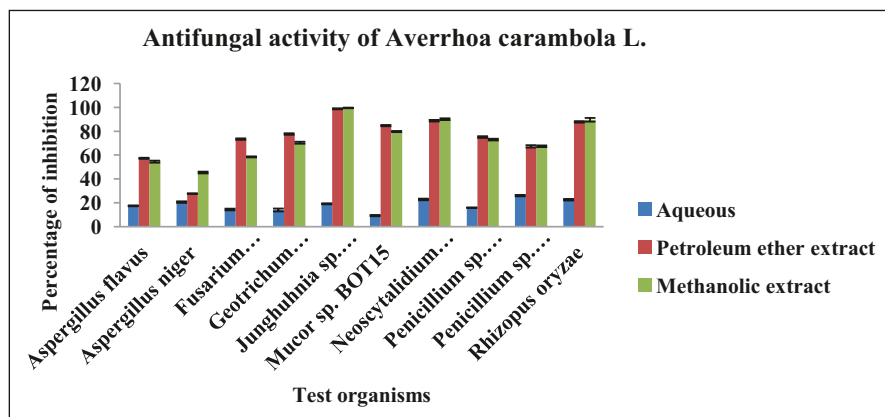


Fig. 15.3 Antifungal activity of *Averrhoa carambola* L. against ten tested fungi

extracts of *Butea monosperma* were fungitoxic to *Cladosporium cladosporioides*. During the present investigations, similar study on comparing the effect of four commercial fungicides with three promising plant extracts could reveal that the activity of a particular plant extract or fungicide effective against a particular test fungus was quite ineffective against others in inhibiting the mycelial growth. For inhibiting the mycelial growth of *Aspergillus flavus*, *Fusarium oxysporum*, and *Penicillium* sp. KBS15, Dhanustin and Mancozeb were found to be superior over the three test plants, but the activity of plant extracts was found to be more efficient than Blitox-50 and Indofil. For growth retardation of *Aspergillus niger*, the plant extracts were found to be more efficient than the test commercial fungicides, except Dhanustin. Against *Geotrichum candidum*, the fungitoxic potential of *Ageratum conyzoides* was found to be equal with Mancozeb, while the effect of the rest of the plant extracts was more than rest of the three commercial fungicides. However, the antifungal activity of plant extracts was found to be superior over fungicides against *Mucor* sp. BOT15. Against *Junghuhnia* sp. AK15, all the test plant's extract and commercial fungicides showed significant inhibitory effect, except Blitox-50. The effect of Dhanustin, which completely inhibited the mycelial growth of *Aspergillus flavus* and *Penicillium* sp. UU15, confirmed its superiority over all the test fungicides and three plant species. For the complete inhibition of mycelial growth in the case of *Geotrichum candidum*, *Junghuhnia* sp. AK15, *Mucor* sp. BOT15, *Neoscytallidium* sp. AM15, *Penicillium* sp. KBS15, *Penicillium* sp. UU15, and *Rhizopus oryzae*, all the three test plant species were significantly superior over all the four commercial fungicides.

Studies on the antifungal properties and potentials of plants and plant products have now been extended in several directions, namely identification of the potential microbiocidal products *in vitro*; their isolation, identification, and characterization in the laboratory; comparison of plant products and extracts with commercial fungicides; and testing the efficacy of plant products in the field, *in vivo*. All these activities put together shall enable us not only locate and use the present herbal

Table 15.5 Comparative efficacy of antifungal activity of sixteen angiospermic plants

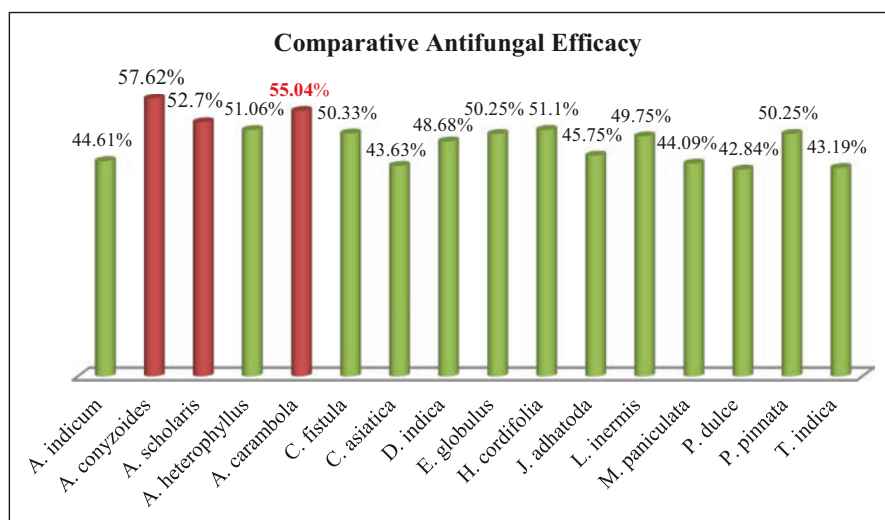
| Growth inhibition of test fungal species (%) | | | | | | | | | | | |
|--|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|--------------|
| Test plants | Af | An | Fo | Gc | Ja | Mb | Na | Pk | Pu | Ro | Mean |
| <i>Abutilon indicum</i> | 37.99 ± 19.24 | 26.4 ± 8.97 | 28.5 ± 4.74 | 43.75 ± 16.8 | 76.72 ± 28.59 | 51.86 ± 24.83 | 58.53 ± 24.42 | 28.59 ± 14.63 | 41.62 ± 11.63 | 52.2 ± 33.95 | 44.61 |
| <i>Ageratum conyzoides</i> | 48.33 ± 16.58 | 43.43 ± 16.99 | 44.3 ± 20.26 | 61.51 ± 32.14 | 77.08 ± 31.09 | 77.05 ± 24.68 | 65.17 ± 31.38 | 51.94 ± 18.38 | 44.21 ± 20.8 | 63.21 ± 36.36 | 57.62 |
| <i>Alstonia scholaris</i> | 38.97 ± 15.93 | 33.3 ± 17.91 | 40.48 ± 24.68 | 57.33 ± 27.63 | 74.97 ± 35.01 | 63.48 ± 35.17 | 60.17 ± 36.21 | 40.69 ± 21.61 | 52.09 ± 27.61 | 65.54 ± 31.38 | 52.7 |
| <i>Artocarpus heterophyllus</i> | 42.53 ± 25.42 | 31.03 ± 12.2 | 40.84 ± 17.07 | 61.77 ± 25.62 | 73.44 ± 34.5 | 63.85 ± 32.76 | 52.54 ± 28.29 | 41.33 ± 18.05 | 38.91 ± 23.46 | 64.42 ± 37.39 | 51.06 |
| <i>Averrhoa carambola</i> | 43.15 ± 18.18 | 31.31 ± 10.42 | 48.83 ± 25.07 | 54.08 ± 28.47 | 72.59 ± 37.76 | 57.98 ± 34.48 | 67.33 ± 31.45 | 54.74 ± 27.45 | 53.65 ± 19.49 | 66.75 ± 31.21 | 55.04 |
| <i>Cassia fistula</i> | 34.27 ± 20.27 | 35.14 ± 11.34 | 41.31 ± 14.25 | 54.07 ± 28.91 | 68.98 ± 33.4 | 63.01 ± 33.39 | 60.8 ± 36.34 | 35.94 ± 16.12 | 46.89 ± 22.98 | 62.94 ± 32.26 | 50.33 |
| <i>Centella asiatica</i> | 38.56 ± 18.0 | 27 ± 13.78 | 22.08 ± 12.46 | 47.99 ± 19.95 | 68.76 ± 33.07 | 50.48 ± 23.06 | 57.16 ± 24.61 | 24.37 ± 15.74 | 40.96 ± 16.1 | 58.95 ± 24.92 | 43.63 |
| <i>Dillenia indica</i> | 34.76 ± 19.98 | 27.5 ± 18.7 | 33.62 ± 17.93 | 51.63 ± 25.85 | 69.07 ± 34.09 | 50.39 ± 19.7 | 63.01 ± 27.96 | 44.51 ± 22.13 | 51.51 ± 25.18 | 60.82 ± 26.06 | 48.68 |
| <i>Eucalyptus globulus</i> | 40.95 ± 16.88 | 37.19 ± 15.11 | 32.52 ± 19.36 | 57.6 ± 27.68 | 72.15 ± 32.4 | 63.85 ± 28.9 | 56.06 ± 27.8 | 39.75 ± 20.06 | 46.82 ± 22.4 | 55.63 ± 24.76 | 50.25 |
| <i>Haldina cordifolia</i> | 36.71 ± 17.23 | 32.09 ± 13.78 | 49.14 ± 23.26 | 53.07 ± 25.58 | 70.16 ± 30.99 | 59.31 ± 35.64 | 65.75 ± 30.3 | 46.64 ± 20.42 | 43.01 ± 24.2 | 55.2 ± 33.05 | 51.1 |
| <i>Justicia adhatoda</i> | 33.55 ± 17.24 | 31.08 ± 12.08 | 37.97 ± 23.38 | 56.01 ± 30.01 | 70.88 ± 33.16 | 59.78 ± 29.62 | 43 ± 25.08 | 29.95 ± 11.8 | 35.63 ± 15.95 | 59.67 ± 31.89 | 45.75 |
| <i>Lawsoniainermis</i> | 33.47 ± 21.8 | 37.68 ± 17.9 | 28.7 ± 18.71 | 55.62 ± 26.17 | 68.69 ± 34.41 | 52.08 ± 24.06 | 62.96 ± 32.67 | 36.06 ± 21.95 | 48.04 ± 23.28 | 74.28 ± 36.36 | 49.75 |
| <i>Murraya paniculata</i> | 40.49 ± 22.33 | 26.81 ± 15.82 | 32.51 ± 12.89 | 48.84 ± 22.56 | 66.98 ± 38.56 | 52.66 ± 26.47 | 56.41 ± 25.05 | 27.56 ± 16.92 | 40.01 ± 22.7 | 48.68 ± 20.57 | 44.09 |
| <i>Pithecellobium dulce</i> | 36.74 ± 17.06 | 22.13 ± 14.93 | 27.41 ± 15.72 | 49.81 ± 26.88 | 68.93 ± 32.57 | 39.7 ± 22.15 | 57.77 ± 29.79 | 26.96 ± 19.58 | 45.71 ± 23.17 | 53.32 ± 24.93 | 42.84 |

(continued)

Table 15.5 (continued)

| Growth inhibition of test fungal species (%) | | | | | | | | | | | |
|--|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|--------------|
| Test plants | Af | An | Fo | Gc | Ja | Mb | Na | Pk | Pu | Ro | Mean |
| <i>Pongamia pinnata</i> | 37.26 ± 18.25 | 35.5 ± 18.5 | 36.52 ± 15.68 | 52.18 ± 28.56 | 74.77 ± 31.27 | 64.74 ± 28.64 | 54.92 ± 26.67 | 49.39 ± 20.67 | 40.35 ± 22.88 | 56.95 ± 28.60 | 50.25 |
| <i>Tamarindus indica</i> | 19.31 ± 14.32 | 16.02 ± 11.33 | 33.06 ± 18.13 | 54.5 ± 27.95 | 68.88 ± 36.15 | 53.51 ± 24.5 | 52.4 ± 23.92 | 35.41 ± 19.01 | 33.32 ± 17.66 | 65.51 ± 32.01 | 43.19 |
| Mean | 37.31 | 30.85 | 36.11 | 53.73 | 71.44 | 57.73 | 58.37 | 38.36 | 43.92 | 60.25 | |

Af *Aspergillus flavus*, An *Aspergillus niger*, Fo *Fusarium oxysporum*, Gc *Geotrichum candidum*, Ja *Junghuhnia* sp. AK15, Mb *Mucor* sp. BOT15, Na *Neoscytallidium* sp. AM15, Pk *Penicillium* sp. KBS15, Pu *Penicillium* sp. UU15, Ro *Rhizopus oryzae*.

**Fig. 15.4** Comparative efficacy of antifungal activity of sixteen medicinal plants

recommendations against postharvest management of root, tuber, and bulb crops, but also the application of biofungicides would certainly be an endeavor for sustainable agricultural practices. As the plant kingdom is a rich reservoir of useful materials, the future of plant products in plant diseases control seems to be very bright.

It is suggested that the extracts of effective experimental plants of the present study may be utilized as postharvest dips for the control of storage decay of root, tuber, and bulb crops in order to avoid the side effects caused due to use of commercial fungicides. The findings of the present investigation will be of immense help to the agrochemical industries to prepare and biofungicides for sustainable agricultural practices.

Table 15.6 Comparative efficacy of three selected plants and four commercial fungicides on mycelia inhibition of ten fungal species

| Percentage of inhibition of mycelial growth | | | | | | | | |
|---|--------------|--------------|--------------|--------------|-----------------------|---------------------|----------------------|------------------|
| Test fungi | Ag | As | Av | Blitox-50 | Dhanustin (0.1 mg/ml) | Indofil (0.1 mg/ml) | Mancozeb (0.1 mg/ml) | Mean (0.1 mg/ml) |
| <i>A. flavus</i> | 48.33 | 38.97 | 43.15 | 21.74 | 100 | 19.63 | 82.5 | 54.99 |
| <i>A. niger</i> | 43.43 | 33.3 | 31.31 | 17.66 | 81.41 | 22.75 | 18.11 | 39.10 |
| <i>F. oxysporum</i> | 44.3 | 40.48 | 48.83 | 0 | 86.7 | 29.33 | 79.53 | 62.81 |
| <i>G. candidum</i> | 61.51 | 57.33 | 54.08 | 17.09 | 31.41 | 31.41 | 61.05 | 53.12 |
| <i>J. sp. AK15</i> | 77.08 | 74.97 | 72.59 | 17.79 | 76.24 | 72.5 | 80.43 | 78.024 |
| <i>M. sp. BOT15</i> | 77.05 | 63.48 | 57.98 | 24.8 | 11.83 | 33.17 | 30.75 | 51.73 |
| <i>N. sp. AM15</i> | 65.17 | 60.17 | 67.33 | 58.33 | 67.13 | 77.28 | 73.2 | 76.69 |
| <i>P. sp. KBS15</i> | 51.94 | 40.69 | 54.74 | 12.5 | 68.58 | 19.25 | 72.57 | 51.78 |
| <i>P. sp. UU15</i> | 44.21 | 52.09 | 53.65 | 56.25 | 100 | 100 | 100 | 78.57 |
| <i>R. oryzae</i> | 63.21 | 65.54 | 66.75 | 100 | 6.48 | 31.75 | 37.16 | 56.04 |
| Mean | 57.62 | 52.70 | 55.04 | 22.61 | 62.98 | 43.7 | 63.53 | |

Ag *Ageratum conyzoides*, As *Alstonia scholaris*, Av *Averrhoa carambola*

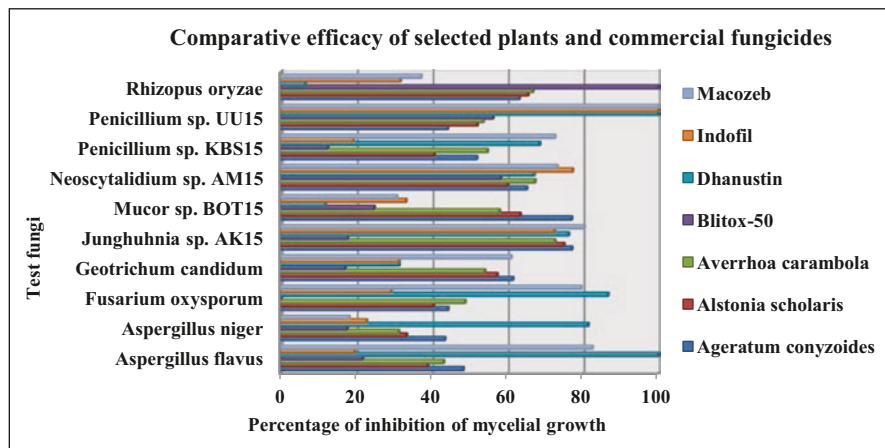


Fig. 15.5 Comparative efficacy of three selected plants and four commercial fungicides on mycelial inhibition of ten test fungal species

15.4 Conclusion

Indian economy is dependent upon agriculture, and agriculture has major problems of fungal diseases. Fungi can cause serious damage in agriculture, resulting in critical losses of yield, quality, and profit. Fortunately, agriculture has made tremendous

progress during the last century, and part of this progress has been the development of modern means of plant disease control. In particular, the introduction of chemicals as disease control agents has contributed to a substantial increase in crop production, to a smoothing of annual undulations in crop yields, and, ultimately, to today's high level of food security. Despite the fact that agrochemicals control diseases of crops, yet the constant and regular use of chemicals has resulted in detrimental effects to the environment and human health since 0.1% of agrochemicals used in crop protection reach the target pest leaving the remaining 99.5% to enter the environment to cause hazards to nontarget organisms including human. Broad-spectrum activity of agrochemicals has increased the risk of residue toxicity, prices, and pathogen resistances. This has urged agriculturists to look for available alternatives. Use of naturally occurring beneficial microorganism may be safe and alternative approach to compensate multiple effects of fertilizers and pesticides. In nature, microorganisms grow in various associations ranging from antibiosis, commensalism, parasitism, and symbiosis. This association can be exploited in biological control of plant pathogen. Advantages of biofungicides are that they are biodegradable in nature, have no effect on nontarget species, are cheaper than agrochemicals, are less toxic, and take less time to develop. The present study was made to evaluate the fungi toxic potential of plant-based bioformulation and to further develop a bio-entrepreneurship protocol of test plants for formulation of biofungicide to be used against storage pathogens associated with decay of root, tuber, and bulb crops in Odisha.

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Chapter 16

Plant Pathogenic Fungi and Their Phytotoxins as Bioherbicides



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16.1 Introduction

Legume crops are the second most important group of food plants and the major source of protein in the predominately vegetarian diet of the people of India. We have the distinction of being the world's single largest producer of legumes/pulses, having an area approximately of 20–24 million hectares under legumes. Madhya Pradesh is contributing a major percentage to the total legume production in the country. Pests especially weeds have increasingly become a major threat to sustainability of legume crops. Weeds are ubiquitous and continually changing pests in agriculture. They claim their own share of soil fertility and productivity at the cost of crop yield. They impose severe allelopathic effects on crops. Conventional

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methods of weed control have failed due to one or other reasons, while microbial management either directly or via their secondary metabolites has received significant momentum as evidenced by commercialization of several products such as mycoherbicides or biorationals.

Weeds are ubiquitous and are considered to be unwanted plants in agriculture and other settings. More specifically, the term is often used to describe native or nonnative plants that grow and reproduce aggressively. About 1800 of the weeds cause serious economic losses in crop production, and about 300 weed species are serious in cultivated crops throughout the world (Burnside, 1979; Holm, 1969; Holm et al., 1977).

Losses in crop yield due to weed infestation are very heavy, and reduction of 20–40% is not uncommon. The range of losses has been reported to be 30–100% in India, 20–40% in USSR, 8–24% in the USA, and 6–50% in the UK (Anonymous., 1979). A variety of microbes are pathogenic to plants. Recently, interest has developed in exploiting these pathogens and their phytotoxin as bioherbicides (Duke et al., 1991).

Some traditional synthetic bioherbicides have had no environmental impact, and some researchers believe that natural products are less toxic or at least more biodegradable. Traditional screening methods for the discovery of bioherbicides have also reached a point of diminishing returns (Wright et al., 1991). Fungal phytotoxic (CFCF) natural products may have novel mechanisms of action that are yet to discover potential new tools to combat bioherbicide resistance. The potential value of bioproducts as a source of compounds for development is widely underappreciated. Types of species of *Alternaria petroselini* (FCLW#23) are known to produce many phytotoxins, but only some have been proven to play roles in pathogenesis (Scheffer, 1992; Otani & Kohmoto, 1992; Montemurro & Visconti, 1992; Kohmoto et al., 1993). Several *Alternaria* sp. have a multitoxin system, and they produce more than one toxin which is important in disease causation.

All the compounds are now low-molecular-weight cyclodepsipeptides (Montemurro & Visconti, 1992; Ayer & Pena-Rodriguez, 1987; Ayer et al., 1987; Bains & Tewari, 1987; Buchwaldt & Jensen, 1991; Buiatti et al., 1987). Destruxin B ($C_{30}H_{51}, N_5O_7$, MW = 593) is the major phytotoxin produced by *Alternaria brassicae*. Two other phytotoxins, homodestruxin B ($C_{31}H_{53}N_{507}$, MW = 607) and destruxin B2 ($C_{29}H_{49}N_7$, MW = 579) are produced in much the same amounts. Desmethyldestruxin B ($C_{29}H_{49}N_{507}$, MW = 579) is also produced in trace quantities because of its phytotoxic activity. All these phytotoxins except dihydrotentoxin have phytotoxic activity.

Fungal phytotoxin is also a source of new bioherbicides (Duke, 1990, 1991; Lydon & Duke, 1989). This review will first address strategies for herbicide discovery. Then, host-specific from *Alternaria petroselini* (FCLW#23) fungi will be discussed with a review of the available literature as well as recent findings from our laboratory. Finally, the development of bioherbicides from phytotoxins of *Alternaria petroselini* (FCLW#23) will be discussed.

Conventional techniques of screening for resistance to pathogens suffer from several drawbacks such as scarce efficiency of selection, high time, and space

requirements. Environmental factors like temperature, Ph, effect of light, and C:N ratio may modify the host–pathogen interaction, often making some disease resistances very difficult to identify. In particular, partial resistances determined by small individual effects of different genes are not easily detected. Uniform inoculation and incubation of large numbers of plants is logistically problematic and may result in high frequencies of escapes. A further limitation of traditional techniques of screening for disease resistance sometimes consists in the lack of standardized infection conditions which cause discrepancies between greenhouse and field disease responses.

16.2 Phytotoxins and Culture Filtrates

The use of (CFCF) phytotoxins or crude extracts of pathogen for the individuation of disease-resistant plants has drawn considerable attention. Produced by many plant pathogens, they may represent practical and appropriate agents for the selection of disease resistance, if *in vitro* response to toxin and *in vivo* reaction to disease at the whole plant are well correlated. Toxins act at cellular level, allowing a uniform exposure of very large populations to the selection pressure. When added to *in vitro* culture, the multiple-step regimen based on gradual increase of the inhibitory level of toxins may often result in more appropriate resistant and vigorously growing cultures (Jones, 1990). This procedure may reduce the risk of physiological habituation of the host tissue to the selective agent.

In vitro selection involves extensive preliminary work, and it is currently restricted to toxigenic pathogens. The poor knowledge of biochemical and pathogenetic events occurring in infected plants is undoubtedly the major obstacle in its development.

When *in vitro* selection with pathogen or toxin is ineffective, artificial inoculation of regenerated weed in the greenhouse or other *in vitro* indicators become of particular relevance and may be used as an alternative (Megnegneau & Branchard, 1991; Storti et al., 1992; Buiatti et al., 1987). Although the screening on regenerated weed at the whole plant level is considerably laborious and requires a large space, increased resistance to specific pathogens, obtained by means of this technique, has been reported.

Some work has been done toward developing bioassay methods for symptom development in seedlings and intact plants, detached and attached leaves, detached branch bearing fruits of rapeseed, and effect on pollen germination (Ayer & Pena-Rodriguez, 1987; Bains & Tewari, 1987; Buchwaldt & Green, 1992). These bioassay methods are less sensitive than the ones based on the effects of other toxins on some physiological or biochemical processes, or on isolated cells and protoplasts (Gardner et al., 1986; Yoder, 1981; Yoder et al., 1977). Since many aspects of toxin research are dependent on the suitability of bioassay methods, there is a need to develop a sensitive, quantitative, reproducible, and easy-to-perform bioassay method.

16.3 Considerations in the Use and Development of Microbial Herbicides by *Alternaria petroselini* (FCLW#23)

There have been basically four types of strategies used to develop bioherbicides. The first type is random screening. Synthesized biochemicals are screened against leguminous weed species for phytotoxic activity. Compounds related to those with activity are then studied for structure/activity relationships to optimize bioherbicidal activity.

A secondary strategy is to design bioherbicides to attack with leguminous weeds a certain molecular site of action and to optimize this activity by the study of structure–activity relationships. This is sometimes called the biorational approach. Metribuzin resulted from this process (Wright et al., 1991). With the third strategy, herbicides may be designed similar to, but beyond the scope of, those patented by competitors. This, however, is unlikely to result in novel compounds or important advances in technology.

The end of the strategy is to isolate natural products from biological sources of fungi and screen for bioherbicidal activity (Tachibana & Kaneko, 1986; Mullner et al., 1993). Although random screening has to date been most rewarding as a source of herbicides, this strategy has approached the point of diminishing returns (Wright et al., 1991).

Sites attacked by natural products may have few and sometimes structurally complex but effective inhibitors. Since these sites may have relatively few effective inhibitors, the odds of discovery of a bioherbicide that attacks such a site by random screening may be very low. The known molecular sites of action of microbial phytotoxins differ in almost every case from the sites of action of commercial bioherbicides (Devine et al., 1993; Duke et al., 1991).

Bioproducts from plant pathogens often are more selective than synthetic compounds, perhaps due to their isolation from host-specific weed hosts. This can be a desirable property, as avoidance of injury to crop plants is a goal of synthetic herbicide development. However, if the compound is too selective killing only one or a few weed species, it may not be a viable herbicide candidate.

16.4 Potential Problems Associated with Bioherbicides

Herbicide discovery can be a much more complicated process with microbially derived herbicides than with synthetic herbicides. First, the microbe must be isolated from its source. A method of growing the fungus in large quantities must be devised. Often, microbes are not stable in culture, and many mutate and lose their virulence. The toxic compound must be isolated and purified prior to toxicity testing. Obtaining sufficient quantities of the toxin may be difficult.

Because of the small quantities obtained, microbioassays have been developed for screening natural phytotoxins. A leguminous weed bioassay has been developed, as well as bioassays involving radicle growth from small-seed plants, leaf discs, and growth and development of intact, small plants (Abbas et al., 1993a,

Table 16.1 Effect of different concentrations of cell-free culture filtrate of *Alternaria petroselini* (FCLW#23) on petriplate bioassay studies of leguminous weeds

| Seed germination inhibited (in %) | | | | | | | |
|------------------------------------|--------|--------|--------|--------|---------|---------|---------|
| Concentration of CFCF (%) | 24 hpt | 48 hpt | 72 hpt | 96 hpt | 120 hpt | 144 hpt | 168 hpt |
| Control | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| 5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| 10 | 0.0 | 0.0 | 0.0 | 0.0 | 10 | 25 | 30 |
| 20 | 0.0 | 0.0 | 0.0 | 10 | 25 | 51 | 65 |
| 25 | 0.0 | 5 | 20 | 35 | 40 | 68 | 75 |
| 50 | 45 | 65 | 88 | 100 | 100 | 100 | 100 |
| 75 | 60 | 75 | 90 | 100 | 100 | 100 | 100 |
| 100 | 80 | 95 | 100 | 100 | 100 | 100 | 100 |

Culture medium used = modified Richard's broth, Temperature = 28 °C, pH = 5

Table 16.2 Effect of different concentrations of cell-free culture filtrate of *Alternaria petroselini* (FCLW#23) on whole plant bioassay studies of leguminous weeds

| Whole plant inhibited (in %) | | | | | | | |
|------------------------------|--------|--------|--------|--------|---------|---------|---------|
| Phytotoxic damage (in %) | 24 hpt | 48 hpt | 72 hpt | 96 hpt | 120 hpt | 144 hpt | 168 hpt |
| Control | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| 5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| 10 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 10 |
| 20 | 0.0 | 0.0 | 0.0 | 10 | 30 | 45 | 50 |
| 25 | 0.0 | 0.0 | 10 | 35 | 50 | 80 | 90 |
| 50 | 30 | 65 | 80 | 90 | 100 | 100 | 100 |
| 75 | 50 | 75 | 85 | 100 | 100 | 100 | 100 |
| 100 | 60 | 85 | 95 | 100 | 100 | 100 | 100 |

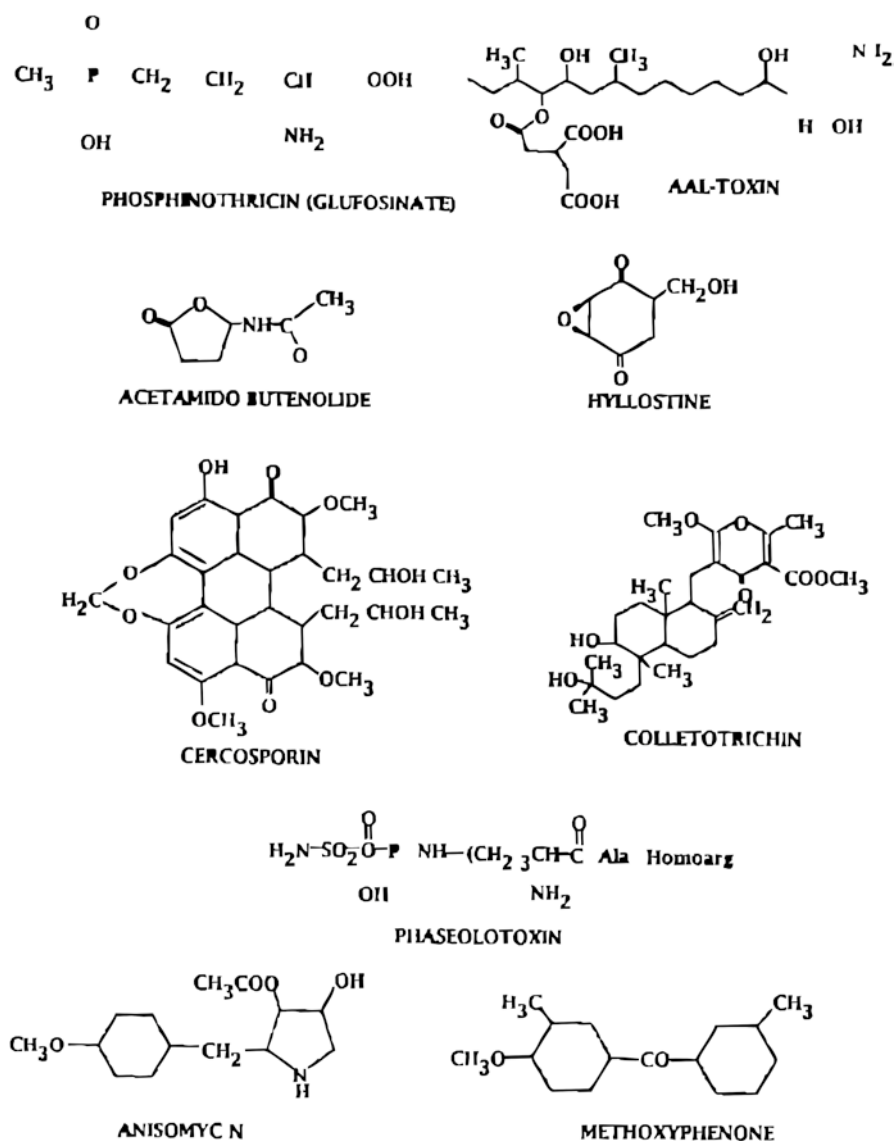
Culture medium used = modified Richard's broth, Temperature = 28 °C, pH = 5

1995; Tanaka et al., 1993). These assays may not be as accurate in predicting the potential of herbicides as greenhouse screening test methods because they may not reflect the effect on intact terrestrial plants in soil (Tables 16.1 and 16.2).

Following isolation of the phytotoxin, determination of its structure is necessary. Because of the complex nature of some of these compounds, this can be difficult. Despite the host specificity of the pathogen, it is possible that the same compound may have been recovered from a different organism. Ayer et al. (1989) showed that, in an herbicide discovery effort based on microbial sources, 72% of compounds, whose structures were determined, were already known. They had already excluded many known compounds through a phytotoxin profile database.

Once a compound is identified, another difficulty is producing it at an economically feasible price. Such compounds may be too complex to lend themselves to chemical synthesis. The cost of production by the microbes themselves may prove prohibitive. It is possible that these compounds may be used as leads to develop structural analogs which are simpler to synthesize (Ito et al., 1974).

Characteristics of microbial products may make their use as herbicides unsuitable due to factors such as extremely short half-lives or rapid degradation in certain environmental conditions. Although the phytotoxins may have activity when introduced by the plant pathogen, they may not be taken up by plant roots or leaf cuticles when applied as herbicides. They also may not be translocated to the site of action. The activity in leaf disc bioassays often does not predict the activity of an herbicide in whole plants (Nandihalli et al., 1992) (Figs. 16.1, 16.2, 16.3 and 16.4).



Cont...

Fig. 16.1 Structures of phytotoxin maintained in the test

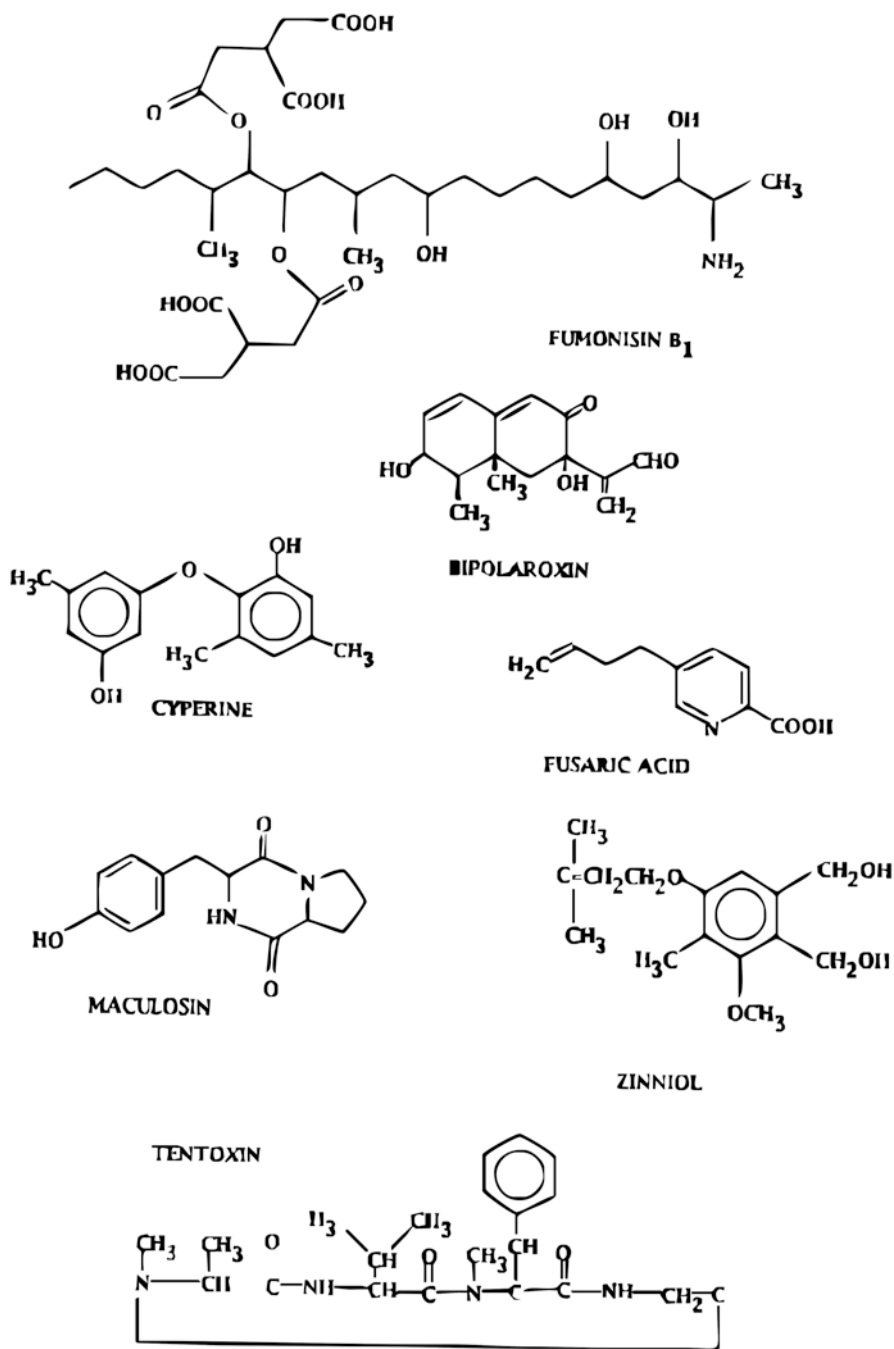


Fig. 16.1 (continued)

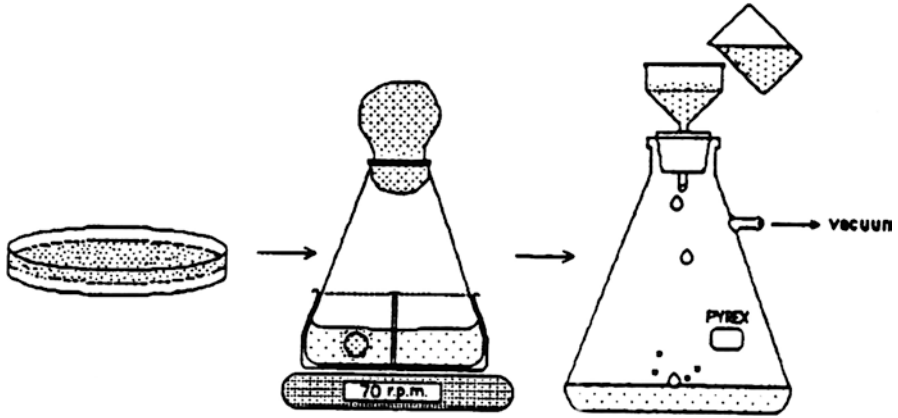


Fig. 16.2 Flow diagram of production of fungal cell-free cultural filtrate (CFCF)

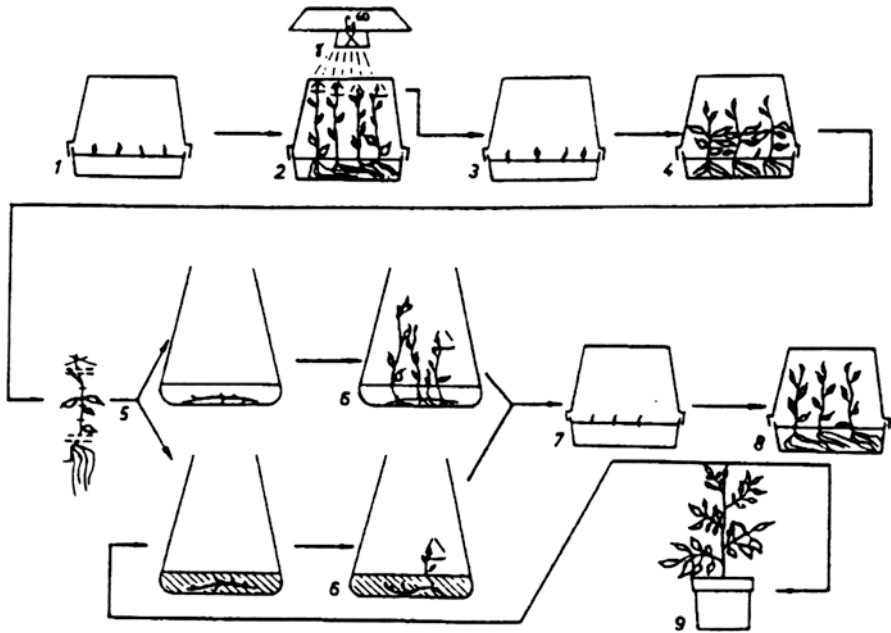


Fig. 16.3 Flow diagram of *in vitro* selection by bioassay

16.5 Use of Live Plant Pathogens *Alternaria petroselini* (FCLW#23) or Their Phytotoxins as Bioherbicides

In many instances, the plant pathogen causes the same effect on weeds as the phytotoxin it produces. It would be possible to use either the pathogen or the phytotoxin as an herbicide. Plant pathogens, however, have limitations that often make their use impractical with current technology (Julien, 1992; Zomer et al., 1993).

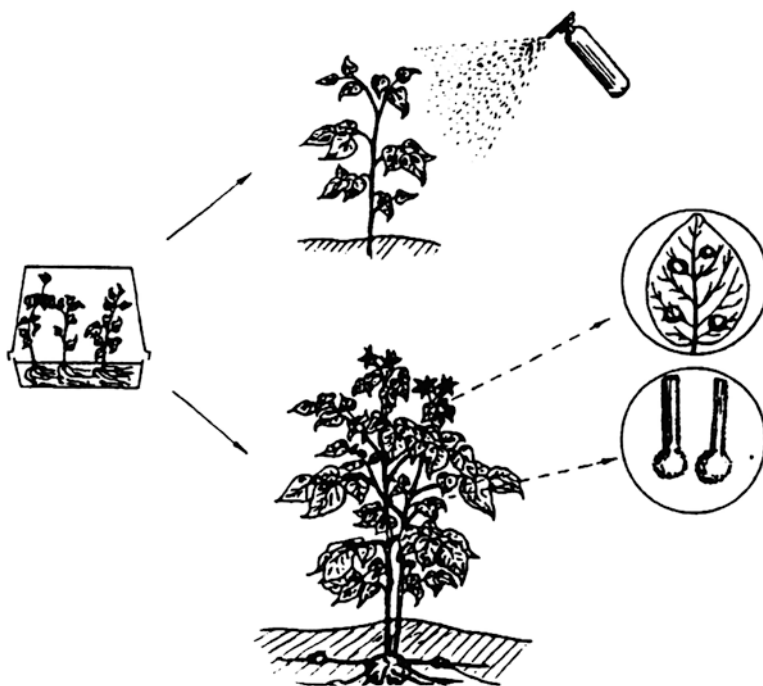


Fig. 16.4 Check of resistance of fungus by different procedures of *in vivo* artificial inoculation

Most plant pathogens that attack weeds are too host-specific, attacking only one or a very few weed species. As a multitude of weed species may affect the same crop, control of only one weed species would be of little interest to the farmer. However, toxins often have broader spectrums than the pathogens that produce them.

For example, AAL-toxin has been isolated from strains of *Alternaria petroselini* that are pathogenic only to certain cultivars of leguminous crops. However, Aflatoxin itself has a much broader range of activity and is toxic to Johnson grass [*Sorghum halepense* (L.) Pers.], black nightshade (*Solanum nigrum* L.), jimsonweed (*Datura stramonium* L.), prickly sida (*Sida spinosa* L.), hemp sesbania (*Sesbania exaltata* Rydb. ex A.W. Hill), and other formulations.

Living organisms also have the disadvantage of requiring special storage conditions such as temperature and humidity. They often have limited periods of viability and may have to be used within one season.

16.6 Host-Specific Phytotoxins

Although most known phytotoxins affect a variety of plant species, host-specific phytotoxins affect only one, or a very few, species (Scheffer & Livingston, 1984). Many are isolated from a pathogen of the affected species, and all known host-specific phytotoxins are from fungal pathogens.

Most known host-specific phytotoxins have been isolated from crop pathogens, and over twenty are known to exist. However, these toxins have not, in most cases, been tested for phytotoxicity to weeds.

AAL-toxin, produced by *Alternaria petroselini* (FCLW#23) lycopersici, is a hydroxylated long-chain alkylamine with a tricarboxylic acid moiety attached. AAL-toxin was initially reported to be host specific to only certain cultivars of leguminous crops, those with the Ase/Ase genotype. Heterozygous or Ase/Ase tomatoes are not affected.

We tested AAL-toxin on a variety of leguminous weed species. We found that AAL-toxin is phytotoxic to a number of weeds [4]. Therefore, AAL-toxin can no longer be considered to be truly host-specific (Tanaka et al., 1993; Duke et al., 1991; Abbas et al., 1993b). It is also likely that, with further testing, other so-called host-specific phytotoxins will be discovered to have a broader host range and may find applications in weed control.

To date, only one truly host-specific phytotoxin has been isolated from a weed pathogen, the cyclic dipeptide maculosin (Stierle et al., 1989; Strobel et al., 1990). Maculosin is derived from a pathovar of *Alternaria petroselini* (FCLW#23) and is host-specific for leguminous weeds. Maculosin is nontoxic to all other weed and crop species tested, including monocots and dicots.

Host-specific phytotoxins will probably be of less use as herbicides than those with broader ranges. Most crops have a combination of problem weed species, and, in most cases, it would be prohibitively expensive to develop and use a different bioherbicide for each weed species. However, relative host selectivity may be an advantage in some situations, and resistant crop plants might be developed or selected.

16.7 Development Considerations

The use of bioproducts as bioherbicides depends on the ability to produce them at a cost that makes the process profitable. If an herbicide is not cost-effective, it should not be produced despite its effectiveness, since these phytotoxins are produced by microbes, and fermentation is one of the methods that could be used for production. Fermentation may also not be feasible in some instances because of instability in the producing strain. However, recent developments in biotechnology have improved the yield and quality of the fermentation process; so costs may decrease with time as technology advances.

The other option for natural products is to produce them by chemical synthesis. Glufosinate, the other commercially viable herbicide derived from microbial sources, is synthetically produced. Some compounds are too complex for economical synthesis. However, active analogs may be produced that are simple to synthesize, as in the case of methoxyphenone which was modified from anisomycin (Yamada et al., 1974; Ito et al., 1974).

16.8 Conclusion

Plant pathogens remain a largely untapped reservoir of potential bioherbicides. Preliminary studies have identified a host-specific phytotoxin isolated from *A. petroselinii* (FCLW#23) that deserve further investigation.

Host-specific phytotoxins are less numerous and sometimes have broader spectra than reported when tested on leguminous weed. This may allow the application of some of these toxins or their analogs in leguminous weed management.

The study of phytotoxins produced by weed-specific pathogens is relatively new, and weed pathogens may be where the greatest chance of developing commercial herbicides lies. Because they are derived from weeds, such phytotoxins are more likely to be toxic to arid weeds less likely to be damaging to crops. However, natural products may seem to be more environment-friendly.

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Chapter 17

Mycotoxins: A Concealed Threat in Agri-Food Sector



Manisha Shukla and M. Mishra

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17.1 Introduction

Like all organisms, fungi manufacture byproducts of metabolism as they grow. The bulk of those low mass secondary metabolites are harmless to humans; however, some will deleteriously have an effect on human health, and these secondary metabolites are named as mycotoxins. These are toxic and have a major impact if they enter the food chain. Mycotoxins have attracted worldwide attention because of their impact on human health, immense economic losses, and domestic and foreign

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trade. It ought to even be remembered that some fungal metabolites are helpful to humans (e.g., antibiotics produced by fungi), whereas there are approximately 300–400 or so recognized mycotoxins, and solely ten are ordinarily ascertained in sickness, that are together called mycotoxicoses. Plant toxin contamination in foods and fodder has been changing into a world concern day by day. In line with Food and Agricultural Organization (FAO) reports, it's calculable that plant toxin affects nearly 25th of the world's crop annually and is inflicting immense agricultural product and industrial losses in billions of dollars (Alshannaq & Yu, 2017).

The best-known mycotoxins are the Aflatoxin family and the ergot alkaloids. This chapter addresses the kind of mycotoxins and role of plant's toxin-detoxifying agents.

There are four major aflatoxins (B1, B2, G1, and G2), but B1 is the most toxic. Most mycotoxins are chemical-stable and survive the food process. Several hundred different mycotoxins have been identified, but the most commonly observed mycotoxins that present a concern to human health and livestock include aflatoxins, ochratoxin A, patulin, fumonisins, zearalenone, and nivalenol/deoxynivalenol.

17.2 Characteristics of Mycotoxins (Cinar and Onbaşı 2018; Etzel 1999)

Mycotoxins are secondary toxic metabolites with a wide variety of chemical structures synthesized by fungi (mold) (Turnera et al., 2009). These secondary metabolites do not seem to be essential for vegetative growth and so could have very little or no primary function inside the organism.

Secondary metabolites are made once the organism enters the stationary phase, once the initial phase of ascension has declined. The metabolites made during this phase are typically related to differentiation and sporulation and might have profound biological activities that in some instances are exploited economically.

Mycotoxins are thought to be a sort of "chemical defence system" to guard mold from insects, microorganisms, nematodes, grazing animals, and humans (Etzel, 1999). Improper storage, transport, and selling may also cause mold growth and synthesis of mycotoxins (Etzel, 1999).

Factors that have an effect on mold growth and mycotoxin production are temperature, relative humidity, fungicides and/or fertilizers, interaction between the colonizing toxigenic fungal species, type of substrate and nutritional factors, geographical location, genetic necessities, and bug infestation (Tola & Kebede, 2016; Deligöz & Smith, 2005) (Fig. 17.1). A type of mold could produce one mycotoxin, a mycotoxin that will be synthesized by several molds.

Mycotoxins cause totally different degrees of toxicity consistent with exposure time, phytotoxin quantity, physiological condition, and sensitivity of the organism in humans and animals. In the majority of animal feed and products like wheat bran, noug cake, pea hulls, maize grain, milk and meat, and conjointly human food like cereal, fruit and vegetables, spice, etc., appearance of mycotoxins was discovered (Tola & Kebede, 2016). These foods create serious health risks in humans and every animal species.

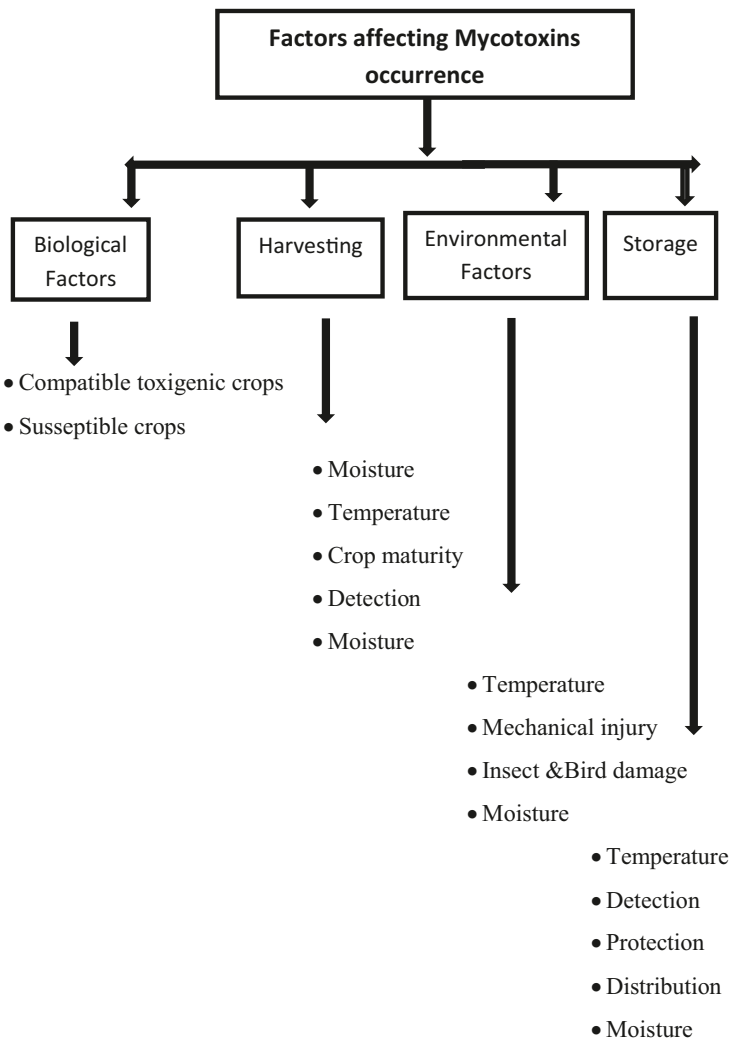


Fig. 17.1 Factors Affecting Mycotoxin occurrence in the agri-food sector

17.2.1 Difference Between Primary and Secondary Metabolites

A number of distinct variations are apparent between primary and secondary metabolites. First, they have been shown to possess an enormous variety of biosynthetic origins and structures that are not in general found among the primary metabolites.

Second, their incidence tends to be restricted to a tiny low variety of organisms and so will vary between isolated strains of constant species.

Finally, their production is characterized by the generation of groups of closely connected compounds which can have terribly totally different biological

properties. Vital samples of secondary metabolites embrace medically vital compounds like antibiotics, statins, cyclosporins, and ergot alkaloids. Agriculturally vital secondary metabolites embrace strobilurubin, associate antifungal compound, and plant hormones like gibberellin.

Secondary metabolites of molds particularly belonging to genera *Aspergillus*, *Penicillium*, and *Fusarium* are called Mycotoxins. Out of 300 identified secondary metabolites, only around 30 have toxic properties which are of some concern.

17.3 Toxicogenic (Mycotoxin-Producing) Fungi Can Be Distinguished into Two Groups

17.3.1 Preharvest

Fungi (such as *Fusarium*) which invade their substrate and produce mycotoxin on the growing plants before harvesting this is the category of field (preharvest) toxins. Aflatoxins and *Fusarium* toxins are included in this group.

17.3.2 Postharvest

These are fungi which produce toxins after harvesting and during crop storage and transportation. These toxins are named storage (or postharvest) toxins, and Ochratoxin A belongs to this group.

Mycotoxins were “discovered” following a sharp and fatal outbreak that occurred in 1960 on Turkey farms in Great Britain (Asao et al., 1963). This acute case led to the identification of aflatoxins and consecutively the link between molds, their toxins, and mycotoxicosis.

It is well known that mycotoxins may also cause chronic diseases in animals and may have completely different effects (hepatotoxicity, genotoxicity, nephrotoxicity, neurotoxicity, nephrotoxicity, immunotoxicity, etc.). It ought to be noted that toxicity might vary significantly at intervals in a structural group of mycotoxins and that the danger might not continuously result from the toxin itself but from its metabolites.

Toxic effects of mycotoxin once present at very low levels don't seem to be investigated till now.

Mycotoxin's area unit is usually present at the same time in staple. Moreover, complete feed is formed of various raw materials. Thus, animals are usually not exposed to at least one mycotoxin but to many toxins at the same time. Once mycotoxins are present at the same time, interactive effects are classified as additive, antagonistic, or synergistic.

Multiple contaminants may additionally occur attributable to the power of a given mold species to provide many varieties of mycotoxins in one type of food

ingredient, or else, many forms of mycotoxins are also found within the same food or feed containing completely different contaminated ingredients or raw materials.

Mycotoxins may occur in conjugated type, either soluble (masked mycotoxins) or incorporated into/associated with/attached to macromolecules (bound mycotoxins). These conjugated mycotoxins will emerge when metabolized by living plants, fungi, and mammals or when food is processed (Berthiller et al., 2009). Masked mycotoxins show lower toxicity than the initial, nonconjugated, toxic parent compounds (Gareis, 1994). The presence of mycotoxins in feed may create a risk to consumers if these toxins and/or their metabolites are excreted and accumulated in animal products like milk and meat. The presence of mycotoxins in milk could be a public health concern and should be regularly monitored. A great majority of mycotoxins found in feeds do not cause issues because either they are not excreted into milk like the bulk of *Fusarium* toxins, or they are excreted as a less toxic metabolite like ochratoxin α .

The mycotoxins of major concern as feed contaminants that are potentially removable from feed are mainly aflatoxins, ochratoxin A, and *Fusarium* toxins (trichothecenes like deoxynivalenol, diacetoxyscirpenol, nivalenol, T2-toxin/HT2-toxin, zearalenone, and fumonisins) (Figs. 17.2 and 17.3). One among the methods

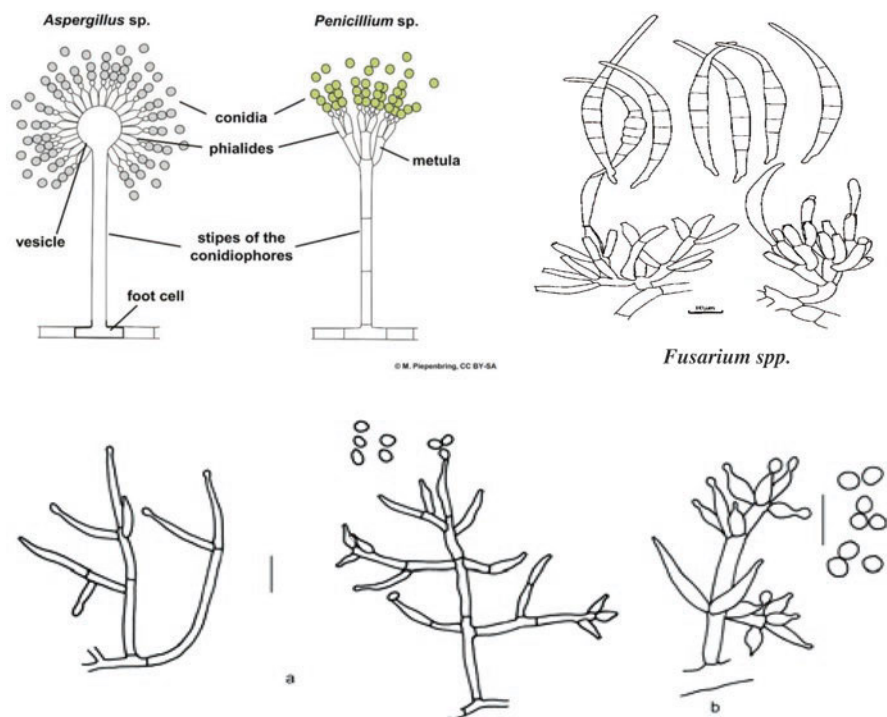


Fig. 17.2 Mycotoxin-producing fungi

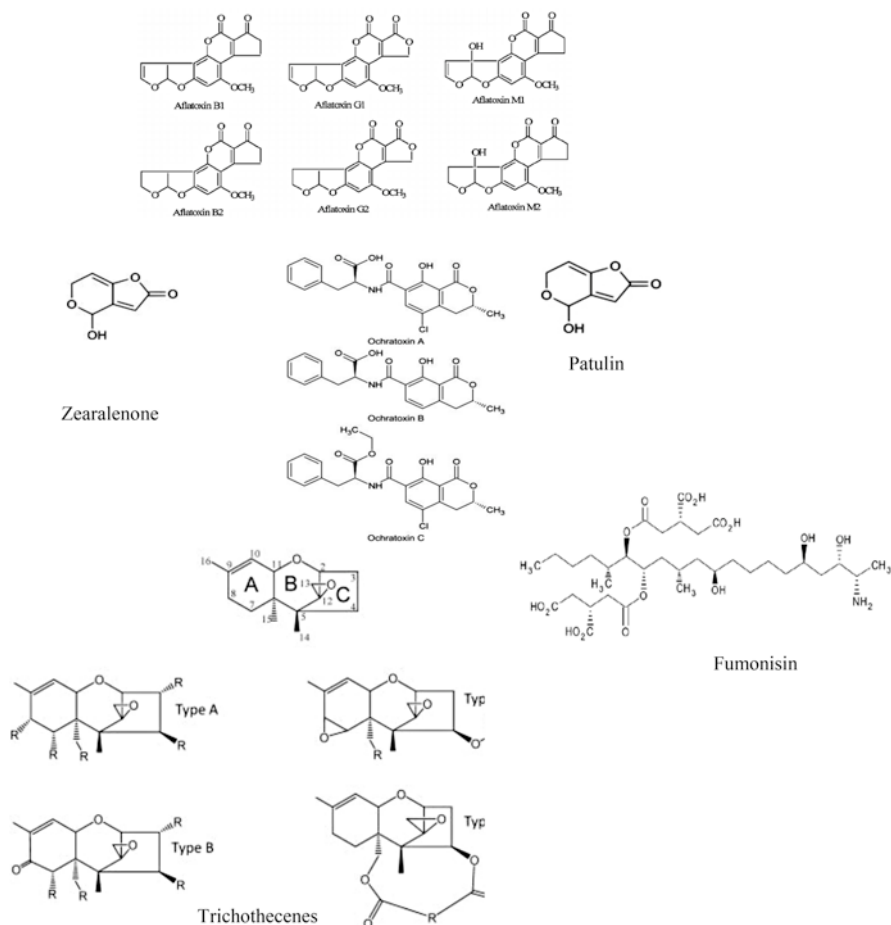


Fig. 17.3 Chemical structure of mycotoxins

for reducing the exposure to mycotoxins is to decrease their bioavailability together with numerous mycotoxin-adsorbing agents in the compound feed, which results in a reduction of mycotoxin uptake as well as distribution to the blood and target organs. Another strategy is the degradation of mycotoxins into nontoxic metabolites by using biotransforming agents like bacteria/fungi or enzymes.

These fungi frequently contaminate and produce toxins while growing on crops like corn and peanuts that are later on consumed by animals and humans. Consumption of meat and milk from cows that are exposed to the mycotoxins also can result in exposure of humans to the toxins, whereas acute aflatoxicosis may be a comparatively rare phenomenon, and eaten aflatoxins are notorious for their carcinogenicity as a result of chronic exposure. The mutagenic nature of aflatoxins is believed to be due to the DNA-damaging properties of aflatoxin metabolic

derivatives. The foremost common type of unwellness associated with dietary exposure to aflatoxins is liver cancer; but it should also be involved in different varieties of cancer. Because genus *Aspergillus* species are present within the surroundings, it's not possible to prevent foodstuff contamination with fungi. However, this contamination is reduced by the utilization of rigorous production, storage, and watching procedures.

Ergotism (also referred to as St Anthony's fire) may be an unwellness related to the uptake of cereals contaminated with the fungus ergot. This fungus, which infects the flowers of grasses and cereals, produces a variety of alkaloids, of which alkaloid is the best known. Alkaloid is said to be the psychoactive drug LSD, and uptake of cereals and cereal-derived merchandise like rye bread that are contaminated with the fungus may end up in serious symptoms of unwellness, like convulsions and gangrene. It's been instructed that mycotoxin may additionally exert a harmful result through inhalation, instead of by uptake. Most homes, workplace buildings, and factories provide several niches appropriate for the expansion of a myriad of filamentous fungi, together with genus *Aspergillus*, *Claviceps*, and *Stachybotrys* spp., and mycotoxins made by these species may contribute to the development referred to as "sick building syndrome." This syndrome has been related to a cluster of nonspecific symptoms (usually together with fatigue, minor metabolism issues, and headache) that are solely practiced inside a specific building. Poor air quality and ventilation, cleansing chemicals, and microbial contamination are the characteristic options of sick building syndrome. The role of fungi within the syndrome is a supply of considerable conjecture.

Mycotoxins are little and quite stable molecules that are very tough to get rid of or eradicate and that enter the feed chain while keeping their unhealthful properties.

Toxicogenic molds could develop underneath all climates on any solid or liquid supports as soon as biological process substances and wet (water activity A_w over 0.6) are present, hence the wide range of contaminated foodstuff substrates (Table 17.1). These toxins are found as natural contaminants in several foodstuffs of plant origin, particularly cereals, however, additionally fruits, hazelnuts, almonds, seeds, fodder, and foods consisting of, or factory-made from, these products and supposed for human or animal consumption (Binder & Binder, 1998).

Mycotoxigenic molds are uncontrollable due to their global presence in nature. Prevention of mycotoxin synthesis in all stages of food processing is an essential point for public health and economic reasons.

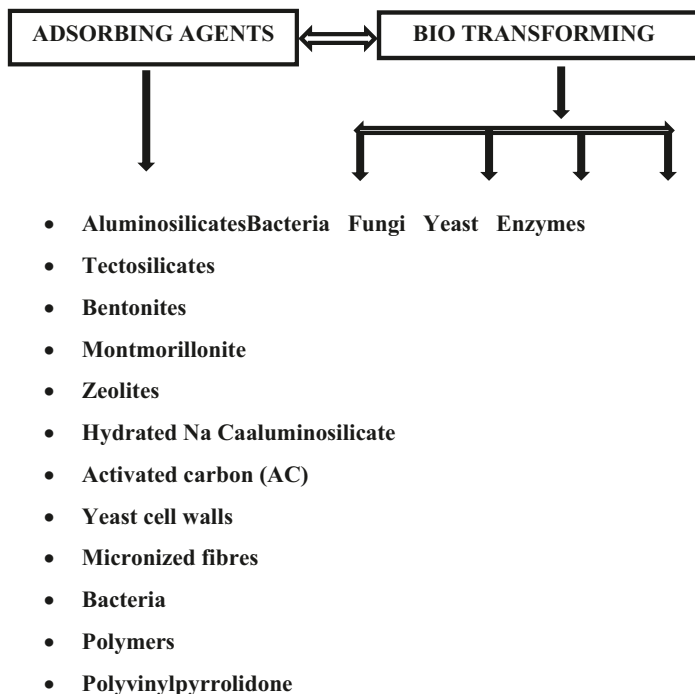
17.4 Mycotoxin-Detoxifying Agents

The Commission regulation (EC) No 386/2009 of May 12, 2006 outlines a new functional group of feed additives as "substances for reduction of the contamination of feed by mycotoxins: substances that may suppress or cut back the absorption promote the excretion of mycotoxins or modify their modes of action" (Kavin, 2018).

Table 17.1 Name of some important mycotoxin-producing fungi, susceptible foods, and mycotoxin effects on humans and animals

| Mycotoxins | Genus/species | Major food | Toxic effects and diseases |
|------------------------------------|--|---|--|
| Aflatoxin | <i>Aspergillus flavus</i> <i>A. parasiticus</i> <i>A. nomius</i> <i>Penicillium</i> | Cereals, feeds, oilseeds and pulp, coconut | Carcinogenic, hepatotoxicity, teratogenicity, decreasing immune systems, affecting the structure of DNA, hepatitis, bleeding, kidney lesions |
| Fumonisin | <i>Fusarium verticillioides</i> <i>F. culmorum</i> | Cereals, corn | Encephalomalacia, pulmonary edema, carcinogenic, neurotoxicity, liver damage, heart failure, esophageal cancer in humans |
| OchratoxinOTA | <i>Aspergillus Penicillium</i> <i>A. ochraceus</i> <i>P. nordicum</i> <i>P. verrucosum</i> | Cereals, herbs, oil seeds, figs, beef jerky, fruits, and wine | Kidney and liver damage, loss of appetite, nausea, vomiting, suppression of immune system, carcinogenic |
| Patulin | <i>Aspergillus terreus</i> <i>A. clavatus</i> <i>Penicillium</i> <i>Penicillium carneum</i> <i>P. clavigerum</i> <i>P. griseofulvum</i> | Silage, wheat, feeds, apples, grapes, peaches, pears, apricots, olives, cereals | Neural syndromes, brain hemorrhage, skin lesions, skin cancer, lung, mutagenicity, antibacterial effect |
| Trichothecenes (T2, DON, DAS, HT2) | <i>Fusarium Cephalosporium</i> <i>Trichoderma</i> <i>Fusarium oxysporum</i> | Cereals, feeds, silage, legumes, fruits, and vegetables | Immune suppression, cytotoxic, skin necrosis, hemorrhage, anemia, granulocytopenia, oral epithelial lesions, GIS lesions, hematopoietic, alimentary toxic aleukia (ATA), hypotension, coagulopathy |
| Zearalenone | <i>Fusarium</i> <i>F. graminearum</i> <i>F. culmorum</i> | Cereals, corn, silage, timothy grass, fodder | Carcinogenic, hormonal imbalance estrogenic effect, reproductive problems, teratogenic |

Table adapted from AycanCinarElif (2018) OnbustMycotoxins



Depending on their mode of action, these feed additives could act by reducing the bioavailability of the mycotoxins or by degrading them or remodeling them into less toxic metabolites.

Therefore, we will outline at least two main categories:

- *Adsorbing agents* One of the methods for reducing the exposure to mycotoxins is to decrease their bioavailability together with varied mycotoxin-adsorbing agents within the compound feed, which ends up in a reduction of mycotoxin uptake, moreover as distribution to the blood and target organs. Adsorbing agents are known as binding agents, adsorbents, or binders.
- *Biotransforming agents* Another strategy is the degradation of mycotoxins into nontoxic metabolites by mistreatment of biotransforming agents like bacteria/ fungi or enzymes. Substances that don't directly act with mycotoxins, i.e., inhibitor agents or immunostimulatory agents, are reviewed; however, they don't seem to be thought of *sensu stricto* as mycotoxin-detoxifying agents. However, we tend to be aware that such compounds could also be terribly economical for reducing the toxicity of mycotoxins.

17.4.1 *Mycotoxin-Adsorbing Agents*

Mycotoxin-adsorbing agents are large giant mass compounds that ought to be able to bind the mycotoxins in contaminated feed while not dissociating within the large number of animals. In this way, the toxin-adsorbing agent advances through the animal and is eliminated via the excreta. This prevents or minimizes exposure of animals to mycotoxins.

Mycotoxin-adsorbing agents are silica-based inorganic compounds or carbon-based organic polymers. The inorganic adsorbing agents presently on the market embrace natural clay products similarly as artificial polymers.

- *Aluminosilicates*: Silicate minerals are the most important category of mycotoxin sequestering agents, and most studies on the alleviation of mycotoxicosis by the utilization of adsorbing agents have centered on aluminosilicates. Inside this group, there is a pair of vital taxonomic categories: phyllosilicate taxonomic category and also the tectosilicate subclass.
- *Phyllosilicates*: They embrace bentonites, montmorillonites, smectites, kaolin-ites, and illites.
- *Tectosilicates*: They embrace zeolites.
- *Bentonites*: They are originally created from the weathering of volcanic ash in place (Ramos et al., 1996). They belong to the phyllosilicate group and are adsorbing agents with a superimposed crystalline microstructure and variable composition. Bentonites are typically impure clay consisting principally of montmorillonite. Because of their montmorillonite content, bentonites swell and form thixotropic gels (Diaz & Smith, 2005).
- *Montmorillonite*: It may be a superimposed salt that absorbs organic substances either on its external surfaces or inside its interlaminal areas (Ramos et al., 1996). Changed montmorillonite nanocomposite (MMN) may be a new sportive additive. Developed with nanomodification techniques, MMN features a sizable area, higher body, and stronger ion exchange activities beside a lot of active sites that build its nanoparticle impact straightforward to exert, and as a result, its surface assimilation effectivity is greatly increased.
- *Zeolites*: They are crystalline hydrous aluminosilicates of alkali and alkaline-earth cations characterized by an infinite three-dimensional structure. Zeolites are a bunch of silicates consisting of interlocking tetrahedrons of SiO₄ and AlO₄ (Kabak & Dobson, 2006; Ramos & Hernandez, 1997). Zeolites have large pores that give area for large cations like Na, potassium, and calcium, and they're characterized by their ability to lose and absorb water and exchange constituent cations while not harming the crystalline structure (Diaz & Smith, 2005; Papaioannou et al., 2002a, b). Clinoptilolite may be a natural mineral whose main application is the surface assimilation of heavy metals from liquid solutions (Papaioannou et al., 2002a, b).
- *Hydrated Na Ca aluminosilicate (HSCAS)*: It is probably the foremost studied mycotoxin-sequestering agent among the mineral clays (Diaz & Smith, 2005; Kabak & Dobson, 2006; Kleiner et al., 2001). It's a naturally occurring and

heat-processed Ca montmorillonite that's ordinarily used as an anticaking additive in animal feed (Galvano et al., 2001).

- *Activated carbon (AC)*: It may be a nonsoluble powder shaped by transmutation of many organic compounds and made by activation processes aimed toward developing an extremely porous structure (Kleiner et al., 2001). C is thought jointly of the foremost effective and nontoxic groups of sorbents and has been shown to be a tenacious sorb agent of a good sort of medication and cytotoxic agents. It has been ordinarily used as a medical treatment for severe intoxications since the nineteenth century (Wang et al., 2008). The sequestrant properties of AC rely on several factors together with pore size, area, structure of the phyto-toxin, and doses. Superactivated carbon differs from AC, wherein the particle size is reduced, thereby increasing area. The precise surface area of AC so varies from 500 m²/g to 3500 m²/g for superactivated charcoals (Ramos & Hernandez, 1997).
- *Yeast cell walls*, derived from the baker's yeast, are used as a dietary mycotoxin-adsorbing agent. Yeast cell walls consist virtually entirely of proteins and carbohydrates. The sugar fraction consists primarily of aldohexose, mannose, and N-acetylglucosamine. Glucans and mannans, the 2 main sugars, are found in equal concentrations in baker's yeast. Yeast mannan chains of varied sizes are exposed on the external surface and are coupled to cell membrane proteins (Huwig et al., 2001). The cell walls harboring polysaccharides, proteins, and lipids exhibit various completely different and straightforward accessible surface assimilation centers.
- *Micronized fibers*: They are obtained from completely different plant materials like cereals (wheat, barley, oat), pea hulls, apple, bamboo, etc. They're deep-rooted mainly of polysaccharide, hemicelluloses, and polymer and may be obtained in ultrafine (<100 μ) or less fine (>100 μ) fractions (Evans & Dawson, 2007).
- *Bacteria*: It has been recommended that cell membrane peptidoglycans and polysaccharides are the 2 most significant parts answerable for binding by carboxylic acid microorganism. Carboxylic acid microorganisms (LAB) are a group of gram-positive, acid-tolerant, typically nonsporulating microorganisms that have common metabolic and physiological characteristics. These microorganisms, sometimes found in rotten plants and drink products, manufacture carboxylic acid because of the major metabolic end-product of sugar fermentation. The most strains that comprise the work are *Eubacteria*, *Leuconostoc*, *Pediococcus*, *Lactococcus*, and true bacteria similarly as a lot of peripheral *Aerococcus*, *Carnobacterium*, *Enterococcus*, *Oenococcus*, and *Sporolactobacillus*. Strains of carboxylic acid microorganisms like *Eubacteria rhamnosus* strain GG and *Eubacteria rhamnosus* strain LC-705 are accustomed to take away mycotoxins. True bacteria thermophilus NG40Z and C5 have additionally been tested for their ability to detoxify mycotoxins (Aoudia et al., 1963).
- *Polymers*: Cholestyramine is an insoluble, quaternary ammonium ion exchange resin that powerfully binds anionic compounds. It's been used as medication in human for gripping digestive juice acids within the canal so as to scale back

sterol (Diaz & Smith, 2005). Polyvinylpyrrolidone may be an extremely polar amphiprotic chemical compound (Underhill et al., 1995).

17.4.2 *Mycotoxin-Biotransforming Agents*

Some studies have shown that some microorganisms have the power to degrade mycotoxins. Biotransforming agents embody bacterium, yeasts, fungi, and enzymes. The concept is for every plant toxin, or category of mycotoxins, to use enzymes that specifically degrade the poisonous substance into a nontoxic compound. Such enzymes are delineated in bacterium, yeast, or fungi. Refined enzymes have additionally been tested for this purpose.

(i) *Bacteria*

1. Gram-positive, anaerobic bacterium. The bacterial strain BBSH 797, a microbial feed additive, was isolated from rumen fluid. It's a gram-positive, nonspore-forming irregular rod living strictly anaerobic. It is 0.2 ± 0.4 and 1 ± 1.5 μm and occurs separately and in long chains up to 100 μm (Celik et al., 2000). Analysis of the 16s rRNA and therefore the guanine/cytosine content beside the specialized nutritional demands and different physiological characteristics indicates a new species of the genus *Eubacterium* (Fuchs et al., 2002).
2. Gram-positive, aerobic bacterium *Nocardia asteroides* may be a species of *Nocardia* that may be a genus of weakly staining gram-positive, catalase-positive, rod-shaped bacterium. It forms partly acid-fast beaded branching filaments. *Nocardia asteroides* are pathogenic: They'll cause nocardiosis, a severe respiratory organ infection in immunocompromised hosts. *Nocardia* are found worldwide in soil that's rich with organic matter (Binder et al., 2007).
3. *Corynebacterium* may be a genus of gram-positive, enzyme positive, nonspore-forming, nonmotile, rod-formed bacterium that is straight or slightly curving. Their size falls between 2.6 μm long and 0.5 μm in diameter. They're defined by high G:C content, with close phylogenetic relationship to *Arthrobacter*, *mycobacterium*, *Nocardia*, and *actinomycete*. They are cosmopolitan in nature and are mostly innocuous. Some are helpful in industrial settings like *C. glutamicum*, whereas others cause human sickness like *C. diphtheria*, the infectious agent liable for contagious disease. *Eubacteria rubrum*, named during this means due to the brilliant red color of its colonies, may be a nonacid fast bacterium capable of manufacturing massive quantities of lipids (Binder et al., 2007).
4. Mycobacteria are aerobic and nonmotile bacteria that are characteristically acid-alcohol fast. Mycobacteria are sometimes classified as gram-positive due to their lack of Associate in Nursing outer semipermeable membrane. *Mycobacterium* strain DSM 44556T was isolated with fluoranthene because

of the single carbon supply from the soil of a former coal gas plant, polluted with polycyclic aromatic hydrocarbons. The physiological properties, carboxylic acid pattern, and therefore the 16S ribosomal polymer gene sequence indicated membership to the genus *Mycobacterium*; however, they were totally different from all types of strains of mycobacterium species. This strain delineated a new species that the name *Mycobacterium fluoranthenorans* sp. nov was proposed for it (Binder et al., 2007).

5. *Rhodococcus erythropolis* is an aerobic, gram-positive, nonmotile, catalase-positive eubacterium that forms rods to extensively branched vegetative mycelium. It's an expedient infectious agent in immunocompromised patients.
6. *Curtobacterium* sp. strain 114-2 belongs to the genus of *Curtobacterium*. It is bacterium of the order actinomycetales. They're gram-positive soil organisms (Wu et al., 2009).
7. Gram-negative, aerobic bacterium *Flavobacterium aurantiacum* (NRRL B-184) may be a species of *Flavobacterium* that may be a genus of gram-negative, nonmotile, rod-shaped bacterium consisting of about 10 recognized species. *Flavobacteria* are found in soil and water in a variety of environments. Many species are famous to cause sickness in seafood (Binder et al., 2007).
8. *Pseudomonas fluorescens* may be a common gram-negative, rod-shaped microorganism. It belongs to the genus *Pseudomonas* bacteria. *P. fluorescens* has multiple flagella. It has a very versatile metabolism and may be found within the soil and in water. It's an obligate organism; however, certain strains are capable of exploiting nitrate rather than oxygen as a final electron acceptor throughout metabolism (Ueno et al., 1983).
9. *Alcaligenes* may be a genus of gram-negative, aerobic, rod-shaped bacterium. *Alcaligenes* species have been used for the economic production of nonstandard amino acids (Ueno et al., 1983). *Flavobacterium*, *Pseudomonas*, *Alcaligenes*, and *Bacilli* are often employed in a combination.

(ii) *Fungi*

1. *Aspergillus* species are highly aerobic and are found in most oxygen-rich environments, wherever they normally grow as molds on the surface of a substrate, as a result of the high oxygen tension. Commonly, fungi grow on carbon-rich substrates like monosaccharides and polysaccharides. *Aspergillus niger* may be a fungus and one of the foremost common species of the genus *Aspergillus*. It's a typical contamination of food. It's omnipresent in soil and is often reported from indoor environments.
2. *Aspergillus flavus* may be a common mold within the surroundings and may cause storage issues in stored grains. *A. flavus* is especially common on corn and peanuts, moreover as water damaged carpets. It can even be a human infectious agent, related to aspergillosis of the lungs and generally causing membrane, otomycosis, and naso-orbital infections. Several strains

produce important quantities of aflatoxin. *A. flavus* spores are allergenic (Megharaj et al., 1997).

3. *Aspergillus candidus* belongs to the genus *Aspergillus* and may be found in warm soils, grain, and within the secondary decay of vegetation (Binder et al., 2007). *Aspergillus parasiticus* may be a mold renowned to provide aflatoxin and generally found on black olives.
4. *Eurotium herbariorum* contains a moderately rapid growth rate with downy to powdery colonies. The *Eurotium* is usually found in tropical and semi-tropical zones. *Eurotium* may be a xerophilic fungus and is isolated primarily from soil, plants, stored grains, and house dirt (Megharaj et al., 1997).
5. *Rhizopus* is a genus of molds that has cosmopolitan thread-like fungi found in soil, decaying fruit and vegetables, animal excreta, and old bread. Mold grows well on general plant media. Mold species might overgrow and inhibit other fungi. Some structures are visible to the naked eye, i.e., sporangia seem macroscopically as black dots within the thick of white, cottony mycelium (Megharaj et al., 1997).
6. *Penicillium raistricki* belongs to *Penicillium* fungus genus that may be a genus of *Ascomycetes* fungi of major importance within the environment, food, and drug production. The mycelium generally consists of an extremely branched network of cell organs, septate, typically colorless hyphae (Binder et al., 2007).
7. *Rhinochadiella atrovirens* colonies are restricted, velvety, or lanose, usually slightly mucoid at the center. Conidiogenous cells are cylindrical, intercalary or free, 9–19 × 1.6–2.2 μm; rough rachis up to 15 μm long, with packed, flat, or butt-shaped, unpigmented conidial denticles. *Rhinochadiella* contains six to eight species, with two species of medical interest, *R. atrovirens* and *R. aquaspersa* (Nakazato et al., 1990).

(iii) Yeast

1. *Trichosporon mycotoxinivorans* belongs to the genus *Trichosporon*, which is characterized by the production of arthroconidia. Thirty-five species are delineated within the genus until now. This genus is monophyletic, on the idea of 18S and 26S rDNA sequences. *T. mycotoxinivorans* may be a yeast strain isolated from the hindgut of the lower termite (Mastotermitidae). The name of this recently isolated strain refers to a vital characteristic of *T. mycotoxinivorans* to detoxify mycotoxins like OTA and ZEA (Blackwell et al., 1999).
2. *Phaffia rhodozyma* and *Xanthophyllomyces dendrorhous* isolates: *Phaffia rhodozyma* was isolated within the 1960s. A minimum of 2 species appear to exist, together with the anamorph *Phaffia rhodozyma* and also the teleomorph *Xanthophyllomyces dendrorhous*. The yeast has attracted considerable biotechnological interest as a result of its ability to synthesize the antioxidant astaxanthin as its major pigment. This property has stimulated research on the biology of the yeast, moreover as development of the yeast as an industrial microorganism for astaxanthin production by fermentation (Molnar et al., 2004).

(iv) *Enzymes*

1. Protease A are enzymes which break down proteins (proteolysis), by reaction of the amide bonds linking amino acids together within the peptide chain. Proteases work best in acidic conditions. Peptidase A is obtained through fermentation processes from selected fungus genus *A. niger* strains (Peteri et al., 2007).
2. Pancreatin may be a mixture of many exocrine gland enzymes made by the gland cells of the pancreas. It's composed of enzyme, amylase, enzyme, and peptidase (Peteri et al., 2007).
3. Carboxypeptidase A typically refers to the pancreatic exopeptidase which hydrolyzes amide bonds of C-terminal residues with aromatic or open-chain facet chains. Most scientists within the field currently refer to this catalyst as CPA1 (Abrunhosa et al., 2006).
4. Epoxidases are enzymes that are able to biotransform epoxide groups into diene groups (Abrunhosa et al., 2006).
5. Lactone hydrolases are enzymes that catalyze the hydrolysis of lactone rings (intramolecular cyclic esters) to provide a hydroxyl group and a carboxyl group (Schatzmayr et al., 2006).

17.5 Impact of Mycotoxins on Livestock and Human

Mycotoxins are toxicant fungal products that are made once fungi grow in human and animal foods. A large range of food and drinkable things is contaminated with mycotoxins. Exposure to mycotoxins causes illness in humans and animals. Recent studies using biomarkers of exposure, internal dose, and adverse effects have shown that mycotoxins are underappreciated as a cause of illness. Some mycotoxins cross the placenta and are present within the craniate at birth, and others are excreted in milk. Some mycotoxins cause pathological processes (cancer) in humans and animals, thus causing kidney and medicine diseases. Animals are exposed to mycotoxins through consumption, inhalation, and/or skin contact. Several mycotoxins are potent in low doses, that is, quite little amounts of the compound will represent vital health effects (Fig. 17.4). Since filamentous fungi are common and expedient organisms, mycotoxins are widespread (Bennett & Krska, 2009). They are found in foods and feeds across the world and are of accelerating concern as potential contaminants within the indoor surroundings. Many of them are involved as warfare agents. Mycotoxins are to blame for cancers in addition to many alternative disorders moving the epithelial duct, urogenital, vascular, kidney, and nervous systems. Some mycotoxins are immunocompromising, thereby reducing resistance to communicable disease. It's estimated that two fifth of the world's crops, as well as several basic foods, are contaminated by mycotoxin-producing fungi (Eaton & Gallagher, 1994). The degree of biological effect caused by mycotoxins varies counting on the kind and supply, exposure route and dose, species status, and underlying subclinical

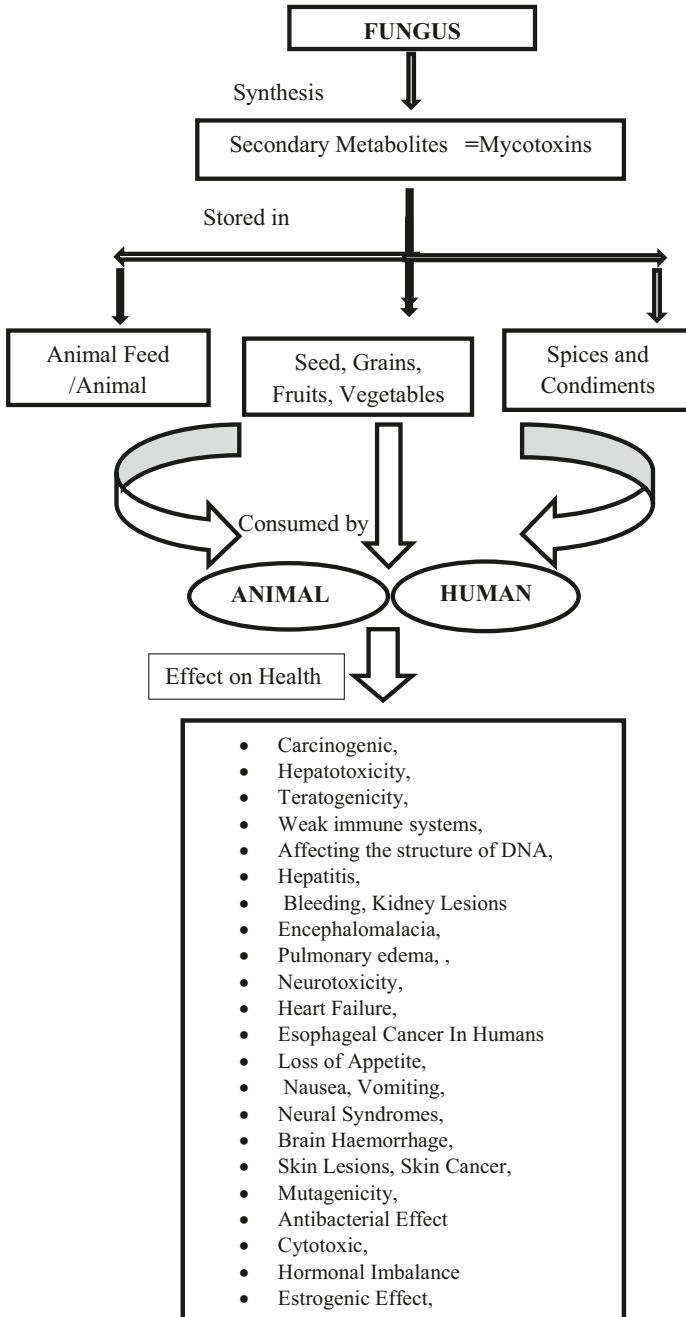


Fig. 17.4 Impact of fungal mycotoxin

conditions of the topic. For example, however, aflatoxin B1 (AFB1) is one in all the foremost potent liver carcinogens in humans (Fletcher & Blaney, 2016). Many countries have set limits for human and animal exposure to those mycotoxins known to cause adverse health outcomes. These laws vary by country, mycotoxin type, species, and food supply. International food laws are expected to be enforced within the close to future; however, adherence to such laws is difficult, particularly in countries where food provides are scarce. Higher quantities of mycotoxins are required in food and feed to produce adverse health symptoms in human and animals. It damages the system. These mycotoxins are restricted by limiting the consumption of contaminated food and feeds. Accessibility of mycotoxins information in numerous commodities is important for the adoption of safety against mycotoxins, and pesticides mycotoxin contamination is managed by spreading data regarding mycotoxins, predicting and preventing contamination, police investigation mycotoxins in grains, effective use of contaminated grain, and breeding crops for fungous resistance (Kam et al., 2007). Mycotoxin plays a big risk and food safety issue to the health and well-being of human and animals. Therefore, though the mycotoxin concentrations might considerably scale back throughout food process by physical treatment as cleansing and edge or by change of state, frying, baking, cooking, and extrusion, i.e., thermal process (Mitchell et al., 2014), nevertheless, it can't be fully eliminated. It's concluded that mycotoxins represent a big downside for the animal feed trade and on-going risk to the safety of the feed supply. Thus, because of the increasing public pressure, it's vital to produce an additional safe and eco-friendly way to manage plant toxin contamination. Use of microorganism antagonists as bacterium, yeast, and fungi is appropriate to cut back the chemical use and facilitate as biomanagement agents. Recent approach for safeguarding animals from harmful effects of mycotoxin contaminated feed is to use substances as mycotoxin binders to cut back the plant toxin contamination. These binders in addition to the diet scale back the absorption of mycotoxins from the alimentary canal and their future distribution to blood and target organs, so preventing or reducing mycotoxicosis in stock.

17.6 Conclusion

Mycotoxin is a well-known food safety risk that could be a threat to human and livestock health and has high economic significance within the food industry. Recently, the food industry has become tuned in to the new term for modified mycotoxins introduced by Rychlik et al. (masked mycotoxin) (Takahashi-Ando et al., 2002).

Prevention is the most vital and effective method in reducing fungal growth and mycotoxin production to ensure food safety, good agricultural practices (GAP) in the field, management measures of harvest and storage, physical processes (cleaning, milling, etc.), implementation of biotechnological application, biological management through the utilization of controlled atmosphere throughout storage, detoxification/degradation, and fermentation techniques are some preventive

measures to regulate mycotoxin prevalence. Physical ways (dehulling, washing, sorting, and cleansing of visible moldy seed) reduce completely different mycotoxin species in foods despite grain genre (Rychlik et al., 2014; AycanCinar Elif, 2018). Scudamore and Patel (2000) and Patel (2011) ascertained a discount of T-2 (62%) and HT-2 (53%) and DON (50%) in wheat seeds after cleaning. Scudamore and Patel also reported a 32% reduction in fumonisin levels in corn in an industrial enterprise (Scudamore & Patel, 2008). Moreover, milling is an important effect in the reduction of *Fusarium mycotoxins* in grains particularly wet milling of maize, which has shown to result in the degradation of mycotoxins (Scudamore & Patel, 2000).

Preharvesting is taken into account initially and one among the foremost necessary stages to forestall mold growth and mycotoxin synthesis. Many methods are out there for the production of healthy products and to cut back the mold formation at preharvesting, together with selection of plants per the soil structure and production capability, use of plant, that is immune to fungi and insects, irrigation time, create fertilization, and use of pesticides to forestall insect injury (Park, 2002).

Harvesting at the suitable time periods (low wetness and full maturity) is important for reducing the chance of a mycotoxin contamination since overmaturity creates sensitivity to mold growth. To boot, appropriate harvest instrumentation and procedures ought to be used, and crops ought to be dried once maturity to each grain wetness reaches safe levels (Barkai & Paster, 2008; World Health Organization, 2018).

17.7 Summary

Mycotoxins are substances produced by fungi and are poisonous to humans and other animals if consumed in sufficient quantities. Toxicity resulting from consumption of mold-contaminated foods has been reported to occur since ancient times. Mycotoxins produced (notably and aflatoxins) have caused many epidemics in the human populations of various countries throughout history and in the current age. They can severely impact our food and must be managed through programs of active surveillance and management.

Toxicity most commonly arises after ingestion; however, dermal and inhalation exposure routes can also result in adverse effects. Different mycotoxins and disrupt and critically damage organs, compromise the immune system, or cause cancer. The degree of biological effect caused by mycotoxins varies depending on the type and source, exposure route and dose, species susceptibility, and underlying subclinical conditions of the subject. However, mycotoxin contamination can be managed by disseminating knowledge about mycotoxins, predicting and preventing contamination, detecting mycotoxins in grains, effective use of contaminated grain, and breeding crops for fungal resistance. But, the efficacy of the preharvest and postharvest methods for mycotoxin reduction varies, and none of them confers absolute protection; therefore, the use of multiple strategies is desirable to minimize mycotoxin exposure.

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Chapter 18

Efficacy of Phytohormones on Mycotoxin Treated Maize Seeds (*Zea mays* L.)



Gajendra Prasad, Nitu Kumari, and Khwaja Salahuddin

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18.1 Introduction

Phytohormones are also known as plant growth regulators that influence several biological activities. These are important for growth and development of plants (Sembedner & Parthier, 1993). Phytohormones function as important chemical messengers as they modulate many cellular processes in plants and also coordinate different signaling pathways during exposure to mycotoxins (Vob et al., 2014).

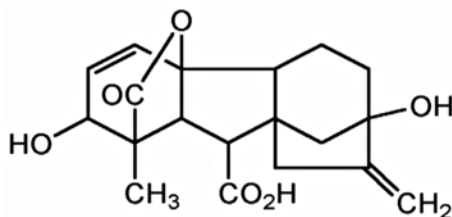
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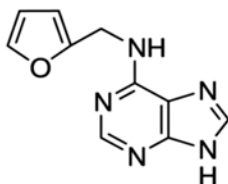
Maize is one of the most important staple food cereals grown throughout the world. It is commonly and directly consumed as processed food, indirectly as additive in most of food products and used as animal fodder (Lutz, 1994). Maize may be contaminated with storage fungi, producing mycotoxin that can be detrimental to health of human beings and animals. It is susceptible to a number of ear and kernel rots that can cause damage in humid areas.

Mycotoxins are naturally occurring toxins by certain fungi. These are found in food and can cause a variety of health effects, posing a serious health threat to both humans and live stocks. These fungi grow rapidly on a variety of natural substrates, and consumption of fungi-contaminated food can be detrimental (Kpodo et al., 1996). The crops that are frequently affected by *Aspergillus* spp. include cereals (corns, wheat, rice, and millet), oilseeds (pea nut, soya bean, sunflower), and spices (coriander, chilly, pepper, black pepper, turmeric, and ginger). The production of mycotoxin depends on high moisture content (20–25%) and high relative humidity (70–90%) (Shah et al., 2010).

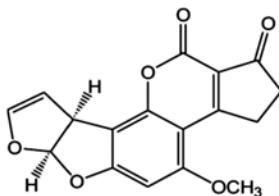
The fungi-producing mycotoxins fall broadly into two groups that invade before harvest commonly called field fungi, and those that occur only after harvest called storage fungi. The four major aflatoxins are called B₁, B₂, G₁, and G₂ based on their fluorescence under UV light. Aflatoxin B₁ is the most potent, natural, highly toxic, and carcinogenic secondary metabolite produced by *Aspergillus flavus* on agricultural crops (Lentopoloys et al., 2003). It is evident that aflatoxin is known for phytotoxicity with respect to seed germination and inhibition of root and hypocotyl elongation (Daskek & Llwellyn, 1983). AFB₁ was found at the highest concentration in contaminated food and feed. The different concentrations of AFB₁ inhibited chlorophyll, carotenoids, protein, and lipid contents and reduced the growth and germination of *Zea mays* L. and *Vicia faba* seeds (El-Naghy et al., 1999). It is properly documented that mycotoxin produced by specific filamentous fungi also causes significant reduction in crop yield and economic losses (Bhatnagar & Garcia, 2001). Aflatoxin content may vary with the season and storage time. It was observed that toxin contamination was higher during the rainy season and increased with increasing storage time (Ahmad, 1993). Aflatoxins are found in food chain in different pathways and make important detrimental damages in the metabolism of organisms (Hela et al., 2000). The seeds have been shown to be naturally contaminated with various levels of aflatoxin in the field (Ayalew, 2010) and during storage (Bilgrami & Sinha, 1992). Aflatoxins cause economic loss at all levels of food and feed production, processing, and distribution.



Chemical structure of Gibberellin (GA₃)



Chemical structure of Kinetin (Kn)



Chemical structure of Aflatoxin (AFB₁)

This article discusses hormonal action on maize seed, the action of mycotoxin, and finally the reversal effect with the application of phytohormones leading to the production of crops in the future.

18.2 Material and Methods

18.2.1 Collection of Healthy Seeds

The healthy seeds of maize (Madhuri-01) were obtained from Dayal Traders Manures and seed storage house, Darbhanga, India. The seeds were evaluated physically and were surface sterilized with 0.1% HgCl₂ for 2 min and then washed with sterilized distilled water.

18.2.2 Collection of Aflatoxin B1 and Phytohormones

The stock solution of AFB₁ was obtained from Sigma, USA. Pure phytohormones such as GA₃ and Kn were obtained from the local scientific stores, Darbhanga, India, and stored before analysis.

18.2.3 Preparation of Stock Solution

The stock solution of AFB₁ and phytohormones were prepared separately in ethanol from, which the dilution (i.e., 0.1, 0.25, 0.5, 1.0, and 2.0 ppm) was made in sterilized distilled water. Solutions of aflatoxin B₁ and phytohormones (2 ppm) were also mixed in different ratios (1:1, 1:2, 2:1, 1:3, 3:1 v/v) in order to record the combined effects.

The seeds were soaked initially in distilled water for 1 h and subsequently in different combinations of AFB₁ and phytohormones for 20 h. For each treatment, 100 seeds were taken in triplicates. The soaked seeds were then placed on moist blotter paper in sterilized petriplates and were kept for germination under automatic regulated seed germinator at 28 ± 2 °C (McLean et al., 1995; Sinha, 1990).

18.2.4 Seed Germination Index (GI)

It was calculated after 5 days of incubation according to the formula as given below:

$$GI = \frac{\text{Number of germinated seeds}}{\text{Number of seeds observed}} \times 100$$

The seedling growth (root and shoot lengths) was determined on the 7th day by measuring the lengths of radicle and plumule.

The data were analyzed statistically, i.e., t-test for seed germination and F-test for seedling growth. Statistical calculations were carried out using the ANOVA test (Dospekov, 1984).

18.2.5 Quantitative Estimation of Starch

About 200 mg freshly grounded sample was thoroughly shaken in 80% warm ethanol. After 5 min, the supernatant was decanted and centrifuged. Starch was estimated from the residue by adding 5 ml distilled water and mixing it thoroughly with 6.5 ml of 52% perchloric acid (prepared by adding 270 ml of 70% perchloric acid

into 100 ml of distilled water). This was followed by constant stirring for 10 min. After 15 min, again 20 ml of distilled water was added and centrifuged and then the supernatant was decanted. The extraction was repeated thrice and the supernatant was mixed together. The volume was raised to 100 ml by adding extra distilled water (stock solution). One milliliter of this stock solution was mixed with 10 ml at 0.1% anthrone reagent (760 ml conc. H_2SO_4 was diluted up to 1 l and 1 g anthrone dissolved) and heated at 100 °C for 12 min. The bluish-green solution was cooled at room temperature. Optical density was recorded at 630 nm under spectrophotometer. Readings were compared with the standard curve of starch prepared through similar procedure.

18.2.5.1 Assay of α -Amylase

It was assayed according to the method of Bernfeld (Bernfeld, 1955). One milliliter of starch solution was taken in experimental and control tubes. Approximately 0.5 ml of enzyme source (homogenate, i.e., 100 mg powder/ml acetate buffer, pH 4.8) was added in experimental and control tubes, but in the latter, the homogenate was previously dipped in boiling water bath for 5 min before being added. Both experimental and control tubes were incubated for 30 min at 37 °C. After incubation, both were dipped in boiling water bath for 5 min and subsequently cooled in running tap water. Three milliliter of KI solution (prepared by adding 25.4 mg I_2 and 400 mg KI in 100 ml of distilled water) was added in both tubes and mixed with the help of cyclomixer. The supernatant was used as crude extract for the extraction of enzyme. Colorimetric reading was taken at 620 nm, and difference of control and experimental reading gave the activity of α -amylase which was expressed in the value of optical density. Data recorded at each stage were the average of the three replicates.

18.2.6 Quantitative Estimation of Protein

It was done by the method of Lowry in both control and treated seeds (Lowry et al., 1951). Approximately 100 mg of seed flour were crushed in 100 ml of acetate buffer (pH 4.8) and centrifuged. The test solution (2 ml) was taken from the supernatant. To this, 10 ml of alkaline reagent (prepared by mixing 50 ml of 2% Na_2CO_3 solution in 0.1 N NaOH solution and 1 ml 0.5% $CuSO_4$ in 1% Na-K tartrate solution) was mixed thoroughly and was allowed to stand at room temperature for 10 min. One milliliter of diluted Folin–Ciocalteu reagent (1:3 in distilled water) was added. After 10 min, the extraction was added at 600 nm against the blank prepared by albumin.

18.3 Results and Discussion

The effects of phytohormones (GA_3 and Kn) and AFB_1 on individual and combined ratios were observed on the seed germination of maize (Table 18.1; Figs. 18.1 and 18.2). A significant fall in seed germination at all concentrations of AFB_1 was noticed, but the maximum inhibition (79%) in all parameters was observed at the highest concentration (2 ppm) of AFB_1 . The seed germination decreased by prolonged soaking intervals than the control treatment. When aflatoxin was mixed with GA_3 , then inhibition was reversed up to 20.40%. The same case was noticed when aflatoxin was mixed with Kn. The toxin functioned as anti-gibberellin by inhibiting DNA synthesis (Cavusoglu & Solusoglu, 2015). This is due to the fact that GA_3 is capable of breaking the dormancy and inducing seed germination and seedling growth.

The effects of phytohormones (GA_3 and Kn) and AFB_1 on the length of roots were observed (Table 18.2). The root length showed positive response to GA_3 and Kn and grew up to 12.11 cm and 11.76 cm, respectively, but when AFB_1 was applied separately, then the root length grew up to 3.42 cm showing 65% inhibition. But when AFB_1 and GA_3 were treated, the root length was reversed up to 8.92 cm, and similar response (8.65 cm) showed if AFB_1 and Kn were applied.

The effects of phytohormones (GA_3 and Kn) and AFB_1 on the length of shoots were observed (Table 18.3). The shoot length showed positive response to GA_3 and Kn and grew up to 8.89 cm and 8.76 cm, respectively, but when AFB_1 was applied separately, then the shoot length grew up only 2.25 cm showing 63% inhibition. AFB_1 restricted the growth of plant by inhibiting seed germination (Reddy et al., 2010). But when AFB_1 and GA_3 were treated, then the shoot length was reversed up to 4.94 cm, and similar response (4.83 cm) showed if AFB_1 and Kn were applied.

The effects of phytohormones (GA_3 and Kn) and AFB_1 on the content of starch were observed (Table 18.4). Starch is accumulated during seed development, and germination is characterized by its degradation. The dissolution of insoluble starch to soluble maltose-dextrins occurs in the presence of α -amylase which catalyzes the hydrolysis of glucosidic linkage (Prasad et al., 2018a). A fluctuation in the starch content was observed in maize seeds due to the treatment of phytohormones and

Table 18.1 Impact of hormones with respect to seed germination on AFB_1 -treated maize seeds

| Observations | Seed germination (Mean value \pm SE) | Difference with control | Percentage Inhibition |
|------------------------------|--|-------------------------|-----------------------|
| Control | 98 \pm 0.47 | – | – |
| GA_3 (2 ppm) | 100 \pm 1 | 2 | 2 |
| Kn (2 ppm) | 100 \pm 1 | 2 | 2 |
| AFB_1 (2 ppm) | 21 \pm 0.57 | 13.50 | 79 |
| AFB_1 + GA_3 (2 ppm v/v) | 78 \pm 0.47 | 22.27 | 20.40 |
| AFB_1 + Kn (2 ppm v/v) | 79 \pm 0.74 | 27.02 | 19.38 |

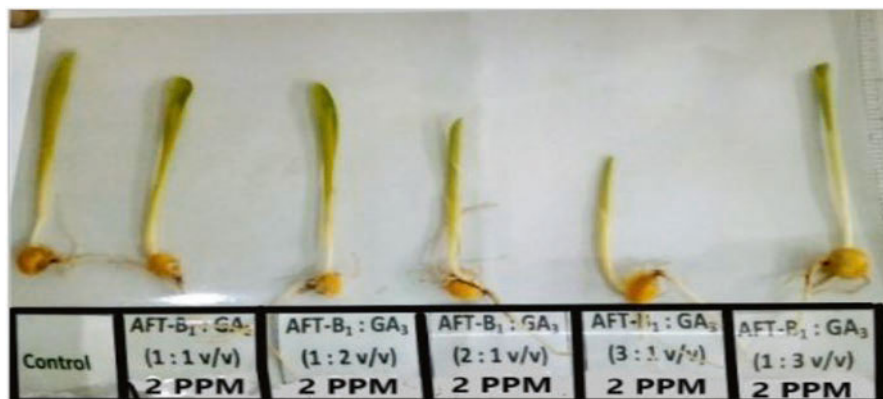


Fig. 18.1 Combined effect of AFB₁ and GA₃ (2 ppm v/v) on maize seedling growth

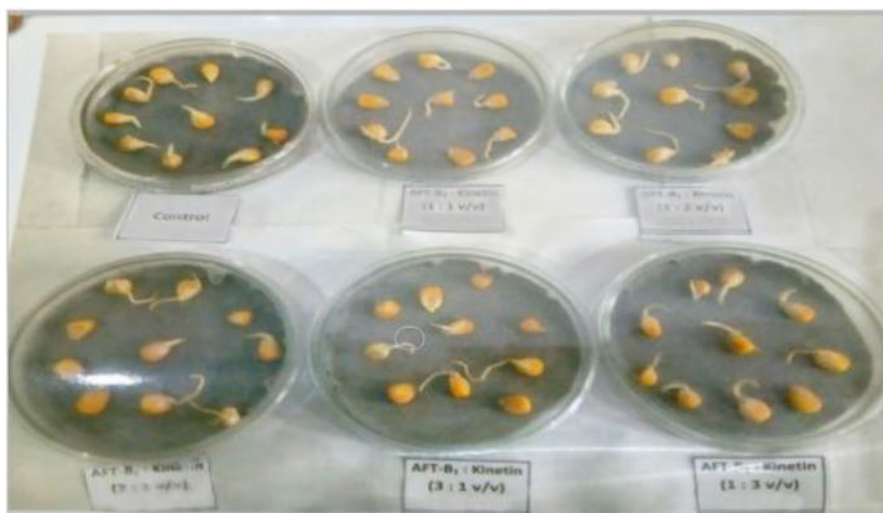


Fig. 18.2 Combined effect of AFB₁ and Kn (2 ppm v/v) on maize seed germination

mycotoxin. The maximum amount of starch (62.41 mg/100 mg) was recorded when maize seeds were treated with AFB₁ at 2 ppm concentration. The amount was decreased considerably up to 12.25 mg/100 mg and 13.12 mg/100 mg when AFB₁ was mixed with GA₃ and Kn, respectively.

The effects of phytohormones (GA₃ and Kn) and AFB₁ on the content of α -amylase were observed in different time duration (Table 18.5). The enzyme activity was completely lost in control set as well as GA₃ and Kn treatment in maize seeds after 90-min interval. However, enzyme activity was also lost in maize seeds due to treatment of GA₃ and Kn individually after 60-min interval. (Sakdeo 2016). The maximum inhibition as a result of AFB₁ treatment was observed. With increase

Table 18.2 Impact of hormones with respect to root length on AFB₁-treated maize seeds

| Observations | Root length (cm) | Difference with control | Percentage Inhibition |
|--|------------------|-------------------------|-----------------------|
| Control | 9.78 ± 0.12 | – | – |
| GA ₃ (2 ppm) | 12.11 ± 1.00 | 2.33 | 23.82 |
| Kn (2 ppm) | 11.76 ± 0.15 | 0.35 | 20.24 |
| AFB ₁ (2 ppm) | 3.42 ± 0.09 | 8.69 | 65.03 |
| AFB ₁ + GA ₃ (2 ppm v/v) | 8.92 ± 0.07 | 3.19 | 8.79 |
| AFB ₁ + Kn (2 ppm v/v) | 8.65 ± 0.04 | 3.46 | 11.55 |

Table 18.3 Impact of hormones with respect to shoot length on AFB₁-treated maize seeds

| Observations | Shoot length (cm) | Difference with control | Percentage inhibition |
|--|-------------------|-------------------------|-----------------------|
| Control | 6.15 ± 0.24 | – | – |
| GA ₃ (2 ppm) | 8.89 ± 0.08 | 2.74 | 44.50 |
| Kn (2 ppm) | 8.76 ± 0.09 | 2.61 | 42.00 |
| AFB ₁ (2 ppm) | 2.25 ± 0.08 | 3.40 | 63.00 |
| AFB ₁ + GA ₃ (2 ppm v/v) | 4.94 ± 0.12 | 0.12 | 19.00 |
| AFB ₁ + Kn (2 ppm v/v) | 4.83 ± 0.12 | 1.32 | 21.00 |

Table 18.4 Impact of hormones on starch content on AFB₁-treated maize seeds

| Observations | Starch content (mg/100 mg) |
|--|----------------------------|
| Control | 4.75 ± 0.13 |
| GA ₃ (2 ppm) | 0.93 ± 0.17 |
| Kn (2 ppm) | 1.14 ± 0.24 |
| AFB ₁ (2 ppm) | 62.41 ± 0.23 |
| AFB ₁ + GA ₃ (2 ppm v/v) | 12.25 ± 0.18 |
| AFB ₁ + Kn (2 ppm v/v) | 13.12 ± 0.10 |

Table 18.5 Impact of hormones on α-amylase content on AFB₁-treated maize seeds

| Observations | α-amylase activity | | | |
|--|--------------------|-------------|-------------|-------------|
| | 0 min | 30 min | 60 min | 90 min |
| Control | 0.73 ± 0.09 | 0.34 ± 0.09 | 0.14 ± 0.08 | – |
| GA ₃ (2 ppm) | 0.56 ± 0.12 | 0.24 ± 0.08 | – | – |
| Kn (2 ppm) | 0.62 ± 0.07 | 0.26 ± 0.12 | – | – |
| AFB ₁ (2 ppm) | 0.97 ± 0.07 | 0.61 ± 0.07 | 0.30 ± 0.07 | 0.12 ± 0.04 |
| AFB ₁ + GA ₃ (2 ppm v/v) | 0.65 ± 0.09 | 0.31 ± 0.10 | 0.13 ± 0.04 | 0.09 ± 0.57 |
| AFB ₁ + Kn (2 ppm v/v) | 0.73 ± 0.17 | 0.34 ± 0.08 | 0.21 ± 0.07 | 0.08 ± 0.04 |

in time incubation, the α-amylase activity gradually decreased in both germinated and nongerminated seeds.

The effects of phytohormones (GA₃ and Kn) and AFB₁ on the content of protein were observed in different time duration (Table 18.6). A highly significant fall in the level of protein in maize seedlings was observed during the treatment of different

Table 18.6 Impact of AFB₁ on protein content of maize seeds

| Concentration of AFB ₁ (ppm) | Protein (mg/100 mg) | Difference with control | Percentage Inhibition |
|---|---------------------|-------------------------|-----------------------|
| 0.00 | 8.43 ± 0.10 | – | – |
| 0.10 | 7.14 ± 0.02 | 1.29 | 15.30 |
| 0.25 | 5.88 ± 0.02 | 2.55 | 30.24 |
| 0.50 | 4.65 ± 0.03 | 3.78 | 44.88 |
| 1.00 | 3.19 ± 0.05 | 5.24 | 62.15 |
| 2.00 | 2.23 ± 0.08 | 6.20 | 73.44 |

Table 18.7 Combined effects of AFB₁ and GA₃ on protein content of maize seeds

| Concentration of AFB ₁ : GA ₃ (2 ppm v/v) | Protein content (mg/100 mg) | | |
|---|-----------------------------|-------------------------|-----------------------|
| | Amount (Mean value ± SE) | Difference with control | Percentage inhibition |
| 0:0 | 8.35 ± 0.04 | – | – |
| 1:1 | 4.86 ± 0.06 | 3.49 | 41.79 |
| 1:2 | 5.59 ± 0.05 | 2.76 | 33.05 |
| 2:1 | 3.87 ± 0.02 | 4.48 | 53.65 |
| 1:3 | 7.12 ± 0.05 | 1.23 | 14.73 |
| 3:1 | 3.19 ± 0.01 | 5.16 | 61.79 |

Table 18.8 Combined effects of AFB₁ and Kn on protein content of maize seeds

| Concentration of AFB ₁ : Kn (2 ppm v/v) | Protein content (mg/100 mg) | | |
|--|-----------------------------|-------------------------|-----------------------|
| | Amount (Mean value ± SE) | Difference with control | Percentage inhibition |
| 0:0 | 8.45 ± 0.04 | – | – |
| 1:1 | 4.76 ± 0.06 | 3.69 | 43.66 |
| 1:2 | 5.56 ± 0.05 | 2.89 | 34.20 |
| 2:1 | 3.85 ± 0.02 | 4.60 | 54.43 |
| 1:3 | 7.10 ± 0.05 | 1.35 | 15.97 |
| 3:1 | 3.17 ± 0.01 | 5.28 | 62.48 |

concentration of AFB₁. The percentage inhibition was found to be the maximum at (2 ppm) concentration of AFB₁. Protein is inhibited due to nonavailability of mRNA.

The biochemical effect of AFB₁ in combination with GA₃ on protein content was noticed when treated in various combination ratios (Table 18.7). The maximum and minimum inhibitions were observed when concentrations of AFB₁ and GA₃ are in the ratio of 3:1 and 1:3, respectively (Prasad et al. 2018b).

The biochemical effect of AFB₁ in combination with Kn on protein content was noticed when treated in various combination ratios (Table 18.8). The maximum and minimum inhibitions were observed when concentrations of AFB₁ and GA₃ are in the ratio of 3:1 and 1:3, respectively (Dilip et al., 2017; Pratiwi et al., 2015).

18.4 Conclusion

Maize is an important agricultural crop and has suffered from various fungal diseases resulting in the loss of its productivity. Phytohormones have played pivotal roles in increasing the length of root, shoot, and seed germination. Definitely aflatoxin has caused much damage to the yield of maize crop if applied separately. The reversal effect was noticed if phytohormones and aflatoxin were applied in combination and has gained much prominence with respect to the yield in such infected maize crops. If this method is applied by farmers, then it will be financially fruitful due to high yields as well as beneficial for the consumers with respect to the protein quality intake.

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Chapter 19

The Untapped Potential of Fungi in Phenol Biodegradation



Reshmi Sasi, Serin Zachariah, and T. V. Suchithra

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19.1 Introduction

Water pollution caused by phenolic pollutants is due to the expulsion of wastewater from industries such as petrochemicals, oil refineries, resin manufacturing, etc. (Villegas et al., 2016). Phenolic compounds are highly toxic even at lower concentrations and have severe effects on humans and animals. Due to the lethal effects of phenol, the US environmental protection agency (EPA) has included phenol and its derivatives in the list of high-priority pollutants (EPCRA, 2014). Removal of phenolic pollutants from the wastewater can be done with the help of numerous methods, among them biological methods gained more attention as it offers complete degradation of pollutants without the use of harmful chemicals and low cost of operation (Mahiudddin & Fakhrudin, 2012). Bacteria and fungi are the major classes of microbes that are being used for phenol degradation. Bacteria are the commonly employed microbes, while fungal degradation of phenolic pollutants is a

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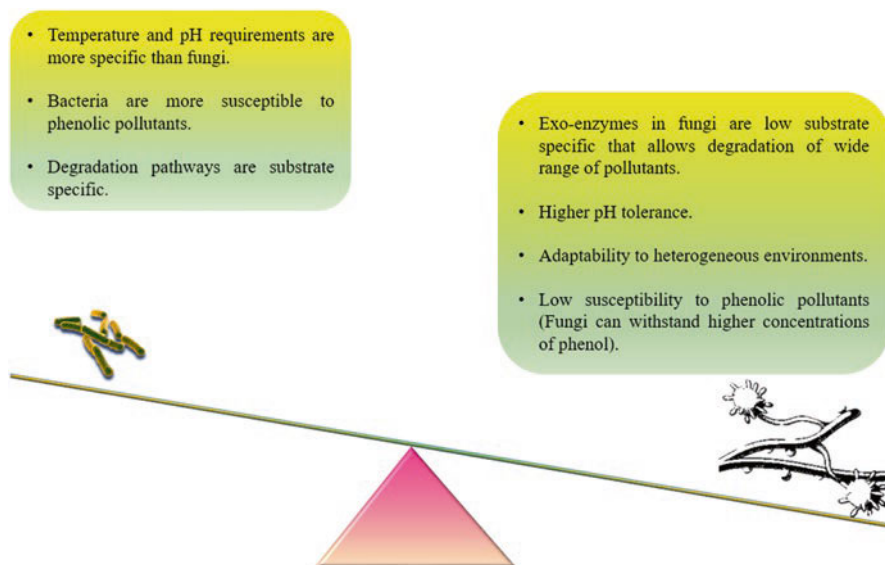


Fig. 19.1 A comparative representation of fungi v/s bacteria in phenol degradation

much less explored area. Fungi have so many advantages over bacteria such as they maintain stability in activity, especially at low pH, and are less susceptible to phenol toxicity when compared with bacteria (Fig. 19.1) (Yordanova et al., 2013). Filamentous fungi are the major groups used for phenol degradation, including white-rot fungi, *Penicillium*, *Fusarium*, etc. White-rot fungi are known for the degradation and mineralization of chlorophenol pollutants, and the process mainly depends on their degradation enzyme systems such as lignin degradation enzymes and cellular degradation enzymes. The lignin degradation system consists of laccase and peroxidase such as manganese peroxidase and lignin peroxidase (Valentín et al., 2013). Yeasts including candida maltose, *Oidium*, *Trichosporon cutaneum*, *Candida albicans*, etc. are good degraders of phenolic pollutants (Santos & Linardi, 2001).

19.2 Major Fungal Species for Phenol Degradation

Fungi are widely used as ideal candidates for the biodegradation of phenolic compounds. Their biochemical, metabolic, and ecological features are making them potential candidates for the biodegradation process. They have the capacity to extend their mycelial network, deeply penetrate the soil, and utilize the organic pollutants as their growth substrates. Despite other microorganisms, the unique characteristic of fungi is the presence of low substrate-specific catabolic enzymes like phenol oxidases in their extracellular matrix of the cell wall. Lignin peroxidases, manganese peroxidases, and laccases are the major ligninolytic enzymes secreted

by certain white-rot fungi. Besides this, some other intracellular enzyme systems consisting of cytochrome P450 monooxygenases (CYP) and glutathione transferases are also present in some fungal species. Adaptability to an immense range of habitats makes them more suitable for large-scale bioremediation of polluted environments (Arisoy, 1998; Pointing, 2001).

Different species of basidiomycetes are one of the major fungal groups involved in the degradation of organic pollutants. The low substrate-specific exoenzymes present in saprotrophic basidiomycetes and white-rot fungi make them best suited for mycoremediation (Lang et al., 1995). Because of the low substrate specificity, ligninolytic enzymes can also target some persistent organic pollutants like polychlorinated biphenyls (PCB), chlordane, and lindane, which have structures similar to lignin (Arisoy, 1998; Pointing, 2001; Mansur et al., 2017). *Phanerochaete chrysosporium* was known as the model white-rot fungus involved in the degradation of many pesticides such as dichlorophenol and pentachlorophenol by utilizing ligninolytic enzymes exuded from the extracellular matrix (Bhalerao & Puranik, 2007; León-Santesteban et al., 2016). *Anthracophyllum discolor*, *Ceriporiopsis subvermispora*, *Lentinus subnudus*, *Phlebia acanthocystis*, *Pleurotus ostreatus*, and *Lenzites betulina* are some other white-rot fungi effective in the biodegradation of organic pollutants such as pentachlorophenol (PCP), chlordane, and PCBs. *Phanerochaete sordida*, another saprotrophic basidiomycete, involves in the PCP degradation initiated by manganese peroxidase. Laccases are oxidases with copper as a cofactor and capable of oxidizing most phenolic and nonphenolic compounds. Laccase activity has been significantly observed in *Tinea versicolor* (Margot et al., 2013).

Mycorrhizal basidiomycetes such as *Gomphidius viscidus*, *Suillus variegatus*, and some edible basidiomycetes like *Agaricus bisporus* and *Lactarius piperatus* utilize the mechanism of biosorption to remove heavy metal contamination from phenolic compounds such as anthracene, TNT (2,4,6-trinitrotoluene), and crude oil. There are no enzyme systems or mediators involved in this degradation process (Margot et al., 2013). *Fusarium solani*, *Penicillium chrysogenum*, and *Scedosporium apiospermum* are some ascomycetes involved in nonligninolytic enzyme-mediated degradation. Cellulose and hemicellulose are the substrates to initiate the degradation process of some organopollutants (Tigini et al., 2009). *Fusarium incarnatum* is another ascomycete species involved in bisphenol degradation by using the laccase enzyme. Lipolytic enzymes and biosurfactants secreted by *Aspergillus niger*, *A. terreus*, *Cladosporium oxysporum*, and *Fusarium ventricosum* are used for the degradation of several pesticides (Bhalerao & Puranik, 2007). Yeasts such as *Oospora*, *Saccharomyces*, *Candida tropicalis*, *Debaryomyces*, and *Trichosporon cutaneum* were reported for active phenol degradation (Table 19.1).

Table 19.1 Different fungal species used for the biodegradation phenol and its derivatives

| Type of fungi | Fungi | Compound | References |
|----------------------------|--|--------------------------|---|
| White-rot fungi | <i>Phanerochaete chrysosporium</i> | Phenol and Chlorophenols | León-Santiesteban et al. (2016), Reddy and Mathew (2001) and Anastasi et al. (2013) |
| | <i>Phanerochaete sordida</i> <i>Ceriporiopsis subvermispota</i> <i>Lenzites betulina</i> | PCP | Lamar et al. (1994), Lamar and Dietrich (1990, 1992) and Tortella et al. (2008) |
| | <i>Anthracoophyllum discolor</i> | Chlorophenols | Acevedo et al. (2011) |
| Brown-rot fungi | <i>Laetiporus sulphureus</i> | Phenanthrene | Sack et al. (1997) |
| Tropical basidiomycetes | <i>Trametes villosa</i> , <i>Agrocybe perfecta</i> <i>Trichaptum bisogenum</i> <i>Peniophora cinerea</i> | PCP | Machado et al. (2005) |
| Mycorrhizal basidiomycetes | <i>Gomphidius viscidus</i> | Anthracene | Huang et al. (2010) |
| | <i>Suillus variegatus</i> | TNT | Meharg et al. (1997) |
| Ascomycetes | <i>Trichoderma harzianum</i> | Phenanthrene | Singh and Singh (2011) |
| | <i>Doratomyces nanus</i> <i>D. purpureofuscus</i> , <i>D. verrucisporus</i> , <i>Myceliophthora Thermophila</i> <i>Phomaeupyrena</i> <i>Thermoascus crustaceus</i> <i>Fusarium solani</i> , <i>Penicillium chrysogenum</i> , <i>Scedosporium apiospermum</i> | PCBs | Mouhamadou et al. (2013) and Tigini et al. (2009) |
| | <i>Aspergillus fumigatus</i> | Phenol | Fernandez-Fernandez et al. (2013) |
| | <i>Fusarium incarnatum</i> | Bisphenol | Chhaya and Gupte (2013) |
| Yeasts | <i>Oospora</i> <i>Saccharomyces</i> <i>Candida tropicalis</i> <i>Debaryomyces</i> <i>Trichosporon cutaneum</i> | Phenol | Harris and Ricketts (1962), Henderson (1961) and Neujahr and Varga (1970) |

19.3 Mycoremediation Pathways

Microbes including fungi degrade phenolic pollutants either aerobically or anaerobically (Fig. 19.2). The initial step in aerobic degradation is oxygenation, in which a monooxygenase enzyme phenol hydroxylase hydroxylates the aromatic ring at the ortho position of the pre-existing hydroxyl group to form 1,2-dihydroxybenzene (Catechol). This reaction requires a reduced pyridine nucleotide (NADH₂). Depending on the microorganism, the resulting catechol is further metabolized via two alternative pathways: β -keto adipate or ortho degradation pathway and metadegradation pathway. In the ortho degradation pathway, the enzyme catechol 1,2-dioxygenase cleaves the aromatic ring between the catechol hydroxyls to produce cis, cis muconate. Cis, cis muconate is further cleaved via β -keto adipate to

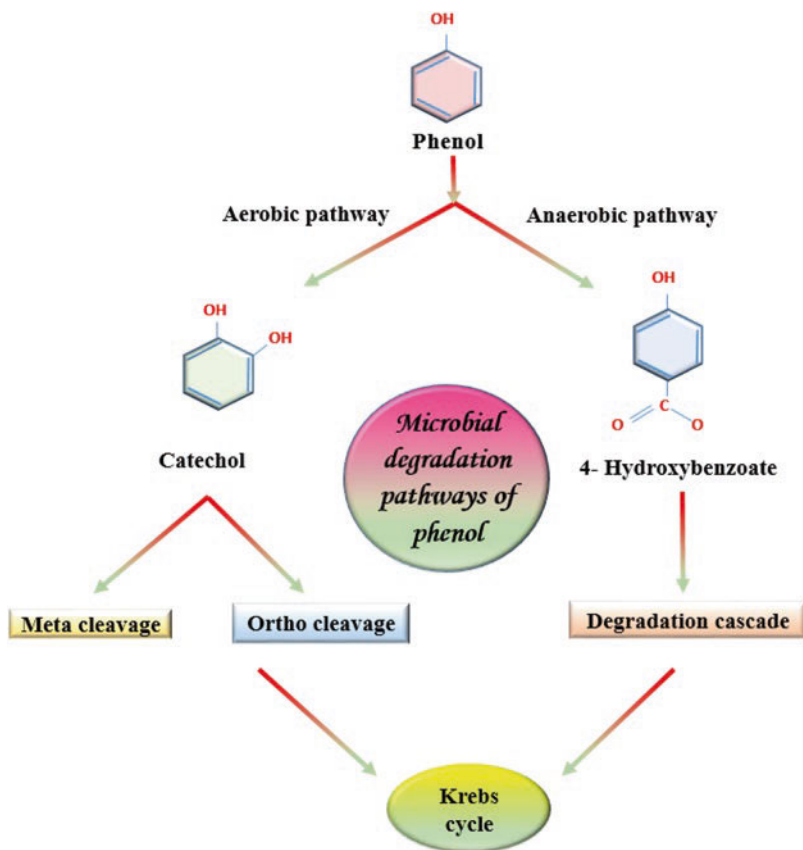


Fig. 19.2 A summary of microbial phenol degradation pathways

Krebs cycle intermediates (Evans, 1947; Kilby, 1948). In the metadegradation pathway, the estradiol fission or ring fusion happens near the two hydroxyl groups of catechol. The catechol 2,3-dioxygenases enzyme converts catechol into 2-hydroxymuconicsemialdehyde, which is further cleaved to form the Krebs cycle intermediates (Harwood & Parales, 1996; Stainer & Ornston, 1973).

Anaerobic phenol degradation is less explored than the aerobic process. This pathway shares similarities with the anaerobic benzoate pathway of *Paracoccus denitrificans*. The initial stage of the anaerobic pathway is the carboxylation of phenol at the para position by the enzyme 4-hydroxy benzoate carboxylase, which produces an intermediate called 4-hydroxybenzoate. The 4-hydroxy benzoate is further metabolized into protocatechuate by the enzyme protocatechuate 3,4-dioxygenase. It undergoes a cascade of metabolic cleavages and finally forms the Krebs cycle intermediates (Basha et al., 2010).

19.4 Fungal Immobilization Techniques for Phenol Degradation

Cell immobilization involves encapsulating cells in polymeric materials to keep them viable for a long-term biochemical process. The polymeric substance restricts the movement of cells by entrapping them inside. During phenol degradation, fungi can be immobilized using different immobilization techniques including encapsulation, entrapment, adsorption, etc. (Fig. 19.3). Several polymeric materials can be used for immobilization, including alginate, agar, collagen, polystyrene, and cellulosic materials. Among all the immobilization techniques, cell entrapment is the most practiced method for fungal immobilization during phenol degradation (Zaushitsyna et al., 2014; Karel et al., 1985).

19.4.1 Cell Entrapment

Cell entrapment is the safest and easiest method of immobilization. This method requires synthetic (polyethylene glycol conjugated PEG–lipid) or natural polymers (agar, alginate, carrageenan, agarose, etc.) to entrap cells and a cross-linking agent to strengthen the polymer network (Teramura et al., 2008). In the cell entrapment method, the cells are not in direct contact with the supporting material instead; they

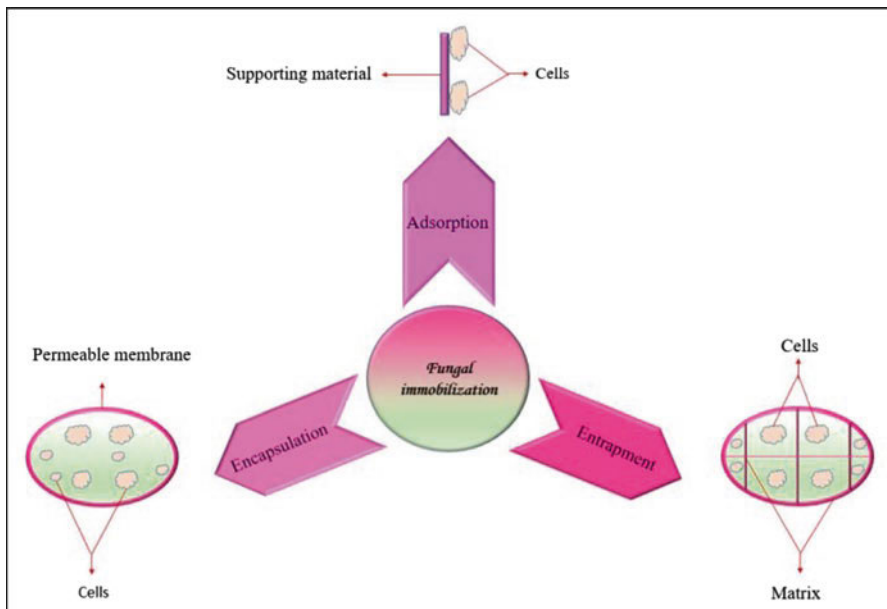


Fig. 19.3 Fungal immobilization techniques

are trapped inside the polymer matrix. The entrapment process involves mixing the cells with the polymeric solution and passing it through a needle and dropping it to a hardening solution (Gray et al., 2013).

19.4.2 Adsorption

As the name suggests, the cells are adsorbed or adhered to the surface of inert material. The adsorption mainly depends on the age of the cells and the chemical nature of the supporting material. The attachment is stabilized and maintained by various forces of attraction such as electrostatic, van der Waals, hydrophobic, and hydration forces. Organic and inorganic supporting materials are available for immobilization. Organic materials include DEAE-cellulose, DEAE-sephadex, and starch, and inorganic materials include silica gel, glass, alumina, etc. (Gray et al., 2013; Cassimjee et al., 2014).

19.4.3 Encapsulation

The encapsulation method is an altered entrapment method where a permeable membrane is used to encapsulate cell suspension. The membrane protects and supports cells during immobilization. It acts as a capsule and the liquid with active substances forms the core component and the membrane forms the outer covering. Different types of encapsulation procedures include emulsification, interfacial polymerization, coacervation, etc. (Karel et al., 1985).

19.5 Fungal Reactor Models for Phenol Removal

Large-scale biodegradation is carried out in large setups called bioreactors. Bioreactors provide optimum growth and degradation conditions for the microbes. Many fungal aerobic bioreactor models have been proposed and widely employed for removing phenol and its derivatives from wastewater. Fluidized bed reactors, batch reactors, bubble column reactors, stirred tank reactors, etc. (Fig. 19.4) are the commonly used models. Holladay et al. in 1978 reported the biodegradation of phenol using an anaerobic packed-bed reactor. They compared phenol degradation efficiency in different bioreactors, including stirred tank, packed bed, and fluidized bed reactor. They could find that the degradation efficiency is higher in stirred tank reactors when compared to packed bed and fluidized-bed bioreactor. The degradation rate inside the bioreactor majorly depends on the biomass development, flow rate, and feed concentration (Holladay et al., 1978).

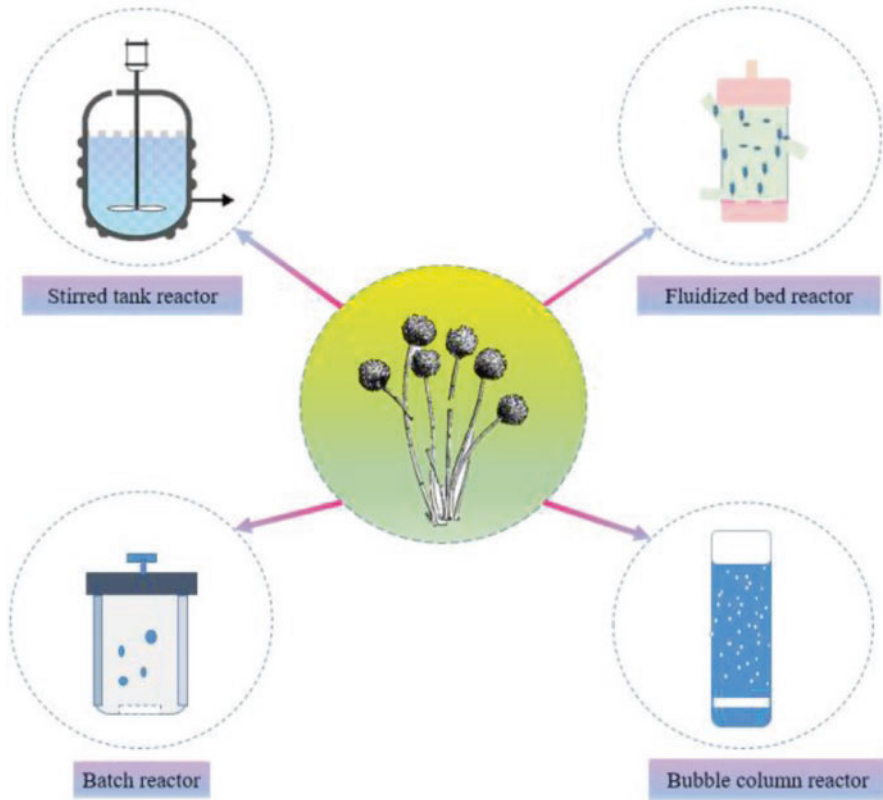


Fig. 19.4 Different bioreactor models for mycoremediation

19.6 Entrepreneurial Aspects of Fungal Bioremediation

Fungal bioremediation is the most suitable green route method for treating contaminated sites with recalcitrant pollutants. It gains more industrial attention among other microbes due to its higher pollutant removal efficiency and less substrate specificity. The environmental factors will significantly influence the industrial aspects of fungal bioremediation. Delayed lag phase, increased sludge generation, difficulty in bioprocess control, and monitoring are the major challenges faced during the scale-up process. The use of enzymes directly into the reactors over fungal biomass will reduce the bioremediation time with no lag phase, minimal sludge generation, and easy process control. Most of the enzymes used for biodegradation are highly expensive and have lesser stability with low shelf life. To tackle this situation, developments in whole-cell and enzyme immobilization techniques can be used to extend their stability and shelf life. These methods are preferably used in large-scale bioreactors like fluidized beds and cost-effective rotating biological

contactors with the reuse of enzymes and by immobilization of fungus (Jebapriya & Gnanadoss, 2013).

Another novel strategy, based on biopurification systems in promoting bioremediation of organopesticides through highly active biological mixtures, particularly white-rot fungi, was significantly highlighted in the advanced studies (Rodríguez-Rodríguez et al., 2013). Mixed filamentous inoculum in a continuous large-scale bioreactor can be used for more sustainable and ecofriendly bioremediation of sewage sludge (Rahman et al., 2014). One of the recent advancements in this area is the introduction of novel permeable reactive biobarriers of *Trichoderma longibrachiatum* on nylon sponge to treat phenolic pollutants (Cobas et al., 2013).

To many entrepreneurs, fungal bioremediation opens a wide window of opportunities. So many start-up companies in the bioremediation sector enjoy the advantages of mycoremediation. Earth Fax engineering groups Pvt. Ltd. is one of the leading companies in the mycoremediation sector. They mainly focus on fungal-based soil remediation which employs the pollutant-degrading abilities of a group of white-rot fungi. They have performed the degradation of many phenolic pollutants including PCP under full-scale conditions with the use of white-rot fungi. Several online consultation companies such as Bloom, Mushroom Mountain, etc. also help and support people in their mycoremediation ventures.

19.7 Conclusion

Fungi are humans' age-old companions. In an entrepreneurial view, its uses are countless, starting from fermentation to bioremediation. The current chapter mainly embraces the role of fungi in phenol biodegradation. In detail, it discusses the essential pathways of fungal phenol degradation, major species used in phenol bioremediation, immobilization techniques for improved degradation, bioreactor models for industrial applications, and the entrepreneurial aspects of fungal remediation of phenol.

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Chapter 20

Mycobased Biorefinery for Gold Nanoparticles Production



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20.1 Introduction

Green synthesis has gained immense recognition as a reliable, sustainable, and eco-friendly strategy for the synthesis of a wide range of nanoparticles (Nasaruddin et al., 2021; Rana et al., 2020; Pei et al., 2017; Molnár et al., 2018). Accordingly, green production approaches are considered important strategies for the mass production of nanomaterials by eliminating the ruinous effects related to the traditional chemical and physical approaches. Gold nanoparticles are interesting because of their unique physicochemical properties along with their oxidation/corrosion resistance, biocompatibility, and stability, which offer a multitude of potential applications in the fields of sensors, electronics, biomedicine, and catalysis (Siddiqi & Husen, 2016).

The production of gold nanoparticles for potential entrepreneurship has progressed dramatically. Both the facile and large-scale productions have aided the reliability of this approach. Further, the wide applications of the products are growing in biomedicine among industrial interests. Fungi are widely studied in the fabrication of metal nanoparticles than bacteria because of their ability to tolerate high metal concentrations (Siddiqi & Husen, 2016; Hyde et al., 2019). Moreover, they

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can release high concentrations of redox agents which ultimately can aid the fabrication of the metal nanoparticles. Hence, fungal protein extracts can produce gold nanoparticles at a large scale and relatively fast for potential entrepreneurship. Several studies supported the formation of small sizes of gold nanoparticles by fungi (Kumar et al., 2018).

There are a few advantages in utilizing fungi such as the thick coating of organic shells because of the redox reactions by the extracellular redox enzymes which are capable to reduce the common gold (III) precursors under static conditions (Pei et al., 2017; Molnár et al., 2018; Siddiqi & Husen, 2016). Overall, the use of microorganisms in gold nanoparticles synthesis is gaining speed because of the negligible toxicity in medical applications. There are a few merits for using fungi for the synthesis of different morphologies (Pei et al., 2017; Molnár et al., 2018). Gold nanoparticles have been fabricated as trigonal, hexagonal, spherical, and other shapes using fungi. The isotropic and anisotropic morphologies permitted wide applications in agriculture, industrial, and medical fields. Bimetallic alloy nanoparticles such as gold-silver were also synthesized using fungi for medical applications (Kumar et al., 2018).

The entrepreneurship based on fungi, apart from the basic laboratory applications, is extremely diverse such as traditional Chinese medicine, control of insects and weeds, environment-friendly biofertilizers, growth-promoting hormones, and many more. Recently, gold nanoparticles fabricated using fungi have been used in food and cosmetic industries (Kumar et al., 2018). In this chapter, we will address the routes for the synthesis of gold nanoparticles using fungi and their applications for entrepreneurship. Highlights on the fundamental mechanisms of the reduction of gold (III) precursors by redox enzymes from fungi will be discussed.

20.2 Mycobased Synthesis of Gold Nanoparticles

Fungi play a pivotal role in tackling enormous global issues for sustainable development and large-scale production. The exploration of the significance of fungi in nanotechnology is considered tremendously important. In this regard, a plethora of fungi have been investigated as they were found to be potential players in the biogenesis of gold nanoparticles. Filamentous fungi have some exceptional advantages over other biosystems like bacteria and algae, because of their high metal tolerance and bioaccumulation capability (Sastry et al., 2003). Since fungi secrete extracellular enzymes effectively, they paved the way for the industrial production of enzymes and proteins (Castro-Longoria, 2012). Moreover, high wall-binding and intracellular metal uptake are the separate privileges of exploiting them in nanoparticle synthesis (Volesky & Holan, 1995). On top of all these, fungi could produce metal nanoparticles by utilizing reducing enzymes intracellularly as well as extracellularly (Ahmad et al., 2003). It is worth mentioning that the effortless propagation and the rapid growth rate are other advantages of fungi for the green choice for the large-scale production of metal nanoparticles (Castro-Longoria, 2012).

Marsili et al., reported that maturation of *Rhizopus oryzae* in the presence of HAuCl_4 solution resulted in an incremental color change of the mycelia over time from off-white to purple. After 48 h, the UV-VIS spectrum of the purple mycelia exhibited a peak at 535 nm, indicating the formation of AuNPs (Das et al., 2012a, b). Sastry et al., confirmed that fungus *Fusarium oxysporum* releases reducing agents into a solution that are responsible for the formation of gold nanoparticles. Moreover, the release of proteins from the biomasses has been confirmed by gel electrophoresis (Mukherjee et al., 2002). Sastry et al., observed that an absorption peak at 270 nm is vivid and is assigned for the aromatic amino acids. Notably, the absorption peak at 270 nm occurs due to tryptophan and tyrosine amino acids (Mukherjee et al., 2002). Fungi capable of producing gold nanoparticles in a wide variety of shapes and sizes are summarized in Table 20.1.

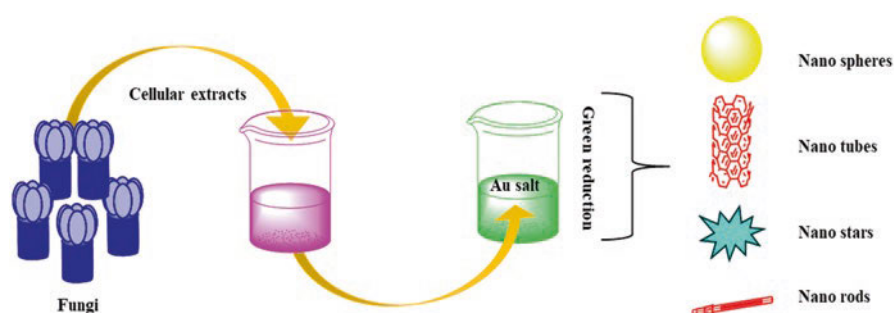
The morphology of gold nanoparticles has an enormous impact on their characteristics and applications. Even though spherical-shaped nanoparticles have been formed most frequently, metallic ion solution, incubation circumstances, and active biomolecules produced by the fungal organisms are the major factors to accomplish a wide variety of morphologies (Moghaddam et al., 2015). Interaction of different metal ions with *F. oxysporum* resulted in the fabrication of nanoparticles in a range of sizes and morphologies. Furthermore, nanoparticles synthesized by *F. oxysporum* had asymmetrical forms displaying a total quasi-spherical morphology in the size range 20–50 nm (Bharde et al., 2006). The utilization of fungal extract could regulate the morphology of gold nanoparticles at room temperature. Consequently, nanoparticles have been fabricated with the alteration of prime growth factors like gold ion concentration, pH, and response time (Gericke & Pinches, 2006; Das et al., 2010; Sun, 2002). The exploitation of an entophytic fungus *Colletotrichum* species for the synthesis of gold nanoparticles has resulted in the formation of different morphologies like rods, flat sheets, and triangles (Shankar et al., 2003) (Fig. 20.1).

20.2.1 Mechanism of Mycobased Synthesis of Gold Nanoparticles

As far as the mechanism of the extracellular and intracellular mycosynthesis is concerned, it is presumed that metabolites such as enzymes, proteins, polysaccharides, flavonoids, alkaloids, phenolic, and acids are secreted as an outcome of the fungus resistance to metal ions when exposed to different environmental stresses and are mostly responsible for the reduction of metal ions to metallic nanoparticles (Srivastava et al., 2019). It is quite noticeable that Au (III) ions penetrated or internalized into the cell membrane and reduced there during intracellular gold nanoparticles fabrication. Das et al. observed that Au (III) has a significant role in the expression of an oxidative stress response gene, which upregulated oxidoreductase enzymes. Also, the same research group discovered that *R. oryzae* reacted with

Table 20.1 Mycosynthesis of Gold nanoparticles

| | Fungi | Size (nm) | Morphology | Mechanism | References |
|----|------------------------------------|-----------|---|----------------------|-------------------------------|
| 1 | <i>Helminthosporium solani</i> | 15–20 | Sphere, rod, triangle, star | Extracellular | Kumar et al. (2008) |
| 2 | <i>Epicoecum nigrum</i> | 5–50 | Spherical, rod | Extracellular | Sheikhloo et al. (2011) |
| 3 | <i>Neurospora crassa</i> | >100 | Quasi-spherical | Intra, extracellular | Castro-Longoria et al. (2011) |
| 4 | <i>Aspergillus clavatus</i> | 48–72 | Triangle, sphere, hexagonal | Extracellular | Verma et al. (2011) |
| 5 | <i>Aspergillus niger</i> | 10–30 | Various shapes | Extracellular | Prakash and Soni (2012) |
| 6 | <i>Candida albicans</i> | 60–80 | Spherical | – | Owais et al. (2011) |
| 7 | <i>Fusarium oxysporum</i> | 8–40 | Spherical, triangular | Extracellular | Mukherjee et al. (2002) |
| 8 | <i>Rhizopus oryzae</i> | 16–25 | Spherical | Extracellular | Das et al. (2012) |
| 9 | <i>Fusarium solani</i> | 20–50 | Spherical | Extracellular | Gopinath and Arumugam (2013) |
| 10 | <i>Sclerotium rolfii</i> | 25 | Triangle, hexagonal, decahedral, rod, isotropic spherical | Extracellular | Narayanan Sakhivel (2011) |
| 11 | <i>Phanerochaete chrysosporium</i> | 10–100 | Spherical | Extracellular | Sanghi et al. (2011) |
| 12 | <i>Hormoconis resinae</i> | 3–20 | Spherical | Extracellular | Mishra et al. (2010) |
| 13 | <i>Yarrowia lipolytica</i> | – | Various shapes depending on Au (III) concentration | Intracellular | Pimprikar et al. (2009) |
| 14 | <i>Fusarium semitectum</i> | 10–80 | Spherical | Extracellular | Sawle et al. (2008) |
| 15 | <i>Saccharomyces cerevisiae</i> | 15–20 | Spherical | Cell wall, cytoplasm | Sen et al. (2011) |

**Fig. 20.1** Schematic representation of the synthesis of gold nanoparticles from fungi

concentration-dependent active biochemical detoxification when interacting with Au (III) (Das et al., 2012a, b).

AuNPs synthesis could take place in either the intracellular or extracellular space. Das et al. observed two reduction methods: (a) electrostatic interaction of Au (III) on the cell wall and (b) internalization of Au (III) into cytoplasm resulted in the synthesis of AuNPs with the aid of proteins or enzymes. Furthermore, Das et al. demonstrated the formation of AuNPs in both cell walls and the cytoplasmic region of *R. oryzae* by TEM micrograph (Das et al., 2012a, b). Even though the exact mechanism of the synthesis of AuNPs has not been elucidated, studies suggested that Au ions require NADH-reliant nitrate reductase enzyme for their reduction, which was secreted by the fungus in the extracellular environment (Rai et al., 2021; Dorcheh & Vahabi, 2017). Another study investigated that cofactors such as nicotinamide adenine dinucleotide (NADH) and reduced form nicotinamide adenine dinucleotide phosphate (NADPH)-dependent enzymes have a crucial role as reducing agents by the transfer of electrons from NADH by NADH-dependent enzymes (Kitching et al., 2014, Fig. 20.2). Observations on AuNPs biosynthesis in the presence of fungi *Fusarium oxysporum* exhibited that NADH-dependent reductase is considerably important in the biomineralization process (Mukherjee et al., 2002). Moreover, Au (III) ions are reduced by NADH or NADPH oxidoreductase either on the cell surface or in the cytoplasm. Narayanan et al. reported the participation of thermostable NADPH-dependent enzymes in the AuNPs biomineralization process (Narayanan & Sakthivel, 2011).

Another recent study on fungal AuNPs biosynthesis investigated the fungal pathogen *candida albicans* as a potential candidate for producing phytochelatins, which are composed of chain links of glucosecysteine and glycine ((c-Glu-cys)

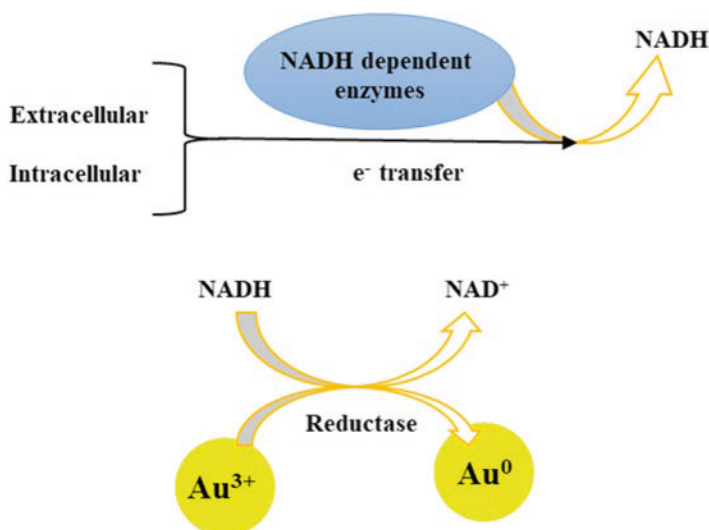


Fig. 20.2 Synthesis mechanism of AuNPs by extracellular or intracellular enzymatic reduction

n-Gly), by the transpeptidation reaction of a c-Glu-Cys dipeptide from a succession of glutathione molecules. Besides, glutathione-capped AuNPs have been formed by the reduction of Au (III) ions in the presence of glutathione (Owais et al., 2011).

Although melanin-mutant and melanin-complemented strains have not been studied, the association of biosynthesis redox mediators in the mycosynthesis of AuNPs has been reported in *Yarrowia lipolytica*. Interestingly, reduction of Au (III) ions to AuNPs occurred in the presence of melanin which was discharged by this fungus (Apte et al., 2013). Even though reduction by melanin has not been proved, further studies from the same research group revealed that melanin could induce the reduction of metal salts to their elemental form as nanomaterials. Remarkably, the phenol form of melanin oxidized to quinone, while Au (III) reduced to AuNPs (Apte et al., 2013).

Szczyziel et al. reported that the formation of AuNPs by *M. plumbeus* is a non-enzymatic process. However, the development of AuNPs occurred by the dead cells of *M. plumbeus* is an implicit indication of the function of polysaccharides in this phenomenon. It is well-known that high temperature results in exposure of polysaccharides, and these molecules become more susceptible to the gold ions for conjugation (Maliszewska et al., 2017) (Fig. 20.3).

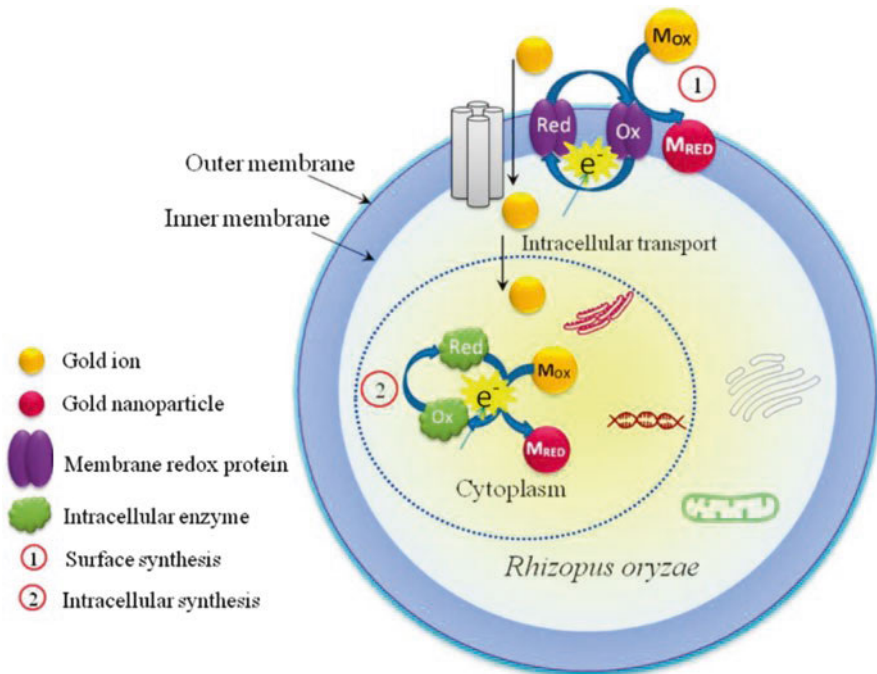


Fig. 20.3 Gold biomineralization mechanism in *R. oryzae*. (Das et al., 2012a, b)

20.3 Challenges and Scale-Up for Entrepreneurship

Mycosynthesis of gold nanoparticles has shown tremendous developments in synthesis and their applications over the last decade. However, there is a large gap between utilizing small-scale synthetic procedures and their commercial utilization. Much work is needed to improve the efficiency of synthesis, decreased toxicity, controlled size, and morphology. For example, mycosynthesis is a relatively slow process compared to other physical and chemical methods. A few challenges in mycosynthesis are addressed below:

1. Pathogenesis from fungi and fungal extracts is a challenge while scaling up for commercialization. Specifically, some extracts might contain pathogenic spores, which transmit to cause hazards to health and the ecosystem (Pantidos & Horsfall, 2014).
2. Particle size and monodispersity are important characteristics that must be controlled in the evaluation of gold nanoparticles for medicine and health care applications. The ability to control the shape, size, and size distribution of gold nanoparticles is more challenging with mycosynthesis since we are still lacking mechanistic insights (Akbari et al., 2011).
3. The stability of mycosynthesized gold nanoparticles is also a concern since the nanoparticles formed by fungi may be decomposed after a certain period.
4. Enzymes involved in the reduction process are not specifically categorized or addressed since the genes involved are hard to identify in fungi as a eukaryote. Thus, the mechanism of the reaction and enzyme roles in this process is hardly understood and is a major challenge in further recognition of mycosynthesis (Thakkar et al., 2010).

Besides the above-mentioned major challenges, gold nanoparticles synthesized using mycological route must be rigorously purified since the major applications are in biomedical fields. Downstreaming of fungal extracts is hard since it is time-consuming and requires several techniques. Even though, mycosynthetic route of gold nanoparticle generation is safer and economical. This is reported to be a non-toxic and eco-friendly “green-route” strategy. Exploiting this mycobased synthetic approach to gold nanoparticles has exciting aspects for addressing the current manufacturing challenges because this encompasses most of the 12 principles of green synthetic route (Patwardhan et al., 2018). Metabolic, zymology, and molecular mechanisms that help in synthesis should be studied in detail to overcome these challenges, increase synthetic rate, and obtain the required size and morphology of gold nanoparticles. Owing to the rich diversity of fungi, their potential as synthetic agents is still to be explored on large scale. Table 20.2 summarizes a few reported mycosynthetic gold nanoparticles with their applications.

Consequently, developments in this field of mycosynthesized gold nanoparticles are important in the industrial and pharmaceutical fields as antimicrobial, antifouling agents, catalysts, drug carriers, anticancerous agents, etc. Considering the immense need for fungi nanotechnology entrepreneurship in medical sciences, gold

Table 20.2 Synthesized gold nanoparticles using various fungus species and their applications

| | Fungus species | Applications | References |
|----|------------------------------------|----------------------------|--------------------------------|
| 1 | <i>Rhizopus oryzae</i> | Pesticides removal | Das et al. (2009) |
| 2 | <i>Fusarium semitectum</i> | Optoelectronics | Sawle et al. (2008) |
| 3 | <i>Helminthosporium solani</i> | Anticancer | Kumar et al. (2008) |
| 4 | <i>Penicillium brevicompactum</i> | Target cancer cells | Mishra et al. (2011) |
| 5 | <i>Volvariella volvacea</i> | Therapeutic | Philip (2009) |
| 6 | <i>Candida albicans</i> | Liver cancer detection | Owais et al. (2011) |
| 7 | <i>Aspergillus japonicus</i> AJP01 | Catalysis | Bhargava et al. (2015) |
| 8 | <i>Aspergillus Niger</i> | Kill mosquito larva | Prakash and Soni (2012) |
| 9 | <i>Mariannaea sp. HJ</i> | Nitrophenol reduction | Pei et al. (2017) |
| 10 | <i>Penicillium aculeatum</i> | Against hydatid cyst | Barabadi et al. (2017) |
| 11 | <i>Cylindrocladium floridanum</i> | Nitrophenol reduction | Narayanan and Sakthivel (2011) |
| 12 | <i>Pycnoporus sanguineus</i> | Catalysis | Shi et al. (2015) |
| 13 | <i>Trichoderma harzianum</i> | Hg ²⁺ detection | Tripathi et al. (2014) |
| 14 | <i>Aspergillus terreus</i> | Bactericidal against | Priyadarshini et al. (2014) |

nanoparticles have their major role in drug delivery, cancer treatment, gene editing, DNA analysis, antibacterial, biosensors, and MRI, which leads us to expand entrepreneurship of mycosynthetic approach in the synthesis of gold nanoparticles. In the energy sector, the role of economically synthesized gold nanomaterials promises to capture light, convert energy, which results in the fabrication of photovoltaics, etc. Promising research and developments will take time to scale up and bring costs down. Hopefully, the proper addressing of these challenges can lead to large-scale and commercial utilization of mycosynthesized gold nanoparticles, especially in medicine and health care system, in the coming years.

20.4 Conclusion

Fungi “nano factories” can produce large-scale materials for gram-scale industrial applications. However, the fabrication speed is still relatively slow compared to the chemical and physical approaches. This area of research is still in need to study the various factors that control the formation of the different morphologies and their effect on the potential toxicity in medical applications. Moreover, the research must expand to include easily reducible metal ions rather than the commonly used high oxidation states which are predicted to oxidize some of the components in the live cells.

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