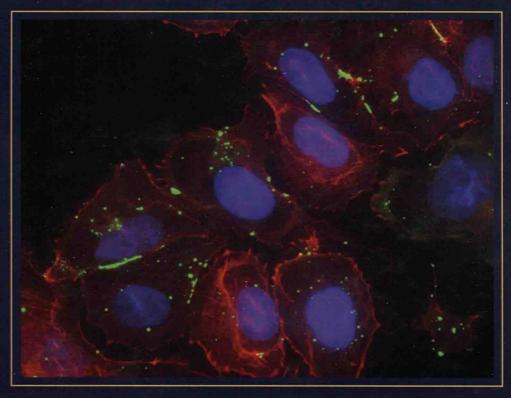
Gap Junction Channels and Hemichannels



Edited by Donglin Bai and Juan C. Sáez



METHODS IN SIGNAL TRANSDUCTION SERIES



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Series Preface

The concept of signal transduction is now long established as a central tenet of biological sciences. Since the inception of the field close to 50 years ago, the number and the varieties of signal transduction pathways, cascades, and networks have steadily increased and now constitute what is often regarded as a bewildering array of mechanisms by which cells sense and respond to extracellular and intracellular environmental stimuli. It is not an exaggeration to state that virtually every cell function is dependent on the detection, amplification, and integration of these signals. Moreover, there is increasing appreciation that in many disease states, aspects of signal transduction are critically perturbed.

Our knowledge of how information is conveyed and processed through these cellular molecular circuits and biochemical switches has increased enormously in scope and complexity since this series was initiated 15 years ago. Such advances would not have been possible without the supplementation of older technologies, drawn chiefly from cell and molecular biology, biochemistry, physiology, and pharmacology, with newer methods that make use of sophisticated genetic approaches as well as structural biology, imaging, bioinformatics, and systems biology analysis.

The overall theme of this series continues to be the presentation of the wealth of up-to-date research methods applied to the many facets of signal transduction. Each volume is assembled by one or more editors who are preeminent in their specialty. In turn, the guiding principle for editors is to recruit chapter authors who will describe procedures and protocols with which they are intimately familiar in a reader-friendly format. The intent is to assure that each volume will be of maximum practical value to a broad audience, including students and researchers just entering this area, as well as seasoned investigators.

It is hoped that the information contained in the books of this series will constitute a useful resource to the life sciences research community well into the future.

Joseph Eichberg Michael Xi Zhu Series Editors

Preface

Gap junction channels are a group of intercellular channels ubiquitously expressed in tissues and organs to synchronize many physiological processes. Hemichannels were discovered about two decades ago (a hemichannel is half of a gap junction channel) as a relevant cell membrane pathway for the release and uptake of ions and small molecules, many of which are autocrine and paracrine signal molecules. Mutations in 14 out of 21 genes encoding gap junction protein subunits, called connexins, have been found to be associated with inherited human disorders and diseases, including hearing loss, skin diseases, peripheral and central neuropathic disorders, lens cataracts, cardiac arrhythmias, and developmental disorders. The prevalence of connexin 26 (Cx26) gene mutation-linked congenital sensorineural hearing loss is so high that complete DNA sequencing of the coding region of GJB2 (the gene encoding Cx26) has become routine for newborns or during pregnancy. Because of the important physiological role of connexins and the linkage of the connexin gene mutations with many diseases, the field of connexin research has recently exploded and has become one of the most active areas of research. Numerous novel approaches and techniques have been developed, but there is no single book dedicated on the unique techniques and protocols for the research on these large pore channels. The last book on gap junction methods and protocols was published 15 years ago, which cannot meet the current need of researchers and trainees in the fields. To fill this gap, we recruited expert researchers in the field to share their state-of-the-art techniques, protocols, and thoughts/reviews on various approaches commonly used in the studies of gap junction channels and hemichannels.

Chapters 1 to 6 in this book are focused on the protocols on, approaches to, and reviews of gap junction channels, and Chapters 7 to 12 are on connexin hemichannels. We hope that this book will be a very useful reference book for graduate students, postdoctoral fellows, and researchers with an interest in gap junction channels and hemichannels. It should be noted that we have covered many common approaches in studying these channels, but some areas/approaches may not be discussed in detail in this book because of the difficulties in finding appropriate contributors.

We would like to thank all the contributors for taking their valuable time to put their chapters together to share their detailed experimental protocols and their views on different approaches. Without their willingness to help, consistent efforts in developing and revising their chapters, and their contribution in helping to reviewing some chapters, we would not be able to complete this book.

A video that relates to Chapter 3 is available from the CRC Press Web site. Interested readers should click on the Downloads/Updates tab at the following URL: https://www.crcpress.com/Gap-Junction-Channels-and-Hemichannels/Bai-Sez/9781498738620.

x Preface

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Editors

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Contents

	ix
	X
	xiii
Chapter 1	Immunofluorescence: Applications for Analysis of Connexin Distribution and Trafficking
	Sandra A. Murray and T. I. Shakespeare
Chapter 2	Imaging Gap Junctions in Living Cells
	Matthias M. Falk, Charles G. Fisher, Rachael M. Kells Andrews, and Tia J. Kowal
Chapter 3	FRAP for the Study of Gap Junction Nexus Macromolecular Organization
	Randy F. Stout, Jr. and David C. Spray
Chapter 4	Patch Clamp Analysis of Gap Junction Channel Properties93
	Donglin Bai and John A. Cameron
Chapter 5	What Do You Need to Measure Gap Junctional Permselectivity?
	José F. Ek Vitorín
Chapter 6	Recording Gap Junction-Mediated Synaptic Transmission in Vivo at Mixed Synapses on the Goldfish Mauthner Cells 151
	Roger Cachope, Sebastian Curti, and Alberto E. Pereda
Chapter 7	Assessing Connexin Hemichannel Function during Ischemic Injury and Reperfusion
	Yeri Kim and Colin R. Green

vi Contents

Chapter 8	Whole-Cell Patch Clamp Recordings of Unitary Connexin 43 Hemichannel Currents
	Nan Wang, Alessio Lissoni, Maarten De Smet, Karin R. Sipido, and Luc Leybaert
Chapter 9	Functional Characterization of Connexin Hemichannels Using Xenopus Oocytes and the Two-Electrode Voltage Clamp Technique
Chapter 10	Functional Assays of Purified Connexin Hemichannels
Chapter 11	Methods to Determine Formation of Heteromeric Hemichannels
Chapter 12	Methods to Examine the Role of Gap Junction and Pannexin Channels in HIV Infection
Index	275

1 Immunofluorescence: Applications for Analysis of Connexin Distribution and Trafficking

Sandra A. Murray and T. I. Shakespeare

CONTENTS

1.1	Summ	lary	4
1.2		uction	
1.3		ials and Reagents	
	1.3.1	Fixatives	
	1.3.2	Permeabilization Solutions	5
	1.3.3	Blocking Solutions	5
	1.3.4	Primary Antibodies	
	1.3.5	Secondary Antibodies	6
	1.3.6	Nuclear Stains	7
	1.3.7	Membrane Stains	7
	1.3.8	Actin Stain	7
1.4	Equip	ment	8
	1.4.1	Humidified Chamber	8
	1.4.2	Microscopes	8
1.5	Locali	zation of Connexin in Cell Cultures	
	1.5.1	Acid Cleaning Coverslips for Growing Cells	8
	1.5.2	Cell Culture Protocol	
	1.5.3	Cell Fixation Protocol	9
	1.5.4	Cell Permeabilization	10
	1.5.5	Blocking	10
	1.5.6	Immunolabeling of Connexin Proteins: Cells on Coverslips	
	1.5.7	Double Labeling of Cells on Coverslips (Optional)	
	1.5.8	Actin Staining (Optional)	
	1.5.9	Nuclear Labeling of Cells on Coverslips (Optional)	
		Mounting the Coverslips on Slides	
1.6		nolabeling of Connexin Proteins: Tissue Sections	
	1.6.1	Quick-Freeze Method	
	1.6.2	Tissue Staining	
	1.6.3	Double Labeling of Tissue Sections (Optional)	13

	1.6.4	Nuclear Labeling of Tissue Sections (Optional)	13
	1.6.5	Mounting Coverslips over Tissue Sections	14
1.7	Local	izing Connexins with Quantum Dots (Qdot): Light and	
	Transi	mission Electron Microscopy	14
	1.7.1	Immunolabeling Connexins	14
	1.7.2	Light Microscopy	16
	1.7.3	Transmission Electron Microscopy	16
	1.7.4	Important Considerations	17
Ackı		gments	
Refe	rences.		17

1.1 SUMMARY

Immunofluorescence methods and techniques for visualizing the subcellular location of gap junction proteins (connexins) and their associated proteins will be presented in this chapter. Specifically, a step-by-step guide for localizing connexin in cell cultures and tissues with immunofluorescence will be presented. The concepts, rationale, and advantages of using particular protocols will be discussed. The methods of preparing and processing cells and tissues for immunofluorescence will be provided. This chapter will serve as a guide for researchers interested in the morphological analysis of gap junctions.

1.2 INTRODUCTION

Gap junctions, which are membrane channels composed of proteins called connexins (Goodenough et al. 1996), permit intercellular communication of regulatory molecules between contacting cells (Decker et al. 1978; Munari-Silem et al. 1995; Oyoyo et al. 1997; Shah and Murray 2001). Information on the assembly of connexins into functional channels and the removal of these channels from the plasma membrane (reviewed in the schematic seen in Figure 1.1) is critical to understanding gap junction turnover, the regulation of cell—cell communication, and the possible role of connexins as anchors for scaffold molecules. To study gap junction assembly and turnover, it is necessary to be able to identify the localization of gap junction proteins at the cell surface and within the cytoplasm (as seen in Figure 1.2). While biochemical methods are powerful tools for detecting the presence, abundance, and types of connexin found in cell populations, immunofluorescence provides information on the specific compartments and cellular locations of connexin and associated proteins needed to study gap junction trafficking and turnover.

Immunofluorescence is a relatively straightforward and inexpensive method for detecting the presence and the subcellular location of connexins within cells. Variations in fluorescent stain intensity and location can be used to gain insight into connexin function and cell-cell communication in cell populations. By using a specific primary antibody that binds to connexin and a secondary antibody that has been conjugated to a fluorophore, connexins can be visualized with a fluorescence microscope (Oyoyo et al. 1997). Colocalization procedures for detecting the

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