Gene Manipulations in **Fungi**

Edited by

J. W. Bennett

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Preface

The fungi constitute a morphologically and physiologically diverse group of eukaryotes, now usually classified in a separate kingdom. Among the attributes that distinguish the fungi are a hyphal or yeastlike form; a rigid, frequently chitinous cell wall; apical growth; absorptive, heterotrophic metabolism; reproduction by spores; and small genome size.

The economic importance of the fungi is a reflection of their diverse, and often unique, metabolic capabilities. In nature, they are major agents of disease and decay. The ability to secrete enzymes into the environment makes filamentous fungi attractive to exploiters of modern biotechnology. In industry, they are the source of many important natural products such as citric acid and penicillin. In the laboratory, they are model systems for basic research.

Scientists who are otherwise separated by discipline, organism, geography, and objectives nevertheless share common goals in applying modern genetic techniques to fungi. With the important exception of Saccharomyces cerevisae, the molecular biology of fungi is just beginning. Chapter after chapter in this volume emphasizes the need for improved transformation systems, appropriate vectors, and broadly applicable selectable markers.

"Gene Manipulations in Fungi" combines a review of classical fungal genetics, contemporary research, and responsible speculation about the future. Yeast is, without question, the primary model system for eukaryotes. It is rapidly replacing *Escherichia coli* as the organism of choice for advanced studies in molecular genetics. Elegant research is also being conducted with the molds *Aspergillus nidulans* and *Neurospora crassa*.

We have included a number of topics of economic importance. Although the ability to manipulate genomes from plant and animal pathogens, or from most industrial fungi, is currently very limited, intense research is under way. It is useful for applied scientists to understand the state of the art. All of these economically important fungi offer special challenges to the virtuosity of the molecular biologist. Novel adaptations of recombinant DNA techniques are required, and we hope this book will stimulate innovative approaches. Where

, xviii Preface

appropriate, useful methodologies are described in the text; elsewhere, literature citations are provided.

Molecular biology generates optimism, even euphoria. Many of our authors project their enthusiasm. We share their optimism and are pleased to have brought together such a diverse group of mycophiles. We predict that "Gene Manipulations in Fungi" will be only the first of many books celebrating the marriage between molecular biology and mycology.

J. W. Bennett Linda L. Lasure

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Contents

	Contributors xiii
	Preface xvii
	Acknowledgments xix
	I Historical Perspective: Mutants to Models
1	From Auxotropic Mutants to DNA Sequences
	John R. S. Fincham
	I. The Establishment of the Paradigm: One Gene-One Polypeptide
2	Molecular Taxonomy of the Fungi
	C. P. Kurtzman
	I. Introduction

1		Cont	tents
	III. IV. V. VI. VII.	DNA Base Composition DNA Relatedness Mitochondrial DNA Relatedness Ribosomal RNA Relatedness Comparison of Relatedness from Nucleic Acid Studies with That Determined by Other Methodologies References	38 41 49 51 53 56
3	Fung	al Mitochondrial Genomes	
	Lawre	nce T. Grossman and Michael E. S. Hudspeth	
	I. III. IV. V. VI. VIII. VIII. IX.	Introduction Physical Organization Genes and the Genetic Code Gene Order Transcription Pseudogenes Mitochondrial Plasmids Effects of Alteration of mtDNA Nuclear-Cytoplasmic Interactions References	66 66 70 74 76 81 83 84 88 91
4	Mode	eling the Environment for Gene Expression	
	Barbar	ra E. Wright	
	I. II. III.	Introduction Modeling Metabolism under Steady-State Conditions Modeling Metabolism during Product Accumulation	106 115
		II Yeasts	
5		naromyces cerevisiae as a Paradigm for Modern cular Genetics of Fungi	
	Jasper	Rine and Marian Carlson	
	I. III. IV. V.	Cloning a Gene	

Conte	nts		vii
	VI. VII.	Other Uses of Cloned Genes An Agenda for Progress References	
6	Yeast	Transformation	
	George	Bogusławski	
	I. II. III. IV.	Introduction Specific Transformation Systems in Yeast Applications Conclusions References	167 178 187
7	Repla	of the LYS2 Gene for Gene Disruption, Gene decement, and Promoter Analysis in paromyces cerevisiae	
	Debra A	A. Barnes and Jeremy Thorner	
	III.	Introduction Characterization of the LYS2 Gene and Its Product Genetic Manipulations Utilizing the LYS2 Gene Conclusions and Prospectus References	203 210 222
		III Molds	
8	Molec	cular Biology of the qa Gene Cluster of ospora	
	Layne F	Huiet and Mary Case	
	II.	Introduction Molecular Analysis of the qa Cluster Transformation of the qa Gene Cluster in Neurospora Summary References	230 236 242
9	Neuro	spora Plasmids	
	John A.	Kinsey	
		Introduction	

viii			Contents

	III. IV.	Replicating Plasmids Constructed in the Laboratory	253
10	Clonin	ng and Transformation in Aspergillus	
	G. Turr	ner and D. J. Ballance	
	I. III. IV. V. VI.	Introduction Transformation Methodology Fate of Transforming DNA after Entry The Quest for Replicating Vectors Development of Vectors Future Prospects References	260 265 270 273 275
11	Expre	ssion of Aspergillus Genes in Neurospora	
	Richard	L. Weiss, Diane Puetz, and Jan Cybis	
	I. III. IV. V. VI.	Transformation of Neurospora crassa Isolation of the Aspergillus nidulans Gene for Ornithine Carbamoyltransferase Transformation of the Neurospora crassa arg-12 Mutant Control of Gene Expression Subcellular Localization of the Active Enzyme Conclusions References	281 282 288 289 290
12	Gene Fungi	Dosage Effects and Antibiotic Synthesis in	
	С. Н. С	O'Donnell, A. Upshall, and K. D. Macdonald	
	I. II. III.	Introduction Methods for the Amplification of Genetic Material in Microorganisms Studies with Disomic Strains of Aspergillus nidulans References	295 297
13		al Genetics and Molecular Biology of the ol of Gene Expression in Aspergillus nidulans	
	Herbert	N. Arst, Jr., and Claudio Scazzocchio	
	I.	Formal Genetic Methodology of Aspergillus nidulans as Applied to the Study of Control Systems	310

Contents		ix

	III. IV. V. VI.	The Metabolic Versatility of A. nidulans and Its Exploitation. Regulatory Genes Putative Receptor Sites The Spatial Organization of Functionally Related Genes At What Level Does Regulation of Gene Expression Occur? References	313 327 330 335
14	A Clo	ning Strategy in Filamentous Fungi	
	P. M. C	Green and Claudio Scazzocchio	
		Text	
		IV Applications	
		The wife of the wife	
15	Prima	ry Metabolism and Industrial Fermentations	
	Ramuna	as Bigelis	
	I. III. IV. V. VI. VII. VIII. IX. X. XI.	Primary Metabolites Genetic Approaches to the Production of Primary Metabolites Organic Acids Amino Acids Polysaccharides Lipids Nucleotides and Nucleic Acid—Related Compounds Vitamins Polyols Ethanol The Promise of Biotechnology References	359 360 368 372 374 376 378 380 382 383
16	Mitoc	hondrial DNA for Gene Cloning in Eukaryotes	
		dzynski and Karl Esser	
	11.	Introduction Fungal Plasmids Cloning Vectors of Mitochondrial Origin Biotechnological Implications References	404 409 412

17	Mole	ecular Bases of Fungal Pathogenicity to Plants	
	O. C.	Yoder and B. Gillian Turgeon	
	I. II. IV. V. VI.	Infection Structures Cutinase	418 421 422 424 424
18	Morp	phogenesis and Dimorphism of Mucor	
	Ronald	d L. Cihlar	
	I. II. IV.	Introduction Dimorphism of Mucor Molecular Analysis of Mucor Perspectives—Molecular Genetics References	450 457 462
19		ard Gene Manipulations with Selected Human al Pathogens	
	W. LaJ	Jean Chaffin	
	II. IV. V. VI.	Introduction Cryptococcus neoformans Histoplasma capsulatum Wangiella dermatitidis Candida albicans Concluding Remarks References A	471 473 474 476 486
20	Funga	al Carbohydrases: Amylases and Cellulases	
	Bland S	S. Montenecourt and Douglas E. Eveleigh	
	II.	Introduction	194 501

V Postscript

21	Prospects for a Molecular Mycology	
	J. W. Bennett	
	I. Introduction	5
	Appendixes	
I	Fungal Taxonomy	
	Linda L. Lasure and J. W. Bennett	
	I. Introduction	ļ
II	Conventions for Gene Symbols	
	J. W. Bennett and Linda L. Lasure	
	I. Aspergillus nidulans 538 II. Neurospora crassa 539 III. Saccharomyces cerevisiae 540 IV. Other Fungi 542 References 542)
	Index	j

I Historical Perspective: Mutants to Models

Mandrins in Wholash