

Exploitation of Fungi

The fungi are a highly diverse kingdom of eukaryotic microbes. Recent advances in molecular genetics, together with the release of whole genome sequences of an increasing number of fungi, are facilitating their exploitation and commercialization. Fungi have the ability to secrete large quantities of proteins of commercial value, and their complex secondary metabolic pathways produce a diverse range of bioactive compounds that have had a major impact in the pharmaceuticals market. In addition, the fungi themselves are increasingly being developed as alternatives to conventional chemically based pest control strategies, and as bioremediation agents capable of transforming pollutants in the soil environment. With chapters written by international experts, this volume highlights current and future biological, biochemical and molecular exploitation of the fungi in biotechnology. It will have broad appeal, not only to mycologists and microbiologists, but also to biomedical scientists, biotechnologists, environmental and molecular scientists, plant pathologists and geneticists.

GEOFF ROBSON is a Senior Lecturer in the Faculty of Life Sciences at the University of Manchester.

PIETER VAN WEST is a Senior Lecturer and a Royal Society University Research Fellow in the Institute of Medical Sciences at the University of Aberdeen.

GEOFF GADD is Professor of Microbiology, Head of the Division of Environmental and Applied Biology, and Deputy Research Director in the School of Life Sciences at the University of Dundee.



Exploitation of Fungi

Symposium of the British Mycological Society held at the University of Manchester September 2005

EDITED BY
G. D. ROBSON, P. VAN WEST
AND G. M. GADD

Published for the British Mycological Society





CAMBRIDGE UNIVERSITY PRESS
Cambridge, New York, Melbourne, Madrid, Cape Town, Singapore, São Paulo
Cambridge University Press
The Edinburgh Building, Cambridge CB2 2RU, UK

Published in the United States of America by Cambridge University Press, New York

www.cambridge.org

Information on this title: www.cambridge.org/9780521859356

© Cambridge University Press 2007

This publication is in copyright. Subject to statutory exception and to the provisions of relevant collective licensing agreements, no reproduction of any part may take place without the written permission of Cambridge University Press.

First published 2007

Printed in the United Kingdom at the University Press, Cambridge

A catalogue record for this publication is available from the British Library

ISBN 978-0-521-85935-6 hardback

Cambridge University Press has no responsibility for the persistence or accuracy of URLs for external or third-party internet websites referred to in this publication, and does not guarantee that any content on such websites is, or will remain, accurate or appropriate.



Contents

| Lis | t of contributors | page vii |
|-----|--|----------|
| Pre | eface | XV |
| I | Comparative and functional fungal genomics | 1 |
| 1 | Genome and proteome analysis of industrial fungi | |
| | S. E. Baker, C. F. Wend, D. Martinez, J. K. Magnuson, | |
| | E. A. Panisko, Z. Dai, K. S. Bruno, K. K. Anderson, | |
| | M. E. Monroe, D. S. Daly and L. L. Lasure | 3 |
| 2 | The rice blast story: from genome sequence to function | |
| | R. A. Dean, T. Mitchell, R. Kulkarni, N. Donofrio, | |
| | A. Powell, Y. Y. Oh, S. Diener, H. Pan, D. Brown, | |
| | J. Deng, I. Carbone, D. J. Ebbole, M. Thon, | |
| | M. L. Farman, M. J. Orbach, C. Soderlund, J-R. Xu, | |
| | Y-H. Lee, N. J. Talbot, S. Coughlan, J. E. Galagan and | 1 |
| | B. W. Birren | 10 |
| II | Bioactive molecules | 23 |
| 3 | The biosynthesis of polyketides, acyl tetramic acids | |
| | and pyridones by filamentous fungi | |
| | R. J. Cox | 25 |
| 4 | Fungal metabolites as lead structures for agriculture | |
| | T. Anke and E. Thines | 45 |
| 5 | Molecular and genetic analysis of symbiosis expressed | |
| | secondary metabolite genes from the mutualistic fungal | |
| | endophytes Neotyphodium lolii and Epichloë festucae | |
| | B. Scott, C. A. Young, A. Tanaka and E. J. Parker | 59 |
| 6 | What can genomics tell us about secondary | |
| | metabolism in Aspergillus? | |
| | G. Turner | 78 |



vi Contents

| III 7 | Protein folding and secretion The role of microtubules and motors for polarized | 93 |
|--------------|--|-----|
| | growth of filamentous fungi | 0.5 |
| 0 | R. Fischer and D. Veith | 95 |
| 8 | The cellular response to protein unfolding stress <i>M. Schröder</i> | 117 |
| 9 | Protein secretion and associated stresses in | 11/ |
| 7 | Aspergillus: a genomic perspective | |
| | D. B. Archer, S. E. Barnes and T. Guillemette | 140 |
| IV | Fungal bioremediation | 159 |
| 10 | Metal stress and the single yeast cell: Berkeley | 137 |
| | Award Lecture | |
| | S. V. Avery | 161 |
| 11 | Bioluminescence-based fungal biosensors | |
| | H. J. Weitz | 187 |
| 12 | White-rot fungi and xenobiotics | |
| | P. J. Harvey and C. E. Scheer | 205 |
| 13 | Metal and mineral transformations: a | |
| | mycoremediation perspective | |
| | M. Fomina and G. M. Gadd | 236 |
| \mathbf{V} | Fungal biocontrol of pests | 255 |
| 14 | Fungal control of subterranean pests | |
| | H. Strasser, S. H. M. Hutwimmer and R. Zelger | 257 |
| 15 | Development of mycoherbicides and evaluation of | |
| | potential risks (using Stagonospora as a model) | |
| | M. Maurhofer, D. Boss, M. O. Ahonsi and G. Défago | 275 |
| 16 | A novel understanding of the three-way interaction | |
| | between Trichoderma spp., the colonized plant and | |
| | fungal pathogens | |
| | S. L. Woo, M. Ruocco, R. Ciliento, P. Ambrosino, | |
| | R. Marra, F. Vinale, S. Lanzuise, D. Turrà, S. Ferraioli, | |
| 1.5 | I. Soriente, S. Gigante, F. Scala and M. Lorito | 291 |
| 17 | Fungal parasites of invertebrates: multimodal | |
| | biocontrol agents? | 210 |
| | L. V. Lopez-Llorca and HB. Jansson | 310 |
| Index | | 336 |



Contributors

M.O. Ahonsi

River Basin Research Center, Gifu University, Yanagido 1–1, Gifu 501-1193, Japan

P. Ambrosino

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale, Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA), Italy

K. K. Anderson

Decision & Sensor Analytics, Pacific Northwest National Laboratory, 906 Battelle Blvd., Richland, WA 99352, USA

T. Anke

Institut für Biotechnologie und Wirkstoff-Forschung, IBWF, Kaiserslautern 67663, Germany

D. B. Archer

School of Biology, University of Nottingham, University Park, Nottingham NG7 2RD, UK

S. V. Avery

School of Biology, University of Nottingham, University Park, Nottingham NG7 2RD, UK

S. E. Baker

Fungal Biotechnology Team MSIN: K2–12,
Chemical and Biological Processes
Development Group,
Pacific Northwest National
Laboratory,
902 Battelle Blvd.,
Richland,
WA 99352,
USA



viii List of contributors

S. E. Barnes

School of Biology,
University of Nottingham,
University Park,
Nottingham NG7 2RD,
UK

B. W. Birren

The Broad Institute,
Massachusetts Institute of
Technology,
77 Massachusetts Avenue,
Cambridge,
MA 02139–4307,
USA

D. Boss

Plant Pathology, Institute of Integrative Biology, ETH Zurich, LFW, 8092 Zurich, Switzerland

D. Brown

North Carolina State University, Department of Plant Pathology, Campus Box 7251, Raleigh, NC 27695–7251, USA

K. S. Bruno

WA 99352,

USA

Fungal Biotechnology Team MSIN: K2–12, Chemical and Biological Processes Development Group, Pacific Northwest National Laboratory, 902 Battelle Blvd., Richland,

I. Carbone

North Carolina State University, Department of Plant Pathology, Campus Box 7244, Raleigh, NC 27695–7244, USA

R. Ciliento

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale, Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA), Italy

S. Coughlan

Agilent Technologies Inc., Little Falls Site, 2850 Centerville Road, Wilmington, DE 19808, USA

R. J. Cox

School of Chemistry and Biological Sciences,
University of Bristol,
Bristol BS8 1TS
UK

Z. Dai

USA

K2–12, Chemical and Biological Processes Development Group, Pacific Northwest National Laboratory, 902 Battelle Blvd., Richland, WA 99352,

Fungal Biotechnology Team MSIN:



List of contributors

ix

D. S. Daly

Statistical Sciences,
Pacific Northwest National
Laboratory,
3180 George Washington Way,
Richland, WA 99352,
USA

R. A. Dean

Center for Integrated Fungal Research, Department of Plant Pathology, 1200 Partners Building II, Box 7251, North Carolina State University, Raleigh, NC 27695, USA

G. Défago

Plant Pathology, Institute of Integrative Biology, ETH Zurich, LFW, 8092 Zurich, Switzerland

J. Deng

North Carolina State University, Department of Plant Pathology, Campus Box 7251, Raleigh, NC 27695–7251, USA

S. Diener

North Carolina State University, Department of Plant Pathology, Campus Box 7253, Raleigh, NC 27695–7253, USA

N. Donofrio

North Carolina State University, Department of Plant Pathology, Campus Box 7251, Raleigh, NC 27695–7251, USA

D. J. Ebbole

Department of Plant Pathology and Microbiology, Peterson Building, Rm 120, MS# 2132 Texas A&M University, College Station, TX 77843–2132, USA

M. L. Farman

Department of Plant Pathology, University of Kentucky, 1405 Veterans Drive, Lexington, KY 40546–0312, USA

S. Ferraioli

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale, Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA), Italy

R. Fischer

Institute for Applied Life Sciences, Applied Microbiology, University of Karlsruhe, Hertzstr. 16, D-76187 Karlsruhe, Germany



x List of contributors

M. Fomina

Division of Environmental and Applied Biology, Biological Sciences Institute, School of Life Sciences, University of Dundee, Dundee DD1 4HN, Scotland. UK

G. M. Gadd

Division of Environmental and Applied Biology, Biological Sciences Institute, School of Life Sciences, University of Dundee, Dundee DD1 4HN, Scotland, UK

J. E. Galagan

The Broad Institute,
Massachusetts Institute of
Technology,
77 Massachusetts Avenue,
Cambridge,
MA 02139–4307,
USA

S. Gigante

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale, Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA),

T. Guillemette

Italy

Laboratoire de Microbiologie, UMR 77 Pathologie Végétale, Université d'Angers, 2 bd Lavoisier, 49045 Angers cedex, France

P. J. Harvey

University of Greenwich, Medway School of Science, Central Avenue, Chatham Maritime, Kent ME4 4TB, UK

S. H. M. Hutwimmer

Institute of Microbiology, University of Innsbruck, Technikerstrasse 25, A-6020 Innsbruck, Austria

H.-B. Jansson

Department of Marine Sciences and Applied Biology, University of Alicante, Apartado 99, E-03080 Alicante, Spain

R. Kulkarni

RTI, 3040 Cornwallis Road, Research Triangle Park, NC 27709, USA

S. Lanzuise

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale, Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA), Italy



List of contributors

хi

L. L. Lasure

Fungal Biotechnology Team MSIN:

K2-12.

Chemical and Biological Processes

Development Group,

Pacific Northwest National

Laboratory,

902 Battelle Blvd., Richland,

WA 99352,

USA

Y.-H. Lee

Seoul National University,

School of Agricultural

Biotechnology, Suwon, 441–744,

Korea

L. V. Lopez-Llorca

Department of Marine Sciences and

Applied Biology,

University of Alicante,

Apartado 99, E-03080 Alicante,

Spain

M. Lorito

Dipartimento di Arboricoltura,

Botanica e Patologia Vegetale,

Sezione Patologia Vegetale,

Universita degli Studi di Napoli

Federico II.

Via Universita,

100-80055 Portici (NA),

Italy

J. K. Magnuson

Fungal Biotechnology Team MSIN:

K2-12,

Chemical and Biological Processes

Development Group,

Pacific Northwest National

Laboratory,

902 Battelle Blvd.,

Richland,

WA 99352.

USA

R. Marra

Dipartimento di Arboricoltura,

Botanica e Patologia Vegetale,

Sezione Patologia Vegetale,

Universita degli Studi di Napoli

Federico II,

Via Universita,

100-80055 Portici (NA),

Italy

D. Martinez

Genome Annotation and Analysis,

Joint Genome Institute.

Los Alamos National Laboratory,

Los Alamos,

NM 87545.

USA

M. Maurhofer

Plant Pathology,

Institute of Integrative Biology,

ETH Zurich,

LFW,

8092 Zurich,

Switzerland

T. Mitchell

North Carolina State University,

Department of Plant Pathology,

Campus Box 7251,

Raleigh,

NC 27695-7251,

USA



xii List of contributors

M. E. Monroe

Biological Separations and Mass Spectrometry, Pacific Northwest National Laboratory 3335 Q Avenue, Richland, WA 99352, USA

Y. Y. Oh

North Carolina State University, Department of Plant Pathology, Campus Box 7251, Raleigh, NC 27695–7251, USA

M. J. Orbach

Department of Plant Pathology, University of Arizona, Forbes Room 105, PO Box 210036, Tucson, AZ 85721–0036, USA

H. Pan *RTI*,

3040 Cornwallis Road, Research Triangle Park, NC 27709,

USA

E. A. Panisko

Fungal Biotechnology Team MSIN: K2–12,

Chemical and Biological Processes
Development Group,

Pacific Northwest National

Laboratory, 902 Battelle Blvd.,

Richland, WA 99352,

USA

E. J. Parker

Institute of Fundamental Sciences, Massey University, Private Bag 11 222, Palmerston North, New Zealand

A. Powell

North Carolina State University, Department of Plant Pathology, Campus Box 7251, Raleigh, NC 27695–7251, USA

M. Ruocco

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale, Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA), Italy

F. Scala

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale, Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA), Italy

C. E. Scheer

Kent ME4 4TB.

BEPHS Business Innovation, University of Greenwich at Medway, Central Avenue, Chatham Maritime.

UK



List of contributors

xiii

M. Schröder

School of Biological and Biomedical Sciences, Science Laboratories, South Road, University of Durham, Durham DH1 3LE, UK

B. Scott

Centre for Functional Genomics,
Institute of Molecular Biosciences
and Institute of Fundamental
Sciences,
Massey University,
Private Bag 11 222,
Palmerston North,
New Zealand

C. Soderlund

Director of Bioinformatics, Department of Plant Science, 303 Forbes Building, Tucson, AZ 85721, USA

I. Soriente

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale, Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA), Italy

H. Strasser

Institute of Microbiology, University of Innsbruck, Technikerstrasse 25, A-6020 Innsbruck, Austria

N. J. Talbot

Department of Biological Sciences,
University of Exeter,
Hatherly Laboratories,
Prince of Wales Road,
Exeter EX4 4PS,
UK

A. Tanaka

Institute of Molecular Biosciences, Massey University, Private Bag 11 222, Palmerston North, New Zealand

E. Thines

Institut für Biotechnologie und Wirkstoff-Forschung, Erwin-Schrödinger-Str. 56, D-67663 Kaiserslautern, Germany

M. Thon

Department of Computer Science, 320C Peterson Building MS# 2132, Texas A&M University, College Station, TX 77843–2132, USA

G. Turner

Department of Molecular Biology & Biotechnology, University of Sheffield, Sheffield SN10 2TN, UK

D. Turrà

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale,



xiv List of contributors

Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA), Italy

D. Veith

Institute for Applied
Life Sciences,
Applied Microbiology,
University of Karlsruhe,
Hertzstr. 16,
D-76187 Karlsruhe,
Germany

F. Vinale

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale, Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA), Italy

H. J. Weitz

School of Biological Sciences, Cruickshank Building, University of Aberdeen, Aberdeen AB24 3UU,

UK

C. F. Wend

Fungal Biotechnology Team MSIN: K2–12, Chemical and Biological Processes

Development Group, Pacific Northwest National

Laboratory,

902 Battelle Blvd., Richland, WA 99352, USA

S. L. Woo

Dipartimento di Arboricoltura, Botanica e Patologia Vegetale, Sezione Patologia Vegetale, Universita degli Studi di Napoli Federico II, Via Universita, 100–80055 Portici (NA), Italy

J.-R. Xu

Department of Botany and Plant Pathology, 915 West State Street, Purdue University, West Lafayette, IN 47906, USA

C. A. Young

Molecular Mycologist,
Forage Improvement Division,
The Samuel Roberts Noble
Foundation, Inc.,
2510 Sam Noble Pky,
Ardmore,
OK 73401,
USA

R. Zelger

The Research Centre for Agriculture and Forestry Laimburg, I-39051 Pfatten/Auer, Italy



Preface

The fungi are a highly diverse kingdom of eukaryotic microbes that have been exploited commercially for decades due to their ability to secrete large quantities of proteins of commercial value, and because they possess complex secondary metabolic pathways producing a diverse range of bioactive compounds that have had a major impact in the pharmaceutical market. For example, penicillin and cephalosporin (antibiotics), cylosporin (immunosuppressant) and more recently the statins (cholesterol reducing agents) are estimated to be worth over \$5 billion per annum, while enzymes and proteins produced commercially in fungal hosts are used in a diverse number of commercial markets including baking, brewing, detergent, textile and animal feed industries. In addition to the exploitation of fungal products, the fungi themselves are increasingly being developed as alternatives to conventional chemically based pest control strategies, as biocontrol agents active against commercially damaging insect pathogens and weeds, and as bioremediation agents capable of transforming organic and inorganic pollutants in the soil environment. Recent advances in the molecular genetics of the fungi together with the recent release of whole genome sequences of an increasing number of fungi will facilitate further the exploitation and commercialization of these important and ubiquitous eukaryotic microorganisms. The objective of this symposium volume is to highlight current and future biological, biochemical and molecular exploitation of the fungi in biotechnology, and act as an interface between current research and future commercialization.

The book is divided into five general parts, each dealing with an important area relevant to the overall topic, but with a considerable degree of complementarity between all the sections. The topics chosen provide a wide-ranging source of examples relating to the exploitation of fungi in a wide range of biomedical, industrial and environmental contexts. The first



xvi Preface

part, Comparative and functional fungal genomics, comprises chapters relating to genomic and proteomic analysis of industrial and plant pathogenic fungi, and how such information can be related to function. The second part on Bioactive molecules contains chapters that discuss the synthesis of some specific metabolites in fungi, e.g. pyridones and tetramic acids, as well as certain metabolites with important applications in agriculture. Molecular analysis of secondary metabolite production from endophytic fungi is also described together with a synthesis of modern genomic approaches that can elucidate secondary metabolism in Aspergillus species. The next part on Protein folding and secretion includes accounts of the roles of microtubules and motors in polarized fungal growth, cellular responses to protein-unfolding stress, as well as a genomic perspective on protein excretion and associated stresses in Aspergillus. The next part on Fungal bioremediation includes the Berkeley Award lecture on stress and the single cell, a topic relevant to many other research areas as well as fungal responses to toxicants. Other chapters provide accounts of novel bioluminescent fungal sensors for pollutants in the environment as well as the theory and practice of fungal bioremediation of xenobiotics and toxic metal pollutants. The final part on Fungal biocontrol of pests has accounts of the fungal control of subterranean pests, the development of mycoherbicides, interactions between *Trichoderma* spp. with plant hosts and fungal pathogens, and fungal parasites of invertebrates.

This book arises from the British Mycological Society Symposium on *Exploitation of Fungi* held at the University of Manchester in September 2005 and the editors would like to thank those authors who contributed enthusiastically to this book. Special thanks also go to Diane Purves in Dundee, who greatly assisted communication, collation, editing and formatting of chapters. The prime objective was to produce a wide-ranging volume that would highlight the importance of fungi in modern biotechnology as well as highlighting the modern approaches and tools that are now used to understand and manipulate fungal biology. We think that this has been achieved and hope the volume has broad appeal not only to mycologists of all persuasions but also to other microbiologists, biotechnologists and molecular biologists, who routinely rely on fungal model systems. It will be interesting to see how genomic revelations influence developments over the next few years.

Geoffrey D. Robson Pieter van West Geoffrey M. Gadd