Dictyostelium

Evolution, Cell Biology, and the Development of Multicellularity

RICHARD H. KESSIN

Columbia University

Bibliography by Jakob Franke *Columbia University*



PUBLISHED BY THE PRESS SYNDICATE OF THE UNIVERSITY OF CAMBRIDGE The Pitt Building, Trumpington Street, Cambridge, United Kingdom

CAMBRIDGE UNIVERSITY PRESS

The Edinburgh Building, Cambridge CB2 2RU, UK
40 West 20th Street, New York, NY 10011-4211, USA
10 Stamford Road, Oakleigh, Melbourne 3166, Australia
Ruiz de Alarcón 13, 28014 Madrid, Spain

© Cambridge University Press 2001

This book 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 2001

Printed in the United Kingdom at the University Press, Cambridge

Typeface Times Roman 10/12 System 3B2 [KW]

A catalog record for this book is available from the British Library

Library of Congress Cataloging-in-Publication Data

Kessin, Richard H., 1944-

Dictyostelium: evolution, cell biology, and the development of multicellularity / Richard H. Kessin; bibliography by Jakob Franke.

p. cm – (Developmental and cell biology series)

Includes bibliographical references.

1. Dictyostelium. I. Title. II. Series.

QK635.D5 K47 2001

579.4'32 – dc21 00-037820

ISBN 0 521 58364 0 hardback

Contents

Preface

A Bri	ief Introduction to Dictyostelium discoideum and its Relatives	1
A History of Research on Dictyostelium discoideum		
2.1	The classical experiments of Kenneth Raper	12
2.2	Chemotaxis and aggregation	16
2.3	Biochemistry and molecular biology	18
The Evolutionary Biology of Dictyostelium		
3.1	A digression into ecology	20
3.2	Soil amoebae have predators	21
3.3	The amoebae can respond to starvation in three ways	22
	3.3.1 The microcyst	25
	3.3.2 The macrocyst	25
	3.3.3 Fruiting bodies	27
3.4	The forms of development evolved in a sequence	27
3.5	Genetic heterogeneity in wild populations	28
3.6		29
3.7	÷ 7	30
3.8		33
3.9		33
3.10	Molecular phylogeny	35
	A His 2.1 2.2 2.3 The I 3.1 3.2 3.3 3.4 3.5 3.6 3.7 3.8 3.9	A History of Research on Dictyostelium discoideum 2.1 The classical experiments of Kenneth Raper 2.2 Chemotaxis and aggregation 2.3 Biochemistry and molecular biology The Evolutionary Biology of Dictyostelium 3.1 A digression into ecology 3.2 Soil amoebae have predators 3.3 The amoebae can respond to starvation in three ways 3.3.1 The microcyst 3.3.2 The macrocyst 3.3.3 Fruiting bodies 3.4 The forms of development evolved in a sequence 3.5 Genetic heterogeneity in wild populations 3.6 The evolution of cooperativity 3.7 The cells in a Dictyostelium aggregate compete to form spores 3.8 Size limitations in an aggregative organism 3.9 The extraordinary parasitism of D. caveatum

xii

viii CONTENTS

4	The	Genome and Genetics	40			
	4.1	The genome is relatively small and will be sequenced soon	40			
	4.2	The ribosomal genes are coded in an extra-chromosomal				
		palindrome	43			
	4.3	Dictyostelium species contain several families of replicating				
		plasmids	44			
	4.4	The genome is littered with transposable elements	45			
		4.4.1 Tdd-2 and Tdd-3	46			
		4.4.2 DIRS-1	46			
		4.4.3 DRE	47			
		4.4.4 Skipper	47			
	4.5	The mitochondrial genome	48			
	4.6	Maintaining the genome – the DNA repair mechanisms of				
		Dictyostelium	49			
	4.7	Molecular genetics	50			
	4.8	Mutagenesis	51			
	4.9	Restriction enzyme-mediated integration	53			
	4.10		55			
		4.10.1 Parasexual genetics	55			
		4.10.2 Sexual recombination	57			
5	Mem	Membranes and Organelles of <i>Dictyostelium</i> 5				
	5.1	The plasma membrane	59			
	5.2	Channels and pumps of the plasma membrane	61			
	5.3	Membrane systems that transiently connect to the plasma				
		membrane	61			
	5.4	Axenic cells feed by macropinocytosis	64			
	5.5	Phagocytosis	66			
	5.6	Lysosomes	67			
	5.7	Endoplasmic reticulum, Golgi, and nuclei	68			
	5.8	Mitochondria and peroxisomes	69			
	5.9	The autophagic vacuole	69			
6	Cell	Motility and the Cytoskeleton	70			
	6.1	Actin and its binding proteins	71			
		6.1.1 G-actin-binding proteins	72			
		6.1.2 F-actin-binding proteins	72			
		6.1.3 Cross-linking proteins	74			
		6.1.4 Attaching to the membrane	75			
	6.2	Myosin motors – conventional myosin	76			
	6.3	Myosin motors – the unconventional myosins	78			
	6.4	Building and retracting the pseudopod	79			
	6.5	Strengthening the filaments of actin	82			
	6.6	Moving the cell	82			
	6.7	Signaling to the cytoskeleton	84			
	6.8	Cytokinesis	85			

			CONTENTS	s ix	
	6.9	The m	nicrotubule cytoskeleton	86	
7			on from Growth to Development: From Starvation to	00	
			ing cAMP Signal Relay	89	
	7.1		can detect imminent starvation	89	
	7.2		th-specific events cease during development	92	
	7.3		rst events after starvation	93	
	7.4	YakA	kinase regulates the growth to development transition	95	
8	Chemotaxis and Aggregation				
	8.1	An ov	verview	98	
8.2		There	are several ways to study the cellular response to		
		cAMI	P binding	99	
8.3 Second messengers and cytoskeletal events can be studied		d messengers and cytoskeletal events can be studied			
		with s	suspended cells	100	
	8.4	*			
		pathw	ay	103	
		8.4.1	The cAR1 receptor is essential to aggregation	105	
		8.4.2	The cAR1 protein has several important domains	108	
		8.4.3	1 1 0		
		0.4.4	pathways	111	
		8.4.4	The $\beta\gamma$ subunit stimulates adenylyl cyclase (ACA) in	112	
		0.4.5	conjunction with other proteins	112	
		8.4.5		114	
		8.4.6			
			affects the activity of cAMP-dependent protein kinase	115	
		0.47	(PKA)	115	
		8.4.7	e		
	0.5	7 01 1	events in the cytoplasm	115	
	8.5		evelopmental regulation of chemotactic components	116	
	8.6		and calcium mobilize the cytoskeleton	116	
	8.7		P controls motility during chemotaxis	118	
	8.8		moebae control extracellular cAMP by secreting a		
			and a PDE inhibitor (PDI)	121	
	8.9	The c	ells can sense density and aggregate size	125	

127

128

131

135

139

140

141

8.10 Auto-induction of the chemotactic response

8.12 Polarity of movement and the polarity of G-protein

8.13 Mathematical simulations to explain signaling in

Amoebae form a sheath during aggregation

The transition from aggregation to loose mound is

mediated by a transcription factor named GBF

8.11 Mobilizing the chemotactic machinery

9 Differentiation and Adhesion in the Aggregate

activation

9.1

9.2

Dictyostelium

x CONTENTS

	9.3	The discovery of cell type-specific genes provided important				
		tools to study pattern formation	143			
	9.4	Pattern formation begins in the mound	144			
	9.5	The position of PstA, PstO, and PstAB cells	146			
	9.6	Differentiation inducing factor (DIF) and the origin of				
		prestalk cells	147			
	9.7	Are any prestalk genes expressed without DIF?	151			
	9.8	The history of a cell affects its fate	152			
	9.9	How are the constant ratios of prestalk and prespore cells				
		to be explained?	154			
	9.10	What is the basis of cell sorting?	156			
	9.11	Overexpression of PKA can compensate for a lack of cAMP	158			
	9.12	The cells induce several adhesion systems during formation				
		of the mound	158			
		9.12.1 Gp24	159			
		9.12.2 Gp80	159			
		9.12.3 Gp64	162			
	9.13		162			
	9.14	The genetic complexity of mound formation	163			
10	The I	he Behavior of Cells in the Slug				
	10.1	The tight aggregate elongates under control of the tip	166			
	10.2		167			
	10.3	Slugs move toward light and heat	171			
	10.4	The unanticipated complexity of cells in the slug	173			
		10.4.1 The prestalk region in the slug can be subdivided	174			
		10.4.2 Cellular traffic and cell-type conversion during slug				
		migration	177			
		10.4.3 How do prespore segments restore their severed				
		prestalk tips?	177			
	10.5	How do the cells in the slug move?	178			
	10.6	Gene regulation within the prestalk and prespore zones	180			
		10.6.1 The <i>ecmA</i> and <i>ecmB</i> promoters are highly regulated	181			
		10.6.2 Dd-STATa binds to the <i>ecmB</i> promoter inhibitory				
		sequences	182			
		10.6.3 Regulation of <i>ecmA</i>	183			
		10.6.4 The prespore gene promoters	184			
	10.7	Slugger mutants maintain their repression of culmination	185			
	10.8	1	186			
11	Culmination 18					
_	11.1	Deciding when migration has gone on long enough	188			
	11.2	Early steps in culmination	189			
	11.3	Movements at the Mexican hat stage	192			
	11.4		193			
		Cellulose synthesis and the formation of stalk	197			

CONTENTS	хi

	11.6	Death comes to the stalk cells	198
	11.7	Participation of the prespore cells in culmination	199
	11.8	The final step in spore formation is regulated exocytosis	199
	11.9	The coordination of spore and stalk formation	201
	11.10	Small peptides and the timing of encapsulation	204
	11.11	Genetic experiments suggest the source of the inducer of	
		spore encapsulation	205
	11.12	Ligands for two-component sensors	206
	11.13	The targets of PKA	208
	11.14	Culmination-defective mutants	208
12	Forma	ation and Germination of Spores	210
	12.1	The spore coat has a complex architecture	210
	12.2	The proteins of the spore coat	211
	12.3	The synthesis of cellulose in spores	215
	12.4	The formation of the spore entails changes in the cytoplasm	216
	12.5	Sorocarps contain inhibitors of germination	216
	12.6	Amoebae emerge with caution	218
	12.7	The program of spore germination	219
13	Resou	irces	223
	13.1	Books	223
	13.2	Articles for the non-scientist	224
	13.3	Films and videos	225
	13.4	The Franke bibliographic database	225
	13.5	Websites	225
Rej	^f erence.	S	227
Ind	lex		285

A Brief Introduction to Dictyostelium discoideum and its Relatives

Dictyostelium discoideum is the most studied species of the social amoebae, which are also known as the cellular slime molds. All of these organisms live in the soil and feed on bacteria, living a solitary life until the bacteria are consumed. The onset of starvation forces a major revision in the life cycle, and the amoebae respond by collecting into aggregates which transform into an organism that undergoes cell differentiation and morphogenesis. The result is a fruiting body consisting of a ball of resistant spores suspended on a stalk. D. discoideum and similar species have evolved strategies to survive in the harsh environment of the soil. A close examination of these strategies raises questions at all levels of biology: How do the amoebae sense starvation and other stresses, and how do they respond? How do they communicate with each other and how do they move? What mechanisms of signal transduction do they use, and how do those resemble the mechanisms of more complex organisms? How did the extraordinary cooperativity of development evolve? Rather than forcing the reader who has no experience with these organisms into details immediately, this chapter will provide a short glossary of terms and an overview of development, first in D. discoideum, and then in a few related species.

The developmental cycle begins when the amoebae consume all of their prey. If they do nothing to protect themselves, they will die from starvation. In response to this pressure, three distinct responses can occur – the amoebae can form microcysts, macrocysts, or fruiting bodies. The last is by far the most studied because it exhibits the fundamentals of all developing organisms. The cells signal each other to insure their correct proportion and pattern and they regulate – creating two full organisms from the two halves of a severed one. Development in *D. discoideum*, and numerous organisms like it, is composed of two phases: an aggregative period during which cells assemble in response to a

chemotactic signal, and a complex fruiting body stage in which cells differentiate and rearrange themselves to form a mass of spores supported by a stalk. The Dictyostelia have strictly separated growth and developmental stages. A cell either consumes bacteria or other nutrients and divides mitotically, or it develops in response to starvation. One of the useful consequences of this separation is that genes that are induced during development are usually not needed for mitotic growth and so they can be mutated without affecting the viability of the growing organism. Starvation induces a variety of new genes whose products are necessary for chemotaxis toward cAMP (or other chemoattractants). These will be presented in detail in future chapters, but for the moment it is sufficient to realize that cAMP is the molecule that the amoebae recognize during chemotaxis. Their ability to synthesize, release, detect, and degrade cAMP is critical for aggregation and none of these capacities exists in the growing amoebae – all are induced as the cells starve.

The amoebae are grown on lawns of bacteria or in a sterile liquid medium. To begin development in the laboratory we remove the source of nutrients and put the cells on a moist solid substratum. The substrates can be agar or filter paper, as long as it is sufficiently moist. No nutrients are provided during development – the amoebae aggregate and make their fruiting structures entirely on metabolic reserves accumulated during the trophic phase. After the washed amoebae are deposited on the substrate, there is a period of apparent inactivity as many of the genes required during growth are down-regulated, or their proteins are degraded and new genes – whose products are essential for aggregation – are induced. Gradually, after a period of hours, occasional amoebae begin to release cAMP into the population and as the macromolecules that produce, detect, and modulate the cAMP signal are made in increasing amounts, the propagation of the signal becomes stronger and stronger.

The signals that D. discoideum amoebae use are relayed, which allows the organism to collect cells from a wide area, so that an aggregate of 100,000 cells can result. The relay system is shown diagrammatically in Fig. 1.1. A single cell in a field of starving cells releases a pulse of cAMP, and other cells detect the cAMP as it binds to their cell surface receptors. First the cells change their shape and then respond in two ways – by releasing another pulse of cAMP, and by moving up the gradient of cAMP released by the first cell. Thus there is an outwardly propagated wave of cAMP and an inward movement of cells. The cAMP propagation and the cell movement happen in steps – the central cells release a pulse of cAMP about every 6 minutes and inwardly moving cells only move as long as the slope of the gradient is positive, so that when the wave decays, the amoebae stop. With dark-field illumination we can see the moving cells because they appear lighter than non-moving cells that are awaiting another wave of cAMP. Such results can be seen in Fig. 1.2. Amoebae can move into aggregation centers from aggregation territories as much as 1 cm across. In the soil, movement of cells into an aggregate would be from three dimensions and over much tougher terrain. The circular patterns of aggregation often evolve into spirals and eventually, as aggregation progresses, the concentric rings break down into long streams of cells, which still move in a

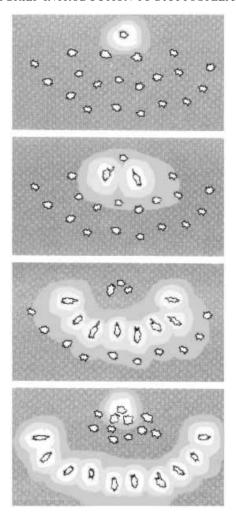


Figure 1.1 During starvation cells develop the capacity to synthesize, detect and destroy cAMP. When one cell releases a pulse of cAMP, neighboring cells detect it, move up its gradient toward higher concentrations and after a minute or so, release cAMP of their own. This attracts more outlying cells. The process is repeated about every 6 minutes under laboratory conditions. (Reprinted by permisson of American Scientist, magazine of Sigma Xi, The Scientific Research Society (Kessin and van Lookeren Campagne, 1992).)

periodic fashion into the center. As the cells move into the aggregate they become adhesive and capable of constructing a three-dimensional structure.

Once the cells have collected into a central point, a process that takes about 8 hours under most laboratory conditions, a stage is attained which is called the loose aggregate. This is shown in the scanning electron microscope figure produced by Grimson and Blanton (Fig. 1.3). The loose aggregate is relatively flat, with indistinct borders, but over the next few hours it changes to a

4 DICTYOSTELIUM

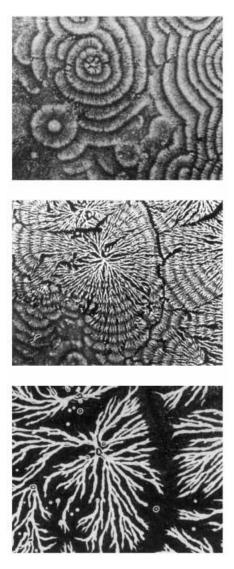


Figure 1.2 The patterns of aggregation can be seen by dark-field illumination because moving and stationary cells reflect light differently. In the region of the immobilized cells (dark bands) there is no longer a cAMP gradient, a situation that will change when the next wave of cAMP is propagated from the center. (Courtesy of Peter C. Newell, University of Oxford.)

hemispherical shape, shown as the second structure in Fig. 1.3, the tight aggregate, or mound. It is covered with a layer of mucopolysaccharide and cellulose that is called the sheath. The third structure in the figure, moving from lower right to left, has formed a critical new element, called the tip. The tip is the source of signals that organize the behavior of cells that are behind it. The tip is

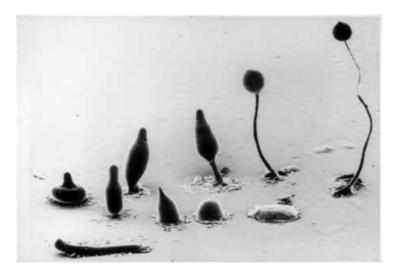


Figure 1.3 The post-aggregation stages of *Dictyostelium* development. Moving counter-clockwise, the stages encountered are the loose aggregate, the tight aggregate, the tipped aggregate, an elongated form called the finger, a slug, and then the stages of culmination leading from the Mexican hat stage to the fruiting body. (This scanning electron micrograph is by R. Lawrence Blanton and Mark Grimson, Texas Tech University.)

an important group of cells that controls development – cut it off and development stops until a new tip is formed. The tip functions like the Mangold/ Spemann organizer in vertebrates. Under the control of the tip, the aggregate elongates and makes a structure called a finger, or standing slug. At this point the organism may proceed directly to the production of fruiting bodies, or the finger may fall over and migrate in a structure called the slug or pseudoplasmodium. This is shown as the structure out of the progression in Fig. 1.3. The slug is wonderfully phototactic and is also capable of migrating up very shallow heat gradients. Once they have aggregated, Dictyostelium aggregates have many of the properties of an embryo – they have polarity, they have exquisite proportioning, they regulate, and they have an organizing center – the anterior tip.

The transition from migrating slug or standing slug to the fruiting body occurs by a process called culmination. The origins of cells that will make the prespore cells or the prestalk cells can be traced to the growing cells and depends to a certain extent, in a way we will describe later, on the position of a cell in the cell cycle when starvation is imposed. Prespore cells are those which have synthesized certain proteins in preparation for becoming spore cells, while prestalk cells are similarly identified by the expression of proteins that contribute to the stalk. As their names imply, each type is a progenitor of a terminally differentiated cell. Whether this separation into two precursor cell types occurs by a positional information mechanism or by a sorting mechanism is a question that we will defer. Neither of these cell types is fully differentiated or committed. Until very late in development, one type can convert to the other and given food, each type will revert to an amoeboid state.

By the time of aggregation, genes that are essential to the formation of the spore or the stalk have been expressed in the two precursor populations, and by the time of the tipped aggregate these two cells types have nearly sorted out, so that the cells that will make the stalk are on top and the cells that will form the spores are on the bottom. The topological problem faced by the developing cells is how to get the spores on top and the stalk on the bottom. The stalk is formed by a movement of cells from the tip through a collar at the apex of the aggregate. A special class of prestalk cells migrates through and down toward the substratum. During this period they become vacuolated as they make cellulose and die. Gradually the prespore cells are lifted up, as shown in the four last structures of Fig. 1.3. The left most structure in Fig. 1.3 is called a Mexican hat, for obvious reasons. It is the earliest of the culminating forms. The next stages (left to right) are called early to late culminants. As the spore mass is pulled up the stalk, the prespore cells undergo encapsulation, in which the contents of internal vesicles are released by exocytosis to form a layered cell wall of mucopolysaccharide and cellulose around each spore. Encapsulation spreads in a wave from the top of the developing spore mass to the bottom. How this is regulated so that it only happens at the very end of culmination will be described in a future chapter. The final structure, the fruiting body, contains a spore mass, supported by a slender stalk. The ratio of stalk to spore cells is about 1:4 and varies little between large fruiting bodies and small ones.

In *D. discoideum*, no mature stalk cells are present in the slug. Some species have a variant on this theme in which the stalk forms in the center of a migrating slug, which appears to have a rod down the middle. Such is the case with *D. mucoroides*, the first species isolated by Brefeld in 1869. As the slug migrates, it extends along the stalk which is left behind and can provide tensile strength, so that migration is in the air as well as on a surface. There are a variety of other species, some much more common than *D. discoideum*. These include *D. purpureum*, *D. giganteum*, *D. lacteum* and many others (Cavender, 1990; Hagiwara, 1989; Raper, 1984).

There are other variations – in the species *Polysphondylium violaceum* and *Polysphondylium pallidum*, culminating fruiting bodies develop delicate whorls, such that a series of secondary stalks and spore masses are produced along the main axis (Byrne and Cox, 1986; Harper, 1929). How this occurs so precisely, so that the secondary spore masses are evenly spaced, is a fundamental question of pattern formation. The various mature fruiting bodies are shown in Fig. 1.4. There is another difference among species, particularly the two species of *Polysphondylium*. Although these are all aggregative organisms, they do not all use cAMP as a chemotactic molecule. *P. violaceum* uses a dipeptide called glorin, which was identified in a tour de force of chemical analysis by Shimomura, Suthers, and Bonner (1982).

The significance of aggregation as a means of producing a mass of cells for differentiation is discussed in the next chapter, but the Dictyostelid species described above have more distant relatives (Cavender, 1990). These are

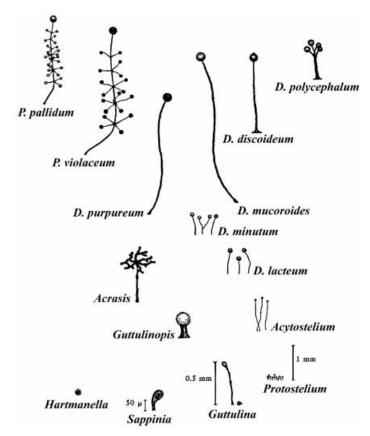


Figure 1.4 The various fruiting body structures of Dictyostelids. Note the distinction between Dictyostelium and Polysphondylium with its secondary stalks. Other much more distantly related organisms are also shown. These share an aggregative life style but may be of independent evolutionary origin. Protostelium forms a stalk and spore from a single cell (Bonner, 1967). (Reprinted by permission of Princeton University Press.)

grouped under the title of Acrasidae and Acytosteliaceae by Raper, whose 1984 book should be consulted for keys and other material for assigning unknown organisms to particular classes or species, as well as excellent descriptions (Raper, 1984). A series of aggregative organisms, all of which have an amoeboid trophic form and a fruiting structure that depends on aggregation, are known. These include species like Acrasis rosea, Copromyxa protea, and Guttulina rosea (Bonner, 1967; Raper, 1984). Most are little studied, especially from the cell biological or molecular point of view, but all share the capacity to assemble multiple cells to form a fruiting structure and all are inhabitants of the soil. Little is known about the details of their development or whether the chemotactic and developmental mechanisms they use resemble those of D. discoideum. There is no reason to believe that these organisms and D. discoi-

8 DICTYOSTELIUM

deum are monophyletic – the aggregative strategy for creating a larger organism could have evolved many times, and probably did (Blanton, 1990).

Since their discovery in 1869 by Brefeld, these organisms have been called slime molds. Phylogenetically they are not fungi, as we will learn in Chapter 3, nor are they slimy, as organisms go. For this reason, and because the title *slime mold* does not sound sufficiently elevated or convey an idea of the true evolutionary niche, many workers in the field have decided to use the designations *Dictyostelium* or social amoeba. Attempts to redirect a language are usually not successful, as the French Academy is no doubt aware, but in this book, we will use *Dictyostelium* or social amoeba.