INNER STRUCTURES OF BACTERIA

Edited by
WOUTERA VAN ITERSON

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Edited by **WOUTERA VAN ITERSON**

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SERIES EDITOR'S FOREWORD

In the 30-odd years since the electron microscope came into general use, there has been a profound change in our concepts of both the inner and outer parts of the bacterial cell. An entirely new understanding of the organization of the bacterial cell resulted from improvements in the electron microscope itself and especially from the introduction of thin sections and positive and negative staining techniques and the introduction of the entirely different freeze-etch technique, which occurred in the late 1960s and early 1970s. In addition, the development of physical fractionation techniques followed by gentler methods permitted a new approach, making it possible to relate structure to function.

Woutera van Iterson. In many papers on cytology it is the selection of the typical photograph that constitutes the prime evidence for the structure reported. To reproduce lengthy papers containing many photographs of the relevant structures would require many volumes to cover the important observed structures. Further, not all bacteria are alike and the structures found in one kind may be completely absent in another. Consequently, the selection of data (consisting almost entirely of electron micrographs) is critical and requires vast experience in the artifacts that can be produced and in the kind of information that can be derived from such micrographs. If, however, one selects only the outstanding micrographs of structures, then increased discussion is also required.

Dr. van Iterson has excerpted many more papers than is usual in a Benchmark volume and because of the nature of the subject she has included a review of an enormous amount of related literature. Indeed, she has produced a monographic discussion of each of the observed structures and has reviewed the literature associated with each throughout the volumes, covering not only cellular cytology but biochemistry and physiology as well. These volumes represent a masterpiece treatment by a recognized world authority of the subject and will stand as benchmarks for years to come.

WAYNE W. UMBREIT

PRFFACE

Bacteria have a solid or semisolid, intricate structure within which their numerous physiological functions are integrated. Structure analysis, with the electron microscope in particular, attempts to examine this structure to supply a basis for integrating biochemical data so that function-structure relationships in microorganisms become more understandable.

In this volume, *Inner Structures of Bacteria*, and its companion volume, *Outer Structures of Bacteria*, I have attempted to review the development of ideas during the 35 years that the electron microscope has been applied to structure analysis so intensely that it created the base for modern bacterial cell biology. However, while the cell's main constituent is water, in the electron microscope the specimen is placed in high vacuum. Membrane-integrated and soluble enzymes, ribosomes, and all elementary biological structures function—and virtually exist—only in water. Fixation, dehydration, infiltration, and curing of resin are parts of embedding techniques applied prior to thin sectioning of cells for examination with the electron microscope. Removal of water affects the cell structures.

The criticism of some 35 years ago that the electron microscopist is a student of artifacts is to some extent still recognized by scientists today. With the present electron microscopical point resolution of ± 0.3 nm, it still is rarely possible to reveal details smaller than 5–10 nm in biological material. The critical investigator knows that what he or she sees in an electron micrograph is not a picture of a life situation, but only an approximation, the best one can get of the true condition. This approximation should be supplemented and checked with data from other disciplines, keeping in mind *their* limitations.

Today's critical attitude toward electron microscope procedures is a challenge for many investigators to reconsider the physical base for improving preservation of fine detail, looking for new ways to overcome the actual present limitations (Sjöstrand, 1978, 1979; Kellenberger, 1979; Kellenberger et al., 1980). Therefore, this moment could well be the time of choice to review our present picture of the bacterial cell structure as deduced from direct and indirect data.

It has been said that fixation with osmium "fossilizes" cell structures, and because proteins would disintegrate into small fragments, leaving only a "skeleton" (Kellenberger, pers. commun.), we are not looking at life, but at a fossil! But has paleontology not offered us a fascinating picture of the living

forms from the past? Then let us look courageously and critically at these new experimentally produced fossils from the laboratory.

New data are appearing with overwhelming rapidity. The newest are not necessarily the truest. Couldn't it be that they contain the same percentage of error found in the older data? In the addenda found in these two volumes on bacterial anatomy, brief mention is made of some of the additional viewpoints that have appeared after completion of the original manuscripts.

Acknowledgements

I feel very grateful to a number of scientists in the Netherlands and to other scientists abroad, for having read through some of the chapters of the present two volumes on the fine structure of bacteria. I hesitate to mention them by name because of all the alterations and rewritings I did afterwards.

How can I thank Dr. H. L. van Vierssen Trip in Ottawa for patiently altering my "Netherlandiisms" and for retyping of the text? Where my ways of expression remain outlandish this must be attributable to my ecological niche on the North Sea shore.

WOUTERA VAN ITERSON

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CONTENTS BY AUTHOR

Altendorf, K. H., 457 Ball, F. L., 126 Beaman, T. C., 451 Bechtel, D. B., 332 Beutner, E. H., 52 Birch-Andersen, A., 54 Birdsell, C. C., 403 Branton, D., 466 Bulla, L. A., Jr., 332 Burdett, I. D. J., 244 Chatterjee, A. N., 406 Cheng, K.-J., 419 Clements, I., 135 Cohen-Bazire, G., 124, 129, 272 Costerton, J. W., 419 Cota-Robles, E. H., 270, 403 Davies, S. L., 197 Doi, R. H., 342 Drews, D., 66 Drews, G., 162, 173 Dugan, P. R., 199 Dunlop, W. F., 132 Engelman, D. M., 476 Fiil, A., 466 Fitz-James, P., 138 Fuller, R. C., 173 Gerhardt, P., 451 Giesbrecht, P., 66 Gilpin, R. W., 406 Golecki, J. R., 173 Graham, L. B., 195 Grilione, P. L., 365 Groen, J. B., 265 Hannay, C. L., 138

Hillier, J., 52

Hitchins, A. D., 328 Hitchins, V. M., 357 Holt, S. C., 164 Ingram, J. M., 419 Kawata, T., 276 Kellenberger, E., 54 Kleinschmidt, A., 62. Kline, B. W., 126 Kunisawa, R., 124, 129, 272 Laishley, E. J., 135 Lake, J. A., 96 Lang, D., 62 Maalge, O., 54 MacAlister, T. J., 135 Mandel; M., 191 Maniloff, J., 91 Marr, A. G., 164 Mason, D. J., 51 Morowitz, H. J., 410, 441 Mudd, S., 52, 276 Oelze, J., 162 Oppenheim, J. D., 452 Pangborn, I., 365 Pankratz, H. S., 451 Payne, J. I., 276 Peterson, J. E., 139 Pfennig, N., 124, 129 Poindexter, J. S., 272 Powelson, D. M., 51 Razin, S., 441, 443 Reichenbach, H., 361 Remsen, C. C., 57, 167, 171, 195 Richards, F. M., 476 Robards, A. W., 132 Robinow, C. F., 48, 50

Contents by Author

Rodwell, A. W., 139 Rodwell, E. S., 139 Rogers, H. J., 244 Rucinsky, T. E., 270 Ryter, A., 54, 254 Sadoff, H. L., 357 Sall, T., 276 Salton, M. R. J., 452 Santo, L. Y., 342 Shively, J. M., 126 Sjöstrand, F. S., 461 Slepecky, R. A., 328 Sleytr, U. B., 60 Smith,\A. G., 52 Staehelin, L. A., 173, 457 Stanier, R. Y., 16 Takagi, A., 276 Terry, T. M., 441 Thornley, M. J., 60 Tourtellotte, M. E., 410 Trüper, H. G., 167, 171

Valois, F. W., 195 van Deenen, L. L. M., 471 van Iterson, W., 2, 17, 235, 265 van Niel, C. B., 16 Verkleij, A. J., 47.1 Ververgaert, P. H. J., 471 Voelz, H., 361 Wake, R. G., 64 Walker, P. D., 303 Wallace, B. A., 476 Waterbury, J. B., 167 Watson, S. W., 167, 191, 195 Weaver, T. L., 199 Weibull, C., 395 Whittenbury, R., 197 Young, C., 135 Young, F. E., 406 Zahn, R. K., 62

CONTENTS

Series Editor's Forewor Preface Contents by Author	rd		xi xiii xv
Contents by Author	PART I: BACTERI	AL CYTOLOGY	XV
Editor's Comments on	Papers 1, 2, and 3		2
1 VAN ITER	RSON, W.: Much Within Lit W. van Iterson, Drukkerij V/H C. de Boer Jr. N.V.,		15
2 STANIER,	, R. Y., and C. B. VAN NIEL: T Arch. Mikrobiol. 4		16
	organisms ;	ructures in Micro- /III, University of Toronto Press, o2, Plate 1	17
	PART II: THE	NUCLEOID	
Editor's Comments on	Papers 4 Through 13	Committee Transport	20
	Bacteria	s and Cell Structure of Rod-Shaped Dubos, Harvard University Press, , 1945, Plate 1	48
5 ROBINON	W, C. F.: Morphology of th Br. Med. Bull. 18:3	e Bacterial Nucleus 2, Plate 1 (1962)	50
6 MASON,	D. J., and D. M. POWELSON: Live Bacteria by a New Te J. Bacteriol. 71		51
7 MUDD, S	Light Microscopic Studie	d E. H. BEUTNER: Electron and s of Bacterial Nuclei. III. The adowed Cells of Escherichia coli 35, 638 (1950)	52
8 RYTER, A	Electron Microscope Stud	dy of Deoxyribonucleic Acid cleoids of Actively Growing	54

Contents

	9	REMSEN, C. C.: Fine Structure of the Mesosome and Nucleoid in Frozen-Etched Bacillus subtilis Arch. Mikrobiol. 61:40, 44-45 (1968)	57
	10	SLEYTR, U. B., and M. J. THORNLEY: Freeze-Etching of the Cell Envelope of an Acinetobacter Species Which Carries a Regular Array of Surface Subunits J. Bacteriol. 116:1394 (1973)	60
	11	KLEINSCHMIDT, A., D. LANG, and R. K. ZAHN: On the Intracellular Configuration of Bacterial DNA Z. Naturforsch. 16B:730, Figure 7 (1961)	62
	12	WAKE, R. G.: Visualization of Reinitiated Chromosomes in <i>Bacillus</i> subtilis J. Mol. Biol. 68: 501, Plate II (1972)	64
	13	GIESBRECHT, P., and D. DREWS: The Nuclear Structures of Bacteria and Their Relation to Those of the Mesokaryotes Translated from Die Zelle, Struktur und Funktion, 3rd ed., H. Metzner, ed., Wissenschaftliche, Stuttgart, 1981, pp. 542-597	66
31		PART III. THE CYTORIAGAA DIROCOARES	
		PART III: THE CYTOPLASM: RIBOSOMES	
Edito		nments on Papers 14 and 15	72
	14	MANILOFF, J.: Analysis of the Helical Ribosome Structures of Mycoplasma gallisepticum Natl. Acad. Sci. (USA) Proc. 68:43-47 (1971)	91
	15	copy of Escherichia coli Small Subunits, Large Subunits and Monomeric Ribosomes 1. Mol. Biol. 105:150, 155 (1976)	96
		PART IV: CYTOPLASMIC INCLUSIONS	
Edito	r's Con	nments on Papers 16 Through 22	100
	16	COHEN-BAZIRE, G., N. PFENNIG, and R. KUNISAWA: The Fine Structure of Green Bacteria J. Cell Biol. 22:207, 208, 209 (1964)	124
	17	SHIVELY, J. M., F. L. BALL, and B. W KLINE: Electron Microscopy of the Carboxysomes (Polyhedral Bodies) of <i>Thiobacillus</i> neapolitanus J. Bacteriol. 116:1405, 1406, 1410 (1973)	126
	18	COHEN-BAZIRE, G., R. KUNISAWA, and N. PFENNIG: Comparative Study of the Structure of Gas Vacuoles J. Bacteriol. 100:1049, 1050, 1059, 1060, 1061 (1969)	129
	19	DUNLOP, W. F., and A. W. ROBARDS: Ultrastructural Study of Poly-β-Hydroxybutyrate Granules from Bacillus cereus J. Bacteriol. 114:1271, 1273, 1280 (1973)	132
14E .	20	LAISHLEY, E. J., T. J. MacALISTER, I. CLEMENTS, and C. YOUNG: Isolation and Morphology of Native Intracellular Polyglucose Granules from Clostridium pasteurianum Can. J. Microbiol. 19:991, Plates III and V (1973)	135

	21	HANNAY, C. L., and P. FITZ-JAMES: The Protein Crystals of Bacillus	
		thuringiensis Berliner Can. J. Microbiol. 1:694, 706, Plate I (1955)	138
	22	RODWELL, A. W., J. E. PETERSON, and E. S. RODWELL: Striated Fibers of the <i>rho</i> Form of <i>Mycoplasma</i> : In Vitro Reassembly, Composition, and Structure J. Bacteriol. 122:1216, 1227, 1228-1229 (1975)	139
		PART V: CYTOMEMBRANE STRUCTURES IN PHOTOTROPHS	
Edito	r's Con	nments on Papers 23 Through 27	144
	23	OELZE, J., and G. DREWS: Membranes of Photosynthetic Bacteria Biochim. Biophys. Acta 265:209, 211, 235 (1972)	162
	24	HOLT, S. C., and A. G. MARR: Location of Chlorophyll in Rhodospirillum rubrum J. Bacteriol. 89:1402, 1407, 1411 (1965)	164
	25	REMSEN, C. C., S. W. WATSON, J. B. WATERBURY, and H. G. TRÜPER: Fine Structure of Ectothiorhodospira mobilis Pelsh J. Bacteriol. 95:2374, 237 , 2383, 2390 (1968)	167
	26	REMSEN, C. C., and H. G. TRÜPER: The Fine Structure of Chromatium buderi	171
		Arch. Mikrobiol. 90: 269, 271, (1973)	
	27	Visualization of the Supramolecular Architecture of Chlorosomes (Chlorobium type Vesicles) in Freeze-Fractured Cells of Chloroflexus aurantiacus Arch. Mikrobiol. 119:269, 271, 272, 276 (1978)	173
	PA	RT VI: CYTOMEMBRANE STRUCTURES IN CHEMOLITHOTROPHS, METHYLOTROPHS, AND IN A FEW ORGANOTROPHS	
Edito	r's Con	nments on Papers 28 Through 31	179
	28	WATSON, S. W., and M. MANDEL: Comparison of the Morphology and Deoxyribonucleic Acid Composition of 27 Strains of Nitrifying Bacteria J. Bacteriol. 107:563, 566-568, 569 (1971)	191
	29	WATSON, S. W., L. B. GRAHAM, C. C. REMSEN, and F. W. VALOIS: A Lobular, Ammonia-Oxidizing Bacterium, Nitrosolobus multiformis Nov. Gen. Nov. Sp. Arch. Mikrobiol. 76:183, 189 (1971)	195
	30	DAVIES, S. L., and R. WHITTENBURY: Fine Structure of Methane and Other Hydrocarbon-utilizing Bacteria J. Gen. Microbiol. 61:227, 231, Plate 1 (1970)	197
	31	WEAVER, T. L., and P. R. DUGAN: Ultrastructure of Methylosinus trichosporium as Revealed by Freeze Etching J. Bacteriol. 121:704, 705, 707, 708, 709 (1975)	199

PART VII: MESOSOMES

Editor	's Con	nments on Papers 32 Through 38		204
	32	van Iterson, W.: Symposium on the Fine Structure and Replication of Bacteria and Their Parts. II. Bacterial Cytoplasm Bacteriol. Rev. 29:299, 302, 305, 306, 307, 309, 310, 311 (1965)		235
	33	BURDETT, I. D. J., and H. J. ROGERS: The Structure and Development of Mesosomes Studied in <i>Bacillus licheniformis</i> Strain 6346 <i>J. Ultrastruct. Res.</i> 38:113-114, 115, 116, 117, 118, 119, 131-133 (1972)		244
	34	RYTER, A.: Association of the Nucleus and the Membrane of Bacteria: A Morphological Study Bacteriol. Rev. 32:39-40, 43-45, 47-48, 50, 51-52, 53, 54 (1968)		254
	35	VAN ITERSON, W., and J. B. GROEN: DNA-Membrane Complexes of Bacillus subtilis: Contact of Mesosomal Vesicles and Nuclear Fibrils J. Ceil Biol. 49:553–554, 555, 556, 557, 558, 559, 560 (1971)		265
	36	RUCINSKY, T. E., and E. H. COTA-ROBLES: Mesosome Structure in Chromobacterium violaceum J. Bacteriol. 118:717, 721, 724 (1974)		270
	37	COHEN-BAZIRE, G., R. KUNISAWA, and J. S. POINDEXTER: The Internal Membrane of Caulobacter crescentus J. Gen. Microbiol. 42:301, 307, 308, Plate 2, Plate 5, Plate 6 (1966)		272
	38	MUDD, S., T. KAWATA, J. I. PAYNE, T. SALL, and A. TAKAGI: Plasma Membranes and Mitochondrial Equivalents as Functionally Coordinated Structures Nature 189:79–80 (1961)	,	276
		PART VIII: SPORES		
Editor	r's Con	nments on Papers 39 Through 42		280
	39	WALKER, P. D.: Cytology of Spore Formation and Germination <i>J. Appl. Bacteriol.</i> 33: 1–12 (1970)		303
	40	HITCHINS, A. D., and R. A. SLEPECKY: Bacterial Sporulation as a Modified Procaryotic Cell Division Nature 233:804–807 (1969)		328
	41	BECHTEL, D. B., and L. A. BULLA, JR.: Electron Microscope Study of Sporulation and Parasporal Crystal Formation in Bacillus thuringiensis J. Bacteriol. 127:1472-1481 (1976)		332
	42	SANTO, L. Y., and R. H. DOI: Ultrastructural Analysis During Germination and Outgrowth of <i>Bacillus subtilis</i> Spores J. Bacteriol. 120 :475-481 (1974)		342
		PART IX: MICROCYSTS AND MYXOSPORES		
Editor	's Con	nments on Papers 43, 44, and 45		350
	43	HITCHINS, V. M., and H. L. SADOFF: Morphogenesis of Cysts in Azotobacter vinelandii J. Bacteriol. 104:492, 493, 494-497, and 498 (1970)		357

			Contents
	44	VOELZ, H., and H. REICHENBACH: Fine Structure of Fruiting Bodies of Stigmatella aurantiaca (Myxobacterales) J. Bacteriol. 99:856, 857, 858, 862, 866 (1969)	361
	45	GRILIONE, P. L., and J. PANGBORN: Scanning Electron Microscopy of Fruiting Body Formation by Myxobacteria J. Bacteriol. 124:1558, 1560-1561, 1565 (1975)	365
		PART X: THE CYTOPLASMIC MEMBRANE IN SITU	
Edito	r's Con	nments on Papers 46 Through 50	370
	46	WEIBULL, C.: The Isolation of Protoplasts from <i>Bacillus megaterium</i> by Controlled Treatment with Lysozyme <i>J. Bacteriol.</i> 66: 688–695 (1953)	395
	47	birdsell, C. C., and E. H. COTA-ROBLES: Production and Ultrastructure of Lysozyme and Ethylenediaminetetraacetate-Lysozyme Spheroplasts of Escherichia coli J. Bacteriol. 93:427, 435, 436, 437 (1967)	403
	48	GILPIN, R. W., F. E. YOUNG, and A. N. CHATTERJEE: Characterization of a Stable L-Form of <i>Bacillus subtilis</i> 168 J. Bacteriol. 113:486, 491, 492, 495, 498, 499 (1973)	406
	49	MOROWITZ, H. J., and M. E. TOURTELLOTTE: The Smallest Living Cells Sci. Am. 206:117-124, 126 (1962)	410
	50	COSTERTON, J. W., J. M. INGRAM, and KJ. CHENG: Structure and Function of the Cell Envelope of Gram-Negative Bacteria Bacteriol. Rev. 38:87, 89 (1974)	419
		PART XI: THE ISOLATED CYTOPLASMIC MEMBRANE	
Edito	r's Con	nments on Papers 51 Through 59	424
	51	RAZIN, S., J. J. MOROWITZ, and T. M. TERRY: Membrane Subunits of Mycoplasma laidlawii and Their Assembly to Membranelike Structures	441
		Natl. Acad. Sci. (USA) Proc. 54: 219, 222 (1965)	
	52	RAZIN, S.: Reconstitution of Biological Membranes <i>Biochim. Biophys. Acta</i> 265: 241, 243, 260-266, 293, 294 (1972)	443
	53	BEAMAN, T. C., H. S. PANKRATZ, and P. GERHARDT: Chemical Composition and Ultrastructure of Native and Reaggregated Membranes from Protoplasts of Bacillus cereus J. Bacteriol. 117:1335, 1338 (1974)	451
	54	OPPENHEIM, J. D., and M. R. J. SALTON: Localization and Distribution of Micrococcus lysodeikticus Membrane ATPase Determined by Ferritin Labeling Biochim. Biophys. Acta 298:297, 305, 307, 309, 312, 314, 318 (1973)	452
	55	ALTENDORF, K. H., and L. A. STAEHELIN: Orientation of Membrane Vesicles from <i>Escherichia coli</i> as Detected by Freeze-Cleave Electron Microscopy J. Bacteriol. 117:888-889, 890, 894, 898-899 (1974)	457
	56	SJÖSTRAND, F. S.: The Problems of Preserving Molecular Structure of Cellular Components in Connection with Electron	

Contents

		Microscopic Analysis J. Ultrastruct. Res. 55:271, 277-280 (1976)	461
	57	FIIL, A., and D. BRANTON: Changes in the Plasma Membrane of Escherichia coli During Magnesium Starvation J. Bacteriol. 98:1320, 1322, 1323, 1326, 1327 (1969)	466
	58	VERKLEIJ, A. J., P. H. J. VERVERGAERT, L. L. M. VAN DEENEN, and P. F. ELBERS: Phase Transitions of Phospholipid Bilayers and Membranes of Acholeplasma laidlawii B Visualized by Freeze Fracturing Electron Microscopy Biochim. Biophys. Acta 288:326, 328, 329-331, 332 (1972)	471
	59	wallace, B. A., F. M. RICHARDS, and D. M. ENGELMAN: The Influence of Lipid State on the Planar Distribution of Membrane Proteins in Acholeplasma laidlawii J. Mol. Biol. 107:255, 266 (1976)	476
Autho Subje		tion Index I - I are the song prescriptions of and a reverse	479 491
About	the E	ditor	501

Part I

BACTERIAL CYTOLOGY: AN INTRODUCTION

Editor's Comments on Papers 1, 2, and 3

- 1 VAN ITERSON Excerpt from Much Within Little
- 2 STANIER and VAN NIEL Excerpt from The Concept of a Bacterium
- 3 VAN ITERSON Excerpt from Membranous Structures in Micro-organisms

During the past thirty years, bacterial cytology has seen a period of rapid development with increasing opportunity to analyze more minute detail. The discipline of bacterial cytology (bacterial anatomy) continues its expansion in the direction of molecular cytology, a subdivision of submicroscopic morphology (see Paper 1).

All physiological processes, studied in biochemistry and its affiliated discipline of molecular biology are, in their final analyses, based on anatomical structure. The processes and reactions studied in living microorganisms may involve components that are integrated in elementary biological structures. When the reactions take place in a water-dissolved state, they may involve endproducts that are determined by the lipoprotein membrane that surrounds the cell or that partitions the cell into membrane-covered parts (organelles). The structures involved in these physiological processes and reactions are so infinitesimally small that it was impossible to think of them realistically prior to visualization in the electron microscope.

It is possible, as has been done in these volumes, to subdivide bacterial morphology into a number of sections, each dealing with a separate structural part, and to present more detailed knowledge in each section. In the end, this should all merge into "The Concept of a Bacterium". This mental integrating activity has graciously been left to the reader.

"The Concept of a Bacterium," a well-known paper by Stanier and van Niel (Paper 2), replaced the old division of the living world into animal and plant kingdoms, to which Haeckel (1866) had added the kingdom of the protists. It is now commonly accepted that the living world may be divided into eukaryotes and prokaryotes, based

on fundamental structural differences between eukaryotic and prokaryotic cell types. The terms eukaryotes (*Eucaryotae*) and prokaryotes (*Procaryotae*) introduced by Dougherty (1957) have proved to represent a sharp demarcation, with few vague intermediate forms.

The only organisms with prokaryotic features are the bacteria and the blue-green algae (*Cyanophyceae*), presently called blue-green bacteria (Cyanobacteria) (see Buchanan and Gibbons, 1974; Stanier et al., 1978). *Inner Structures of Bacteria* and *Outer Structures of Bacteria* deal with bacteria in the original sense; the blue-greens are not discussed in either volume.

The triumph of thirty years of ultrastructural research is our present awareness of a basic unity in cell structure, five decades after Kluyver and Donker (1926) advocated unity in biochemisty. In eukaryotic and prokaryotic cells, analogous functions take place on analogous elementary structures, on fibrillar macromolecules such as the nucleic acids or the flagellar proteins, and on membranes and ribosomes. But the analogies are accompanied by constant fundamental differences in ultrastructural anatomy (see Stanier and van Niel, Paper 2; van Iterson, Paper 3).

SITE OF RESIDENCE OF GENETIC INFORMATION

The activities of the cell are directed by genetic information, which resides in its deoxyribonucleic acid (DNA).

In the eukaryotic cell, the genetic information is primarily contained in a membrane-covered cell nucleus; it also is contained in smaller cell organelles.—the mitochondria and the plastids (chloroplasts). In the nucleus, the genetic information is spread over a large number of highly organized elements called chromosomes, which consist of DNA, histones, and nonhistone proteins. The chromosomes replicate and divide by mitosis, which involves a spindle composed of a number of microtubules. During mitotic division, the nuclear envelope (membrane) disappears in most types of cell, to be formed again after the two daughter sets of chromosomes have parted. The nuclear division is followed by the cell division in a plane coinciding with the spindle's equatorial plane. The DNA in mitochondria and plastids, on the other hand, resembles bacterial DNA in that it is not organized in a composite chromosome. It is more or less a "naked" molecule, usually in a closed loop structure, shorter than bacterial DNA and dividing by an amitotic process.

In the prokaryotic cell (bacteria and blue-greens), the site of genetic information is mainly in a small number of nuclear equivalents—the nucleoids—devoid of a nuclear membrane. The