

CONTEMPORARY STUDIES IN BIOLOGICAL SCIENCES I

Editors

Semra BENZER

Ali GÜL



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PREFACE

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CHAPTER 1

A NEW SEARCH OF BIOACTIVE COMPOUNDS: ALGAE

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INTRODUCTION

The study of metabolic products of plants and other groups of organisms is not only for a better understanding of nature, but also includes the discovery of possible useful metabolites in different areas of interest to humans. In recent years, the discovery of some antiviral, antimicrobial, anticoagulant and growth-inhibiting metabolites, especially from macroalgae, has led to the emergence of a new industrial field.

Since algae have been used in traditional medicine for a long time, they have been intensively researched by various researchers (Evrim et al., 2010). Chemicals responsible for antibiotic activity are widely found in macroalgae. Of particular interest among these are haloforms, halogenated compounds such as halogenated alkanes and alkenes, alcohols, aldehydes, hydroquinones and ketones. There is also a long list of terpenoids with antibiotic properties, many of which are halogenated. Sterols, heterocyclic and phenolic compounds also sometimes show antibiotic properties.

However, for most of them, it can only show antimicrobial activity at concentrations where it is toxic (Evrim et al., 2010).

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detected. In the methanol extract, in addition to P-hydroxybenzoic acid, caffeic acid, p-coumaric acid, protocatechuic acid ethyl ester and vanillic acid, salicylic acid, ferulic acid metabolites, specifically caftaric acid and campferol were found. P-hydroxybenzoic acid, caffeic acid, p-coumaric acid, protocatechuic acid ethyl ester, protocatechuic acid and vanillic acid were found in the acetone extract of *Chara hispida*. The phenolic metabolites found in the DMSO extract of this algae were determined as gallic acid, gentisic acid, catechin, rutine and cinnamic acid, as well as the phenolics in the acetone extract. It can also be emphasized that gallic acid and gentisic acid interact with other phenols in the *Chara hispida* DMSO/acetone extract mixture.

As we mentioned before, no other study was found regarding the antimicrobial activities of the algae extract mixtures included in the study and the antagonistic or synergistic interactions of these extracts. Therefore, the findings in our study can be claimed as the first records on the subject.

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CHAPTER 2

THE MEDICINAL-AROMATIC PROPERTIES of GINGER (*Zingiber officinale*) PLANT and TISSUE CULTURE STUDIES

**Çiğdem Alev ÖZEL¹
Yiğit Buğra AKYÖN²**

INTRODUCTION

Throughout history, human beings have used plants for nutrition, shelter, warmth, sheltering their wounds and treating their diseases (Acibuca and Budak, 2018). The first records of plant-based treatment date back to around 5000 BCE in the Mesopotamian civilization, where the use of 250 herbal drugs for therapeutic purposes has been identified (Demirezer, 2010). The aromatic use of plants and the extraction of essential oils can be traced back to ancient times, with a notable example found in the Ebers Papyrus, dating back to the 16th century BCE, discovered by the German Egyptologist Georg Ebers in 1872. The concept of aromatic medicine is evident in the papyrus, where it is discussed as a concept, and various treatment methods are described (Carpenter et. al., 2006). Over time, with advancements in technology, the addition of synthetic substances to food and beverages, as well as the introduction of synthetic drugs into production, has led to a decline in the traditional use of medicinal and aromatic plants. In the twentieth century, the discovery of side effects associated with synthetic drugs and the awareness of the harm caused by synthetic substances in food and beverages to human health have come to light. The increasing demand for traditional medicinal and aromatic plants has led to a return to natural products, with people shifting their focus towards natural and herbal products, primarily for health reasons. As a result, there has been a renewed interest in traditional medical practices, and herbal treatments have started to gain greater acceptance in the field of alternative

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CHAPTER 3

THE MEDICINAL-AROMATIC PROPERTIES of TURMERIC (*Curcuma longa*) PLANT and TISSUE CULTURE STUDIES

**Yiğit Buğra AKYÖN¹
Çiğdem Alev ÖZEL²**

INTRODUCTION

Throughout history, it is unimaginable to think of a time when humans and plants were separate entities. Starting with the history of humanity, plants have been utilized for various purposes such as food, clothing, fuel, and shelter. In the subsequent periods, humans have discovered additional properties of plants, incorporating the therapeutic benefits of specific plants into their lives (Jamshidi-Kia, Lorigooini, and Amini-Khoei, 2017). A Sumerian clay tablet dating back approximately 5000 years is considered one of the oldest written evidences indicating the use of medicinal plants for preparing remedies. Discovered in Nagpur, this tablet references over 250 different plants, including poppy, henbane, and mandrake, all containing alkaloids. It provides detailed instructions for preparing medicines from these plants, marking a significant historical record of early pharmacological knowledge (Castiglioni, 2019). In the ancient Chinese medical text “Pen T’Sao,” believed to have been written around 2500 B.C., information is provided about the use of plants, including drugs that are still in use today, along with their respective applications (Wiart, 2007). In the wisdom of ancient Indian culture, as documented in the Vedic teachings and sacred texts, there is a detailed account of plants used in Ayurvedic medicine. The extracts of these plants are intricately described, often associated with deities in their belief system. Many plants, such as turmeric, ginger, and basil, are revered for their sacred qualities (Tarun et al., 2016). Many ancient civilizations, much like those mentioned earlier, placed sig-

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Turmeric is one of the most important spices used today and is one of the food additives used safely all over the world. The reason why turmeric is used as a food additive is its main component, curcuminoids. Curcuminoids show anti-inflammatory, antitumor and antioxidant effects (Pistelli et al., 2012). They measured the content of curcuminoids in plant organs obtained by *in vitro* propagation. Plant parts obtained *in vitro* were examined with curcumin, bis-demethoxy curcumin, demethoxy curcumin by LC-DAD-ESI-MS method. It was observed that a significant amount of curcumin was produced from microrhizomes grown in MS medium containing 0.1 mg/L Kn, 1 mg/L NAA, 6% sucrose, 5% agar and 0.1% activated charcoal, thus success was achieved in the production of curcumin with tissue cultures. These results showed that it is safe to produce *in vitro* (Pistelli et al., 2012).

Tyagi et al. (2007) addressed the cost implications of sucrose as a carbon source and agar as a gelling agent in tissue culture studies conducted for plant production. In an effort to find a solution, the researchers utilized alternative sources, reducing the cost of turmeric plant regeneration and *in vitro* conservation by up to 73%. Isubgol, a colloidal and polysaccharide structure containing over 30% mucilage, was used as a low-cost substitute for agar as a gelling agent (Babbar and Jain, 1998; Ozel, Khawar and Arslan, 2008). In this cost-effective medium, no adverse effects on shoot regeneration and conservation were observed compared to the agar-gelled control medium. *C. longa* cultures, when conserved in the isabgol-gelled medium, exhibited 33–56% survival for up to 12 months, while the survival rate in the control group was 16%. After one year, the genetic stability of the *in vitro*-preserved plants was evaluated using the RAPD technique, revealing no significant variation in the RAPD profiles of the *in vitro*-preserved seedlings.

In conclusion, the tissue culture production of turmeric and related studies play a crucial role in the sustainability of agriculture, the development of disease-resistant plants, and the enhancement of agricultural production.

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CHAPTER 4

ONE HEALTH: LOSS OF BIODIVERSITY

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INTRODUCTION

Zoonotic infectious diseases, which emerged in the late twentieth century and mostly originate from wildlife animals, reveal that animal, human and ecosystem health are interrelated. A multidisciplinary approach involving animal, human and ecosystem health has been developed to conduct risk assessments, take preventive measures and develop control plans. Following the emergence of various viral diseases such as bird flu and Zika virus, the interrelatedness of animal, human and ecosystem health has become even more important. The “One Health” concept was developed in 2004, based on the “One Medicine” concept of human and veterinary medicine (Destoumieux-Garzón et al., 2018; Mackenzie and Jeggo, 2019).

According to United States Centers for Disease Control and Prevention, One Health means “a collaborative, multisectoral, and transdisciplinary approach—working at the local, regional, national, and global levels—with the goal of achieving optimal health outcomes recognizing the interconnection between people,

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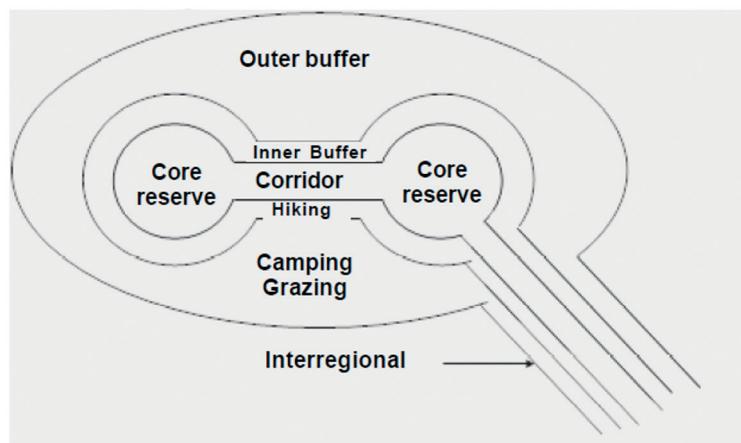


Figure 5. Protected area template (Nigam, 2019)

CONCLUSION

This review examined biodiversity loss, one of the ecosystem health components within the concept of One Health. The definition of biodiversity, the relationships between species, the causes of biodiversity loss, especially its effects on ecosystem health, and the measures taken to prevent biodiversity loss are mentioned. Today, when biodiversity is rapidly decreasing day by day, our understanding and conservation efforts in this context need to be further strengthened. Future studies should take a comprehensive approach to more deeply understand the relationships between biodiversity sustainability, ecosystem services, and human health.

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CHAPTER 5

EFFECTS OF ANTIFOULING SUBSTANCES ON AQUATIC ORGANISMS

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INTRODUCTION

Organisms such as mussels, algae and bacteria that live in aquatic ecosystems cause damage by adhering to vehicles such as boats, ships, oil wells, or piers (Figure 1). The adhesion of these organisms to an area is a fast and complex process (Li et al., 2023). In the first stage, organic substances such as proteins, polysaccharides, and inorganic substances adhere to a solid surface immersed in water, forming a thin film layer. This substrate film layer is a prerequisite for the adsorption of microorganisms (Jain and Bhosl, 2009). In the second stage, bacteria or diatoms adhere to the substrate film layer and form the biofilm layer. In the third stage, microfouling, as a result of the excessive proliferation of microorganisms in the biofilm layer, the biofilm is fixed and prepared for adhesion to different organisms. In the fourth and final stage, macrofouling, different species adhere to the membrane surface. Therefore, a biological pollution occurs (Li et al., 2023) (Figure 2).

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CONCLUSION

The effects of antifouling paints, which have an important place especially in the marine industry, on aquatic organisms are shown in this review. The fact that organotin compounds are extremely toxic and show both acute and chronic toxic effects on aquatic organisms has led to these substances being banned by international agreements. However, studies have revealed the acute toxic effects of alternative antifouling substances, the use of which has started to increase, and harmful effects on non-target species have also been observed in acute and chronic exposure studies. For this reason, some alternative antifouling agents are also banned. In the long term, the effects of new or existing alternative antifouling substances on non-target aquatic species are of concern.

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CHAPTER 6

Anti-Quorum Sensing Activity and Anti-Biofilm Potential of Extracts Isolated from Fish Associated Bacteria

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INTRODUCTION

Antibiotic-resistant bacterial strains have emerged due to the misuse of antibiotics, which are used to treat and prevent diseases. Misuse of antibiotics results in selection pressure, leading to drug resistance (Chu et. al., 2014).

As multidrug-resistant infections increase, antibacterial therapies become less effective (Rama Devi et al., 2016). The scientific community is concerned about developing novel antimicrobial agents and studying their mechanisms of action as potential precursors to therapeutic compounds to address this widespread issue. Bacterial quorum sensing (QS) is an intercellular signals and gene control process that relies on density of population for cell-to-cell communication.

The cell density-dependent bacterial communication system is used to regulate the virulence phenotype that contributes to the pathogenicity of many pathogenic microorganisms (Rajkumari et al., 2019).

In collaborative activities such as colony formation and motility, many bacteria release metabolic intermediates that are considered signals, which have low molecular weights.

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CHAPTER 7

BENEFITS AND DAMAGES OF LACTIC ACID BACTERIA BIOFILM IN THE FOOD INDUSTRY

Tugba KILIC¹

INTRODUCTION

Lactic acid bacteria (LAB) are bacteria that are within the phylum of Firmicutes, Gram-positive, rod or cocci-shaped, facultatively anaerobic, low G+C%, catalase-negative, non-motile (except for a few privileged members), lack cytochrome, do not form spores (except *Sporolactobacillus inulinus*), and generate lactic acid as the end-product during carbohydrate fermentation. The main LAB genera are *Lactobacillus*, *Lactococcus*, *Leuconostoc*, *Pediococcus*, *Oenococcus*, *Enterococcus*, and *Streptococcus* (Gómez et al., 2016; Arena et al., 2017). LAB is isolated from different sources such as milk, meat, wine, soil, silages, bread, sauerkraut, and sausages as food-related habitats (Arena et al., 2017; Mgomi et al., 2023).

LABs are usually recognized as safe (GRAS) and, therefore, recognized as food-grade bacteria. Thus, from a legal perspective, they can easily be used in foods as probiotics (Abid et al., 2018). Furthermore, LABs serve as starter cultures in the fermentation of various foods and often help modify and preserve flavor and texture. *Lactobacillus*, *Pediococcus*, *Oenococcus*, *Lactococcus*, *Leuconostoc*, and *Streptococcus* are important genera for food biotechnology (Rothstein et al., 2020). Most LAB transform lactose and other sugars into lactic acid (Aoudia et al., 2016). Lactic acid limits the growth of food spoilage microorganisms, lowers the pH of the food substrate, and provides foods with a characteristic sour taste

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CHAPTER 8

SOME MERISTIC AND MORPHOMETRIC CHARACTERISTICS OF *Arabibarbus grypus* (Heckel, 1843) LIVING IN KARAKAYA DAM LAKE

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INTRODUCTION

The body of *Arabibarbus grypus* is from the Cyprinidae family, is flattened on the sides and covered with large scales. The body is seen as round from the front of the dorsal fin. Next to the large fleshy lips, the mouth is located ventrally. The dorsal fin is further forward than the ventral fin. The abdominal areas are dirty yellow, the sides tend to dark brown, and the back is dark brown. Sometimes there are greenish reflections on the sides of the fish's body. The fins except the caudal and anal fins are light colored and these are dark colored. Its distribution areas are the Euphrates and Tigris river systems (Geldiay & Balık, 2007; Düşükcan, Çalta & Eroğlu, 2015; Düşükcan & Çalta, 2018; Yüksel, Demirok, Gündüz & Çoban, 2020).

Within the scope of biological characteristics of fish, determining their morphometric and meristic properties in different water systems is important for ichthyofauna. In order to reveal the ichthyofauna in a wetland, the biological characteristics of all fish species must be known and periodic monitoring studies must be carried out. Determining the morphometric and meristic properties of fish in different water systems within the scope of their biological characteristics will be important for ichthyofauna. Studies on fish populations should be evaluated from

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CHAPTER 9

THE USE OF ARTIFICIAL INTELLIGENCE TECHNOLOGIES IN THE BIOLOGICAL SCIENCES

**Recep BENZER¹
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INTRODUCTION

Artificial Intelligence (AI) is one of the fundamental research areas in computer science. With its rapid technological advancements and broad application scope, artificial intelligence has become widespread very quickly, especially due to its strong applicability in solving problems that are not easily addressed by traditional computing structures, particularly those challenging for humans (Rich and Knight, 2009).

The progress of science and technology has brought about developments in various fields, from unraveling the genetic structures of living organisms to the creation of satellites for geographic information systems. The new knowledge emerging in every aspect of our lives, ranging from natural sciences to social sciences, is processed through the use of mathematical models and computer software, facilitating the refinement of large amounts of data (Eaton et al., 2020). Therefore, in the 21st century where the organization of any acquired information is increasingly crucial, it is evident that mathematics and computerized processes will play a prominent role in understanding scientific data. Particularly, the field of biomathematics (mathematical biology), a subset of applied mathematics, is gaining more attention as it quantitatively evaluates data in biological sciences, which are integral to daily life. Interdisciplinary studies encompassing biology, mathematics, and computer science are becoming more prevalent. To the extent that the applications of biomathematics are becoming inevitable in the career plans of young individuals (Akman et al., 2020).

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tutions towards investing in research and development within this field. Several major tech companies have made strides and introduced new approaches in deep learning, fostering the evolution of new applications and perspectives. Entities such as Google, Microsoft, Baidu, IBM, Apple, Nvidia, Facebook, Twitter, Amazon, among many others, have actively engaged in research and development pertaining to deep learning.

When examining the studies conducted, it becomes evident that deep learning architectures have introduced a new perspective and breakthrough in artificial intelligence technologies. The research on future predictions using deep learning architectures is believed to propel current technologies several years forward. The active integration of deep learning architectures into our lives is noticeably geared towards simplifying human life and facilitating the pursuit of a healthy lifestyle. With a particular focus on biology, the utilization of deep learning architectures is seen as a pivotal driver in advancing predictive capabilities and discoveries. Especially within the realm of biology, the application of deep learning architectures is increasingly playing a more effective role in diagnostics, treatment procedures, and disease prognostication. This is perceived as a significant stride towards enhancing human life quality concerning health-related aspects.

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