

Genetics of Morphogenesis in Basidiomycetes

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In this chapter, our aim is to discuss the current knowledge of the genetics of morphogenesis in basidiomycetes. We begin by outlining some features that are shared among fungi in general and those that are unique to basidiomycetes. With this background of basic fungal morphology and physiology, we focus our discussion on the genetics of morphogenesis and the fascinating biology of three broad groups of basidiomycetes: the smuts, the rusts and the mushrooms. Where the smuts and rusts are considered, there is ample discussion on pathogenesis, as these two broad groupings of fungi are particularly destructive to plants worldwide. Of course, mushrooms represent a valuable commodity in their own right. Each of these broad sections encapsulates a current assessment of the environmental cues and the genes regulating morphological change during development. Each of these three taxon-focused sections also concludes with a section discussing current trends and/or future directions.

1. INTRODUCTION

The kingdom fungi lies on the opisthokont lineage of the Eukaryotes and thus is a closer relative to the animals than to the plants. The fossil record of the fungi is estimated to date back as far as 900 million years with all major classes represented about 300 million years ago (Taylor et al. 1994). The basidiomycetes are one of the two ultimate branches of the true fungi, the other being the ascomycetes. About 30,000 basidiomycete species have been described. Among the basidiomycetes are the well known mushrooms of the Hymenomycetes, the rusts of the Urediniomycetes known through recorded history as devastating pathogens of staple grain crops, and the smuts of the Ustilaginomycetes that cause persistent crop losses. In this chapter, we present an overview of some of the central processes particular to the basidiomycetes. We then

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Table of Contents Genetics And Morphogenesis In The Basidiomycetes

1. Understanding the eBook Genetics And Morphogenesis In The Basidiomycetes
 - The Rise of Digital Reading Genetics And Morphogenesis In The Basidiomycetes
 - Advantages of eBooks Over Traditional Books
2. Identifying Genetics And Morphogenesis In The Basidiomycetes
 - Exploring Different Genres
 - Considering Fiction vs. Non-Fiction
 - Determining Your Reading Goals
3. Choosing the Right eBook Platform
 - Popular eBook Platforms
 - Features to Look for in an Genetics And Morphogenesis In The Basidiomycetes
 - User-Friendly Interface
4. Exploring eBook Recommendations from Genetics And Morphogenesis In The Basidiomycetes

- Personalized Recommendations
 - Genetics And Morphogenesis In The Basidiomycetes User Reviews and Ratings
 - Genetics And Morphogenesis In The Basidiomycetes and Bestseller Lists
5. Accessing Genetics And Morphogenesis In The Basidiomycetes Free and Paid eBooks
 - Genetics And Morphogenesis In The Basidiomycetes Public Domain eBooks
 - Genetics And Morphogenesis In The Basidiomycetes eBook Subscription Services
 - Genetics And Morphogenesis In The Basidiomycetes Budget-Friendly Options
 6. Navigating Genetics And Morphogenesis In The Basidiomycetes eBook Formats
 - ePub, PDF, MOBI, and More
 - Genetics And Morphogenesis In The Basidiomycetes Compatibility with Devices
 - Genetics And Morphogenesis In The Basidiomycetes Enhanced eBook Features
 7. Enhancing Your Reading Experience
 - Adjustable Fonts and Text Sizes of Genetics And Morphogenesis In The Basidiomycetes
 - Highlighting and Note-Taking Genetics And Morphogenesis In The Basidiomycetes
 - Interactive Elements Genetics And Morphogenesis In The Basidiomycetes
 8. Staying Engaged with Genetics And Morphogenesis In The Basidiomycetes
 - Joining Online Reading Communities
 - Participating in Virtual Book Clubs
 - Following Authors and Publishers Genetics And Morphogenesis In The Basidiomycetes
 9. Balancing eBooks and Physical Books Genetics And Morphogenesis In The Basidiomycetes
 - Benefits of a Digital Library
 - Creating a Diverse Reading Collection Genetics And Morphogenesis In The Basidiomycetes
 10. Overcoming Reading Challenges
 - Dealing with Digital Eye Strain
 - Minimizing Distractions
 - Managing Screen Time
 11. Cultivating a Reading Routine Genetics And Morphogenesis In The Basidiomycetes
 - Setting Reading Goals Genetics And Morphogenesis In The Basidiomycetes
 - Carving Out Dedicated Reading Time
 12. Sourcing Reliable Information of Genetics And Morphogenesis In The Basidiomycetes

- Fact-Checking eBook Content of Genetics And Morphogenesis In The Basidiomycetes
- Distinguishing Credible Sources
- 13. Promoting Lifelong Learning
 - Utilizing eBooks for Skill Development
 - Exploring Educational eBooks
- 14. Embracing eBook Trends
 - Integration of Multimedia Elements
 - Interactive and Gamified eBooks

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