RNAi

METHODS IN MOLECULAR BIOLOGYTM

John M. Walker, Series Editor

- **465. Mycobacteria**, *Second Edition*, edited by *Tanya Parish and Amanda Claire Brown*, 2008
- Essential Concepts in Toxicogenomics, edited by Donna L. Mendrick and William B. Mattes, 2008
- **459. Prion Protein Protocols**, edited by *Andrew F. Hill*, 2008
- **458.** Artificial Neural Networks: Methods and Applications, edited by David S. Livingstone, 2008
- **457. Membrane Trafficking**, edited by *Ales Vancura*, 2008
- **456. Adipose Tissue Protocols**, *Second Edition*, edited by *Kaiping Yang*, 2008
- **455.** Osteoporosis, edited by Jennifer J. Westendorf,
- 2008
 454. SARS- and Other Coronaviruses: Laboratory
- Protocols, edited by Dave Cavanagh, 2008
 Bioinformatics, Volume 2: Structure, Function, and Applications, edited by Jonathan M. Keith, 2008
- **452. Bioinformatics, Volume 1:** Data, Sequence Analysis, and Evolution, edited by Jonathan M. Keith. 2008
- **451.** Plant Virology Protocols: From Viral Sequence to Protein Function, edited by Gary Foster, Elisabeth Johansen, Yiguo Hong, and Peter Nagy, 2008
- **450. Germline Stem Cells**, edited by *Steven X. Hou and Shree Ram Singh*, 2008
- 449. Mesenchymal Stem Cells: Methods and Protocols, edited by Darwin J. Prockop, Douglas G. Phinney, and Bruce A. Brunnell, 2008
- 448. Pharmacogenomics in Drug Discovery and Development, edited by *Qing Yan*, 2008
- **447. Alcohol:** *Methods and Protocols*, edited by *Laura E. Nagy*, 2008
- 446. Post-translational Modification of Proteins: Tools for Functional Proteomics, Second Edition, edited by Christoph Kannicht, 2008
- **445. Autophagosome and Phagosome**, edited by *Vojo Deretic*, 2008
- **444. Prenatal Diagnosis**, edited by *Sinuhe Hahn and Laird G. Jackson*, 2008
- 443. Molecular Modeling of Proteins, edited by Andreas Kukol, 2008
- **442. RNAi:** *Design and Application*, edited by *Sailen Barik*, 2008
- **439. Genomics Protocols:** *Second Edition*, edited by *Mike Starkey and Ramnanth Elaswarapu*, 2008
- **438.** Neural Stem Cells: Methods and Protocols, Second Edition, edited by Leslie P. Weiner, 2008
- **437. Drug Delivery Systems**, edited by *Kewal K. Jain*, 2008
- **436. Avian Influenza Virus**, edited by *Erica Spackman*, 2008
- **435.** Chromosomal Mutagenesis, edited by *Greg Davis* and *Kevin J. Kayser*, 2008
- **434. Gene Therapy Protocols: Volume 2:** Design and Characterization of Gene Transfer Vectors, edited by Joseph M. LeDoux, 2008
- 433. Gene Therapy Protocols: Volume 1: Production and In Vivo Applications of Gene Transfer Vectors, edited by Joseph M. LeDoux, 2007

- **432.** Organelle Proteomics, edited by *Delphine Pflieger* and *Jean Rossier*, 2008
- **431. Bacterial Pathogenesis:** *Methods and Protocols*, edited by *Frank DeLeo and Michael Otto*, 2008
- **430. Hematopoietic Stem Cell Protocols**, edited by *Kevin D. Bunting*, 2008
- **429. Molecular Beacons:** Signalling Nucleic Acid Probes, Methods and Protocols, edited by Andreas Marx and Oliver Seitz, 2008
- **428.** Clinical Proteomics: Methods and Protocols, edited by Antonio Vlahou, 2008
- **427. Plant Embryogenesis**, edited by *Maria Fernanda Suarez and Peter Bozhkov*, 2008
- **426. Structural Proteomics:** *High-Throughput Methods*, edited by *Bostjan Kobe, Mitchell Guss, and Huber Thomas*, 2008
- **425. 2D PAGE: Volume 2:** Applications and Protocols, edited by Anton Posch, 2008
- **424. 2D PAGE: Volume 1:** Sample Preparation and Pre-Fractionation, edited by Anton Posch, 2008
- **423.** Electroporation Protocols, edited by *Shulin Li*, 2008
- 422. Phylogenomics, edited by William J. Murphy, 2008
- **421. Affinity Chromatography:** *Methods and Protocols, Second Edition,* edited by *Michael Zachariou, 2008*
- **420. Drosophila:** *Methods and Protocols*, edited by *Christian Dahmann*, 2008
- **419. Post-Transcriptional Gene Regulation**, edited by *Jeffrey Wilusz*, 2008
- **418. Avidin-Biotin Interactions:** *Methods and Applications,* edited by *Robert J. McMahon, 2008*
- **417.** Tissue Engineering, Second Edition, edited by Hannsjörg Hauser and Martin Fussenegger, 2007
- **416.** Gene Essentiality: Protocols and Bioinformatics, edited by Svetlana Gerdes and Andrei L. Osterman, 2008
- **415. Innate Immunity**, edited by *Jonathan Ewbank and Eric Vivier*, 2007
- **414. Apoptosis in Cancer:** *Methods and Protocols*, edited by *Gil Mor and Ayesha Alvero*, 2008
- Protein Structure Prediction, Second Edition, edited by Mohammed Zaki and Chris Bystroff, 2008
- 412. Neutrophil Methods and Protocols, edited by Mark T. Quinn, Frank R. DeLeo, and Gary M. Bokoch, 2007
- Reporter Genes for Mammalian Systems, edited by Don Anson. 2007
- **410.** Environmental Genomics, edited by *Cristofre C. Martin*, 2007
- **409.** Immunoinformatics: Predicting Immunogenicity In Silico, edited by Darren R. Flower, 2007
- **408.** Gene Function Analysis, edited by *Michael Ochs*, 2007
- **407. Stem Cell Assays**, edited by *Vemuri C. Mohan*, 2007
- **406. Plant Bioinformatics:** *Methods and Protocols*, edited by *David Edwards*, 2007
- **405. Telomerase Inhibition:** *Strategies and Protocols*, edited by *Lucy Andrews and Trygve O. Tollefsbol*, 2007

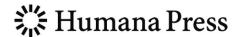
RNAi

Design and Application

Edited by

Sailen Barik

Department of Biochemistry and Molecular Biology, University of South Alabama, College of Medicine, Mobile, Alabama



Editor
Sailen Barik
Department of Biochemistry
and Molecular Biology
College of Medicine
University of South Alabama
Mobile, Alabama

Series Editor
John. M. Walker
School of Life Sciences
University of Hertfordshire
Hatfield, Hertfordshire, UK

ISBN: 978-1-58829-874-4 e-ISBN: 978-1-59745-191-8

ISSN:1064-3745 e-ISSN: 1940-6029

Library of Congress Control Number: 2007940759

© 2008 Humana Press, a part of Springer Science+Business Media, LLC

All rights reserved. This work may not be translated or copied in whole or in part without the written permission of the publisher (Humana Press, 999 Riverview Drive, Suite 208, Totowa, NJ 07512 USA), except for brief excerpts in connection with reviews or scholarly analysis. Use in connection with any form of information storage and retrieval, electronic adaptation, computer software, or by similar or dissimilar methodology now known or hereafter developed is forbidden.

The use in this publication of trade names, trademarks, service marks, and similar terms, even if they are not identified as such, is not to be taken as an expression of opinion as to whether or not they are subject to proprietary rights.

While the advice and information in this book are believed to be true and accurate at the date of going to press, neither the authors nor the editors nor the publishers can accept any legal responsibility for any errors or omissions that may be made. The publisher makes no warranty, express or implied, with respect to the material contained herein.

Cover illustration: Figure 3, Chapter 16, "Temporal Control of Gene Silencing by in ovo Electroporation" by Thomas Baeriswyl, Olivier Mauti, and Esther T. Stoeckli. Figure 3, Chapter 17, "Altering Flower Color in Transgenic Plants by RNAi-Mediated Engineering of Flavonoid Biosynthetic Pathway," by Yoshikazu Tanaka, Noriko Nakamura, and Junichi Togami.

Printed on acid-free paper

987654321

springer.com



Preface

RNA interference (RNAi), in which RNA silences RNA, is the most recent discovery to revolutionize biology and to be recognized by a Nobel Prize (in 2006, to Andrew Fire and Craig Mello). It is a story that began with historic observations in plants and fungi and eventually worked its way up to humans. If one were to describe the major steps of RNAi very briefly, it would read as follows: RNAi is triggered by double-stranded RNA (dsRNA), produced endogenously or introduced by scientists —> Long dsRNA is trimmed into short interfering RNA or microRNA (siRNA or miRNA) by Dicer —> The individual strands of the si/miRNA then guide the assembly of a multiprotein complex, known as RISC, the key constituent of which is Argonaute —> Depending on the extent of homology of the guide RNA to the target, RISC either destroys the target RNA or suppresses its translation, leading to gene silencing.

The chapters in *RNAi*: *Design and Application*, contributed by leaders in the field, sum up the state-of-the-art methods on practical, everyday use of RNAi in biological research. Although multiple books and monographs have been published on RNAi, there is a noticeable dearth of bench protocols that can be used quickly and easily by beginners aspiring to enter this new field. This volume aims to fill that void.

RNAi: Design and Application is divided into two parts. The first and smaller part (chapters 1–4) covers the fundamentals including designs of RNAi, biochemical assay protocols for the major components of RNAi, and study of potential off-target effects. The larger second part (chapters 5–18) covers various applications of RNAi in diverse model organisms and systems, from antiviral and anticancer applications to altering flower color in plants. Armed with this volume, a researcher with standard molecular biological training should be able to perform today's major RNAi-related experiments and carry out gene knock-down analyses in virtually any cell line or species of interest. In the established tradition of the Methods in Molecular BiologyTM series, each chapter contains step-by-step protocols, extra notes, and problem-solving tips, which are usually not found in original research papers. As the horizon of RNAi application is rapidly broadening, we have strived to offer the most recent protocols in each area so that they remain useful for years to come.

viii Preface

My sincere thanks go to all the authors and the Humana staff for bringing it all together, and to Professor John M. Walker for his guidance. I remain indebted to my wife, Kumkum, and my children, Titus and Tiasha, for their immeasurable support and encouragement.

Sailen Barik

Contents

Prefac	e	vii
Contri	butors	хi
Part 1.		3
2.	Expression, Purification, and Analysis of Recombinant Drosophila Dicer-1 and Dicer-2 Enzymes Xuecheng Ye and Qinghua Liu	11
3.	In vitro RNA Cleavage Assay for Argonaute-Family Proteins Keita Miyoshi, Hiroshi Uejima, Tomoko Nagami-Okada, Haruhiko Siomi, and Mikiko C. Siomi	29
4.	Identifying siRNA-Induced Off-Targets by Microarray Analysis *Emily Anderson, Queta Boese, Anastasia Khvorova, *and Jon Karpilow***	45
PART 5.	II: APPLICATION OF RNAI IN DIVERSE ORGANISMS Hydrodynamic Delivery of siRNA in a Mouse Model of Sepsis Doreen E. Wesche-Soldato, Joanne Lomas-Neira, Mario Perl, Chun-Shiang Chung, and Alfred Ayala	67
6.	Nasal Delivery of siRNA Vira Bitko and Sailen Barik	<i>75</i>
7.	RNA Interference as a Genetic Tool in Trypanosomes Vivian Bellofatto and Jennifer B. Palenchar	83
8.	Lentivirus-Mediated RNA Interference in Mammalian Neurons Scott Q. Harper and Pedro Gonzalez-Alegre	95
9.	Silencing Genes by RNA Interference in the Protozoan Parasite Entamoeba histolytica Carlos F. Solis and Nancy Guillén	113

x Contents

10.	Use of RNAi in <i>C. elegans</i> Tsuyoshi Ohkumo, Chikahide Masutani, Toshihiko Eki, and Fumio Hanaoka
11.	Application of siRNA Against SARS in the Rhesus Macaque Model
	Qingquan Tang, Baojian Li, Martin Woodle, and Patrick Y. Lu 139
12.	siRNA and shRNA as Anticancer Agents in a Cervical Cancer Model
	Wenyi Gu, Lisa Putral, and Nigel McMillan159
13.	Identification and Expression Analysis of Small RNAs During Development
	Toshiaki Watanabe, Hiroshi Imai, and Naojiro Minami 173
14.	Screening and Identification of Virus-Encoded RNA Silencing Suppressors
	Sumona Karjee, Mohammad Nurul Islam, and Sunil K. Mukherjee187
15.	Application of RNA Interference in Functional Genomics Studies of a Social Insect
	Michael E. Scharf, Xuguo Zhou,
	and Margaret A. Schwinghammer
16.	Temporal Control of Gene Silencing by <i>in ovo</i> Electroporation <i>Thomas Baeriswyl, Olivier Mauti, and Esther T. Stoeckli</i>
17.	Altering Flower Color in Transgenic Plants by RNAi-Mediated Engineering of Flavonoid Biosynthetic Pathway
	Yoshikazu Tanaka, Noriko Nakamura, and Junichi Togami 245
18.	Transgenic RNA Interference in Mice
	Pumin Zhang 259
Index .	

Contributors

- Mohammed Amarzguioui The Biotechnology Centre of Oslo, Oslo, Norway
- EMILY ANDERSON Dharmacon, ThermoFisher Scientific, Lafayette, CO
- Alfred Ayala Division of Surgical Research, Department of Surgery, Rhode Island Hospital/Brown University School of Medicine, Providence, RI
- Thomas Baeriswyl Institute of Zoology, University of Zürich, Zürich, Switzerland
- Sailen Barik Department of Biochemistry and Molecular Biology, University of South Alabama, College of Medicine, Mobile, AL
- VIVIAN BELLOFATTO Department of Microbiology and Molecular Genetics, UMDNJ-NJ Medical School, International Center for Public Health, Newark, NJ
- VIRA BITKO Department of Biochemistry and Molecular Biology, University of South Alabama, College of Medicine, Mobile, AL
- QUETA BOESE Dharmacon, ThermoFisher Scientific, Lafayette, CO
- Chun-Shiang Chung Division of Surgical Research, Department of Surgery, Rhode Island Hospital/Brown University School of Medicine, Providence, RI
- Toshihiko Eki Department of Ecological Engineering, Toyohashi University of Technology, Toyohashi, Japan
- Pedro Gonzalez-Alegre Department of Neurology, Carver College of Medicine at The University of Iowa, Iowa City, IA
- Wenyi Gu Cancer Biology Program, Centre for Immunology and Cancer Research, Princess Alexandra Hospital, University of Queensland, Brisbane, Australia
- NANCY GUILLÉN Unité de Biologie Cellulaire du Parasitisme, Institut Pasteur, Paris, France
- Fumio Hanaoka Graduate School of Frontier Biosciences, Osaka University, and SORST, Japan Science and Technology Agency, Osaka, Japan

xii Contributors

Scott Q. Harper • Center for Gene Therapy, Department of Pediatrics, The Ohio State University, Columbus, OH

- HIROSHI IMAI Laboratory of Reproductive Biology, Department of Agriculture, Kyoto University, Kyoto, Japan
- Mohammad Nurul Islam International Center for Genetic Engineering and Biotechnology, PMB Lab, New Delhi, India
- Sumona Karjee International Center for Genetic Engineering and Biotechnology, PMB Lab, New Delhi, India
- JON KARPILOW Dharmacon, ThermoFisher Scientific, Lafayette, CO
- Anastasia Khvorova Dharmacon, ThermoFisher Scientific, Lafayette, CO
- BAOJIAN LI Top Genomics, Ltd., and College of Life Sciences, Sun Yat-sen University, Guangzhou, China
- QINGHUA LIU Department of Biochemistry, University of Texas Southwestern Medical Center, Dallas, TX
- Joanne Lomas-Neira Division of Surgical Research, Department of Surgery, Rhode Island Hospital/Brown University School of Medicine, Providence, RI
- PATRICK Y. Lu Sirnaomics, Inc., Rockville, MD
- Chikahide Masutani Graduate School of Frontier Biosciences, Osaka University; and SORST, Japan Science and Technology Agency, Osaka, Japan
- OLIVIER MAUTI Institute of Zoology, University of Zürich, Zürich, Switzerland
- NIGEL McMILLAN Cancer Biology Program, Centre for Immunology and Cancer Research, Princess Alexandra Hospital, University of Queensland, Brisbane, Australia
- Naojiro Minami Laboratory of Reproductive Biology, Department of Agriculture, Kyoto University, Kyoto, Japan
- Keita Miyoshi Institute for Genome Research, University of Tokushima, Tokushima, Japan
- Sunil K. Mukherjee International Center for Genetic Engineering and Biotechnology, PMB Lab, New Delhi, India
- Tomoko Nagami-Okada Institute for Genome Research, University of Tokushima, Tokushima, Japan
- NORIKO NAKAMURA Institute for Advanced Core Technology, Suntory Ltd., Osaka, Japan
- Tsuyoshi Онкимо Graduate School of Frontier Biosciences, Osaka University; and SORST, Japan Science and Technology Agency, Osaka, Japan

Contributors xiii

Jennifer B. Palenchar • Department of Chemistry, Villanova University, Villanova, PA

- Mario Perl Universitätsklinikum Ulm, Zentrum für Chirurgie, Klinik für Unfallchirurgie, Hand-, Plastische- und Wiederherstellungschirurgie, Ulm, Germany
- LISA PUTRAL Cancer Biology Program, Centre for Immunology and Cancer Research, Princess Alexandra Hospital, University of Queensland, Brisbane, Australia
- JOHN J. ROSSI Division of Molecular Biology, Beckman Research Institute of the City of Hope, Duarte, CA
- MICHAEL E. SCHARF Molecular and Applied Insect Toxicology, Entomology and Nematology Department, University of Florida, Gainesville, FL
- Margaret A. Schwinghammer Department of Entomology, Purdue University, West Lafayette, IN
- HARUHIKO SIOMI Keio University School of Medicine, Tokyo, Japan
- MIKIKO C. SIOMI Institute for Genome Research, University of Tokushima, JST, CREST, Tokushima, Japan
- CARLOS F. Solis Unité de Biologie Cellulaire du Parasitisme, Institut Pasteur, Paris, France
- Esther T. Stoeckli Institute of Zoology, University of Zürich, Zürich, Switzerland
- Yoshikazu Tanaka Institute for Advanced Core Technology, Suntory Ltd., Osaka, Japan
- QINGQUAN TANG OriGene Technologies, Inc., Rockville, MD
- Junichi Togami Institute for Advanced Core Technology, Suntory Ltd., Osaka, Japan
- HIROSHI UEJIMA Institute for Genome Research, University of Tokushima, Tokushima, Japan
- Toshiaki Watanabe Division of Human Genetics, Department of Integrated Genetics, National Institute of Genetics, Research Organization of Information and Systems; and Department of Genetics, School of Life Science, The Graduate University for Advanced Studies (SOKENDAI), Mishima, Japan
- Doreen E. Wesche-Soldato Division of Surgical Research, Department of Surgery, Rhode Island Hospital/Brown University School of Medicine, Providence, RI
- MARTIN WOODLE Nanotides Pharmaceuticals, Inc., Rockville, MD
- Xuecheng Ye Department of Biochemistry, University of Texas Southwestern Medical Center, Dallas, TX

xiv Contributors

Pumin Zhang • Department of Molecular Physiology and Biophysics, Baylor College of Medicine, Houston, TX

Xuguo Zhou • Molecular and Applied Insect Toxicology, Entomology and Nematology Department, University of Florida, Gainesville, FL