

CELL BIOLOGY

BASICS TO BREAKTHROUGHS

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Cell Biology: Basics to Breakthroughs

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CONTENTS

FOREWORD	i
PREFACE	ii
ACKNOWLEDGEMENTS	iii
LIST OF CONTRIBUTORS	iv
CHAPTER 1 FOUNDATIONS OF LIFE: CELLS AND ORIGIN	1
<i>Meghana A. Shakthi, Harin N. Ganesh, R. Kirubakaran and K.N. Aruljothi</i>	
INTRODUCTION	1
Theory of Spontaneous Generation	1
Abiogenesis for the Origin of Life	2
Cell and its Contents	3
<i>Cell Organelles</i>	5
Organelle Biogenesis and the Endosymbiotic Theory	6
<i>Chemical Composition of Cell</i>	7
<i>Cytoplasmic Composition of the Cell</i>	8
Structural and Functional Aspects of the Cell	8
<i>Cell Shape and How it Impacts Function</i>	8
<i>Cell Size Variation</i>	9
Integration of Cells into Tissues	10
Extracellular Matrix and Connective Tissues	11
Plant and Animal Cells	12
<i>Plant Cell</i>	12
<i>Evolution of Photosynthesis</i>	14
<i>Animal Cell</i>	14
Cells Under the Microscope	15
CONCLUSION	16
ACKNOWLEDGEMENTS	16
REFERENCE	16
CHAPTER 2 PROKARYOTES VS. EUKARYOTES: COMPARATIVE STRUCTURAL AND FUNCTIONAL INSIGHTS	20
<i>Kruthika Prakash, Sayantani Chattopadhyay, T. Sasitharan, Soumyadeep Maiti, Sanjana Dhayalan, K. Kumaran, R. Kirubakaran and K.N. Aruljothi</i>	
INTRODUCTION TO PROKARYOTES	21
Prokaryotic Diversity	22
Classification of Prokaryotes	22
Morphological Diversity	22
Metabolic Diversity	23
Ecological Diversity	23
Cell Envelope	24
Cytoplasm and External Structures	24
Prokaryotic Genetics	25
Genome Structure	25
Gene Transfer Mechanisms	25
Mutation and Adaptation	25
Prokaryotes and Human Health	26
Introduction to Eukaryotes	26
Characteristics of Eukaryotes	26

WHITTAKER CLASSIFICATION - A BREAKTHROUGH IN THE STUDY OF EUKARYOTE ANATOMY	27
DEVELOPMENT OF MULTICELLULAR ORGANISMS	28
Biotechnological Applications of Eukaryotes	30
CONCLUSION	30
REFERENCES	31
CHAPTER 3 NUCLEUS: THE CONTROL CENTRE OF THE CELL	34
<i>K. Kumaran, Harin N. Ganesh and K.N. Aruljothi</i>	
INTRODUCTION	34
History	35
STRUCTURE AND FUNCTION	36
Structure of Nucleus in Eukaryotes	36
Function of Nucleus in Eukaryotes	38
Structure of Genetic Material in Prokaryotes	38
Function of Genetic Material in Prokaryotes	39
DISORDERS ASSOCIATED WITH NUCLEUS	39
RECENT RESEARCH IN THE NUCLEUS	41
Therapeutics	42
CONCLUSION	43
ACKNOWLEDGEMENTS	44
REFERENCES	44
CHAPTER 4 ENDOPLASMIC RETICULUM: THE CELLULAR FACTORY	47
<i>Shanmuga Priya, K. Kumaran, Sanjana Dhayalan, Krishnan Anand and K.N. Aruljothi</i>	
INTRODUCTION	47
FUNCTIONS OF RER	48
The Process of Protein Folding Associated with Molecular Chaperones	49
<i>ER as the Quality Control Center</i>	50
<i>ER Stress</i>	50
Diabetes Type II and ER Stress in Autophagy	52
Therapeutics	53
<i>ER Stress – Therapeutics and Curative Treatment for Acute Spinal Cord Injury</i>	53
Therapeutic for Ulcerative Colitis	54
SMOOTH ENDOPLASMIC RETICULUM	55
Evolutionary Aspects of SER	56
<i>Calcium Storage in Smooth Endoplasmic Reticulum</i>	56
The Mitochondrial-Associated Endoplasmic Reticulum Membrane (MAM)	57
<i>Understanding the Role of MAM in Cardiovascular Disease</i>	57
RECENT TRENDS IN ER RESEARCH	58
CONCLUSION	59
ACKNOWLEDGEMENTS	59
REFERENCES	59
CHAPTER 5 GOLGI APPARATUS: SHAPING AND SHIPPING CELLULAR PROTEINS	61
<i>S. Sahasra, Sanjana Dhayalan, K. Kumaran and K.N. Aruljothi</i>	
INTRODUCTION	61
STRUCTURE OF GOLGI APPARATUS	62
CIS Face	62
MEDIAL Face	63
TRANS Face	63

FUNCTION OF THE GOLGI APPARATUS	64
Protein Glycosylation	64
Lipid and Polysaccharide Metabolism	67
Protein Sorting and Export	68
GOLGI APPARATUS DYSFUNCTION AND ASSOCIATED DISEASES	68
Neurodegenerative Diseases	68
Infectious Disease	69
Cancer	69
Genetic Mutation	69
THERAPEUTICS	70
Gaucher Disease (GD)	70
Parkinson's Disease	70
<i>GTPase Rab1</i>	71
<i>LRRK2 Mutations</i>	71
<i>Rab8 and Rab10</i>	71
<i>PLA2G6 Mutations</i>	71
<i>α-Synuclein Aggregates</i>	71
Cancer Therapy	72
<i>Inhibitors of Peripheral Golgi-Associated Proteins - TAS-116</i>	72
<i>Compounds Affecting GA Protein Localization-2-(Substituted Phenyl)-Benzimidazole (2-PB)</i>	72
<i>Nanodelivery Approaches</i>	72
RECENT TRENDS	73
COVID-19	73
Golgi Dynamics and Regulation	73
GOLGI AND DISEASE	73
Golgi and Cell Signalling	73
Golgi and Infectious Diseases	74
CONCLUSION	74
ACKNOWLEDGEMENTS	74
REFERENCES	74
CHAPTER 6 RIBOSOMES: ENGINES OF PROTEIN SYNTHESIS	77
<i>Sanjana Dhayalan, Shalini Roy, K. Kumaran and K.N. Aruljothi</i>	
INTRODUCTION	77
Ribosome in Prokaryotes and Eukaryotes	78
Origin and Evolution of the Ribosome	79
Ribosomes Fingerprint	81
STRUCTURE OF RIBOSOME	82
Composition of the Ribosome	84
FUNCTION OF RIBOSOME	85
Ribosome Biogenesis (Ribi)	86
<i>Ribi in Eukaryotic Cells</i>	86
<i>Ribi in Prokaryotic Cells</i>	86
Role of the Ribosome in Translation – Initiation	87
Role of the ribosome in Translation - Elongation	88
Role of the Ribosome in Translation – Termination	89
Ribosome Recycling	90
Regulation of Termination	91
DYSFUNCTION AND DISEASE OF THE RIBOSOME	91
	95

<i>RP Mutations and Cancer</i>	95
<i>Onco-Ribosomes and Cancer Development</i>	95
DIAGNOSTIC AND THERAPEUTIC PROPERTY OF RIBOSOMES	96
RP as Anti-microbial	96
Therapeutic Targeting of Ribi and mTORC1 in Cancer	97
RECENT TRENDS	97
Advancements in Ribosome Profiling and Cancer	97
Ribosome Profiling Reveals Non-Canonical ORF Translation in High-Risk Childhood Medulloblastoma	98
CONCLUSION	99
ACKNOWLEDGEMENTS	99
REFERENCES	99
CHAPTER 7 LYSOSOMES: THE CELL'S DIGESTIVE SYSTEM	102
<i>B. Surya, Faizaan Khan, Abhinav Roy, Sanjana Dhayalan, K. Kumaran, Krishnan Anand and K.N. Aruljothi</i>	
INTRODUCTION	102
Discovery of Lysosomes	103
Evolution of Lysosomes	104
<i>Early Endosomes</i>	105
<i>Recycling Endosomes</i>	105
<i>Late Endosomes</i>	105
FUNCTION	106
Autophagy (Process of Self-destruction)	106
Phagocytosis [Function of Specialized Cells (macrophages, Neutrophils)]	106
Lysosomal Membrane Proteins	107
Lysosome Reformation	108
Endocytic Lysosome Reformation (ELR)	109
Autophagic Lysosome Reformation (ALR)	109
Phagocytic Lysosome Reformation (PLR)	110
Key Players of Membrane Fusion	111
LYSOSOMAL DYSFUNCTION	114
Lysosome Storage Disorders (LSD)	115
<i>Parkinson's Disease</i>	116
<i>Muscular Dystrophy</i>	118
FUTURE DIRECTIONS FOR LYSOSOME RESEARCH	119
CONCLUSION	120
ACKNOWLEDGEMENTS	120
REFERENCES	120
CHAPTER 8 VACUOLES: THE STORAGE VAULTS OF THE CELL	122
<i>S. Sahasra, S. Aswini, K. Kumaran, Sanjana Dhayalan and K.N. Aruljothi</i>	
INTRODUCTION	122
Types of Vacuoles	123
<i>Plant Vacuoles</i>	123
<i>Animal Vacuoles</i>	124
<i>Fungal Vacuoles</i>	124
Differences from Plant and Animal Vacuoles	125
<i>Protist Vacuole</i>	125
Structure	126
Tonoplast	126
Morphological Changes	127

Dynamic Structures	127
Components	128
Inclusion Bodies in the Vacuole	128
<i>Pigments</i>	129
<i>Vacuoles vs. Vesicles</i>	129
Structure	129
Function	129
FUNCTION OF VACUOLES	130
Vacuolar Membrane-collapse System	131
<i>Vacuole-plasma Membrane-fusion System</i>	131
Autophagosome-lysosome Fusion	132
<i>Formation of Autolysosomes</i>	132
DIAGNOSTIC AND THERAPEUTIC APPLICATION OF VACUOLES	134
RECENT TRENDS	134
CONCLUSION	135
ACKNOWLEDGEMENTS	135
REFERENCES	136
CHAPTER 9 PLASMA MEMBRANE: GATEWAY AND SENTINEL OF CELLULAR EXCHANGE	138
<i>S. Aswini, K. Kumaran, Sanjana Dhayalan, Prakash Gangadaran and K.N. Aruljothi</i>	
INTRODUCTION	139
STRUCTURE	139
Gorter and Grendel Membrane Theory (1920)	139
Pauic-molecular Theory	140
Fluid Mosaic Model (1972)	140
Peripheral Proteins (Extrinsic Proteins)	141
<i>Integral Proteins (Intrinsic Proteins)</i>	141
Handerson and Unwin's Membrane Theory	142
Evolutionary Aspects of Plasma Membrane	143
FUNCTIONS OF PLASMA MEMBRANE	143
Acting as a Physical Barrier	143
Selective Permeability	143
Endocytosis by the Plasma Membrane	143
Exocytosis	144
Facilitating Communication and Signalling among Cells	144
Providing Shape to the Cytoskeleton and Maintaining Cell Potential	144
Membrane Transport Mechanism	145
<i>Fick's First Law</i>	145
<i>Osmosis</i>	146
<i>Passive Diffusion</i>	146
<i>Facilitated Diffusion</i>	147
<i>Glucose Transporter</i>	148
Potassium Channels	149
<i>Calcium-activated Potassium Channel</i>	149
<i>Inwardly Rectifying Potassium Channel</i>	149
<i>Tandem Pore Domain Potassium Channel</i>	149
<i>Voltage-gated Potassium Channel</i>	150
Sodium Channel	150
Aquaporin	151

<i>Active Transport</i>	151
<i>Group Translocation</i>	151
ABNORMALITIES IN PLASMA MEMBRANE	153
Hypertension	153
Sphingolipids-related Disorders	153
<i>Basis for Sphingolipidoses-neuronal Vulnerability</i>	154
Neurotrophic Signalling	154
Insulin Signalling	155
Impacts of Sphingolipid Accumulation on the use of Cellular Energy	155
DIAGNOSTIC AND THERAPEUTIC ROLE OF PLASMA MEMBRANE	156
RECENT TRENDS	156
CAR-T Therapy	156
Liposomal Drug Delivery Systems	157
CONCLUSION	158
ACKNOWLEDGEMENTS	158
REFERENCES	158
CHAPTER 10 MITOCHONDRIA AND CHLOROPLASTS: EVOLUTIONARY ENGINES OF THE CELL	161
<i>.K. Kumaran, Keerthana Ramesh, Sanjana Dhayalan, Gargii Chatterjee and K.N Aruljothi</i>	
MITOCHONDRIA	162
Introduction	162
<i>Origin and Evolution of Mitochondria</i>	163
<i>Mitochondrial Inheritance in Eukaryotes</i>	163
Structure of Mitochondria	164
Function of Mitochondria	166
<i>Electron Transport Chain and Oxidative Phosphorylation</i>	166
<i>Mitochondria-associated Membranes</i>	168
Mitochondrial DNA Mutations	168
<i>Role of Mitochondria in Cancer and Aging</i>	169
Recent Trends in the Biology of Mitochondria	170
CHLOROPLAST	173
Introduction	173
<i>Origin and Evolution of Chloroplasts</i>	173
Structure of Chloroplast	174
<i>Dynamic Thylakoid Architecture</i>	175
<i>Structural Plasticity</i>	175
<i>Light Intensity and Thylakoid Structure</i>	175
<i>Light and Dark Reactions</i>	176
Recent Trends in Malfunctions in Chloroplasts	177
<i>Autophagy and Chloroplast Degradation</i>	177
<i>Sensitivity to Thermal Stress</i>	177
<i>Production of Reactive Oxygen Species (ROS)</i>	178
<i>Impact on Mitochondrial Function</i>	178
CONCLUSION	178
ACKNOWLEDGEMENTS	179
REFERENCES	179
CHAPTER 11 CYTOSKELETON: THE CELL'S BACKBONE AND HIGHWAY	182
<i>Danyal Reyaz, Sparshika Mishra, Ranjini Sengupta, T. Sasitharan, S. Gnanavel and K.N. Aruljothi</i>	

INTRODUCTION	182
Types of Cytoskeletal Fibers	183
MICROFILAMENTS	184
Structure	184
Organisation	185
Treadmilling	185
Organization of Actin Filaments	185
Functions	186
<i>Protrusions of the Cell Surface</i>	186
<i>Muscle Contraction</i>	186
<i>Contractile Assemblies of Actin and Myosin in Non-muscle Cells</i>	188
Cytokinesis	188
<i>Locomotion</i>	189
INTERMEDIATE FILAMENTS	189
Intermediate Filaments in the Cell	189
Composition	190
Types of Intermediate Filaments	191
<i>TYPE I: Acidic Keratins</i>	191
<i>TYPE II: Basic Keratins</i>	191
<i>TYPE V: Nuclear Lamins</i>	194
Structure And Assembly of Intermediate Filaments	194
<i>Intermediate Filament: Organization And Intracellular Localization</i>	195
<i>Intermediate Filaments and Cell Signalling</i>	196
MICROTUBULES	196
Microtubules Assembly	197
<i>Role of MAPs in the Organization of Microtubules in Cells</i>	197
<i>Microtubule Motors</i>	198
<i>Cargo Transport by Microtubules</i>	199
<i>Diseases Associated with Microtubules</i>	199
CONCLUSION	201
ACKNOWLEDGEMENTS	201
REFERENCES	201

CHAPTER 12 SIGNAL TRANSDUCTION PATHWAYS ORCHESTRATE CELLULAR COMMUNICATION: A NARRATIVE REVIEW	206
<i>Kruthika Prakash, Raksa Arun, Srisri Satishkartik, Sayantani Chattopadhyay, Mashira Rahman, Prakash Gangadaran and K.N. Aruljothi</i>	
INTRODUCTION	206
TYPES OF CELL SIGNALING	207
Autocrine Signaling	207
Paracrine Signaling	207
Endocrine Signaling	208
Juxtacrine Signaling	208
Role of Signal Transduction in Cellular Communication	208
Receptors	208
Signaling Molecules	208
Signal Transduction Proteins	208
Effector Proteins	209
CELL-CELL SIGNALING	209
Modes of Cell-cell Communication	209
<i>Direct Cell-cell Contact</i>	209

Types of Signaling Molecules	212
<i>Proteins and Peptides</i>	212
<i>Steroids</i>	212
<i>Small Molecules</i>	212
Importance in Tissue Homeostasis	212
<i>Immune Response</i>	212
<i>Growth Regulation</i>	213
<i>Tissue Repair</i>	213
PATHWAYS OF INTRACELLULAR SIGNAL TRANSDUCTION	213
Phases of Signal Transduction	214
<i>Reception</i>	214
<i>Transduction</i>	214
<i>Response</i>	214
<i>Amplification</i>	215
<i>Termination</i>	215
CELL SURFACE RECEPTORS	215
Types of Cell Surface Receptors	215
<i>GPCRs</i>	216
<i>RTKs</i>	217
<i>Ligand-gated Ion Channels</i>	217
<i>Cytokine Receptors</i>	217
Receptor–ligand Interaction	218
GPCR PATHWAYS	219
Mechanism of GPCR Activation	219
Signaling by GPCRs	220
cAMP Pathways	220
IP3/DAG Pathway	221
Physiological and Pathological Roles	222
MAPK PATHWAYS	223
Components of MAPK Pathways	223
MAPK Pathways	224
Extracellular Signal-regulated Kinase (ERK1/2) Pathway	225
JNK Signaling Pathway	226
p38 MAPK Pathway	227
Dysregulated MAPK Signaling in Disease	228
CROSS-TALK BETWEEN GPCRS AND THE MAPK PATHWAY	229
REGULATION AND TERMINATION OF SIGNALING PATHWAYS	230
Feedback Mechanisms	231
Role of Phosphatases and Ubiquitination	231
Implications for Drug Development	232
CONCLUSION	232
CONSENT FOR PUBLICATION	233
CONFLICT OF INTEREST	233
ACKNOWLEDGEMENTS	233
REFERENCES	233

CHAPTER 13 CELL DEATH: MECHANISMS AND MYSTERIES BEYOND APOPTOSIS ... 245

Shambhavi Jha, Rohan Vyas, S. Manvi, Vasanth Kanth T.L., Keerthivasu Ramasamy, K.N. Aruljothi and Ramya Lakshmi Rajendran

INTRODUCTION	246
The Historical Tapestry of Cell Death Research	247

Exploring Diverse Cell Death Pathways	248
PROGRAMMED CELL DEATH	249
Delving into Apoptosis	250
Extrinsic and Intrinsic Pathways in Apoptosis: Dual Routes to Cellular Demise	251
<i>Extrinsic Pathway</i>	251
<i>Intrinsic Pathway</i>	252
Convergence and Caspase Activation	253
<i>Cell Shrinkage</i>	253
<i>Membrane Blebbing</i>	253
<i>DNA Fragmentation</i>	253
<i>Recent Advancements</i>	255
Anoikis: Cell Death due to Detachment	256
<i>Extrinsic and Intrinsic Pathways in Anoikis</i>	257
<i>Integrin-Mediated Signalling Pathway</i>	259
<i>PI3K/Akt Pathway</i>	259
<i>MAPK/ERK Pathway</i>	260
<i>Focal Adhesion Kinase (FAK) Signalling</i>	260
<i>SRC Kinase Pathway</i>	260
<i>Anoikis and Cancer</i>	260
<i>Recent Advancements</i>	261
Ferroptosis: Iron-Dependent Cell Death Mechanism	262
<i>Iron Metabolism and Iron Regulatory Proteins</i>	264
<i>Lipid Metabolism and Lipid Peroxidation</i>	265
<i>Glutathione and Glutathione Peroxidase 4 (GPX4)</i>	265
<i>Nuclear Factor Erythroid 2-Related Factor 2 (NRF2) Pathway</i>	265
<i>Ferroptosis Regulators</i>	265
<i>Recent Advancements</i>	266
Parthanatos: Death by PARP-1 Overactivation	267
<i>PARP Activation</i>	268
<i>Poly(ADP-ribose) (PAR) Polymer Synthesis</i>	268
<i>AIF Translocation and Nuclear Events</i>	268
<i>Involvement of Other Proteins</i>	268
<i>Mitochondrial Dysfunction</i>	268
<i>Interactions with Apoptotic Pathways</i>	269
<i>Recent Advancements</i>	269
NETosis: The Unique Cell Death Pathway	269
<i>Mechanism of NETosis</i>	270
ADCC: The Immune Response Cell Death	270
NON-PROGRAMMED CELL DEATH	272
Necrosis and its Implications for Cell Death	272
Coagulative Necrosis	272
<i>Preservation of Tissue Structure</i>	273
<i>Denaturation of Proteins</i>	273
<i>Nuclear Changes</i>	273
Liquefactive Necrosis	273
<i>Liquid Formation</i>	274
<i>Inflammatory Response</i>	274
Caseous Necrosis	274
<i>Cheese-Like Appearance</i>	275
<i>Loss of Tissue Structure</i>	275
<i>Distinct Granulomas</i>	275

<i>Tissue Discoloration</i>	276
MECHANISMS OF NECROSIS	277
<i>Recent Advancements</i>	279
Exploring mPTP-Mediated Necrosis	279
CELL DEATH: KEY FINDINGS AND FUTURE PROSPECTS	280
CONCLUSION	282
CONSENT FOR PUBLICATION	282
CONFLICT OF INTEREST	282
ACKNOWLEDGEMENTS	282
REFERENCES	282

CHAPTER 14 STEM CELLS: BREAKTHROUGHS IN MEDICINE AND THERAPEUTICS 289

Vanshikaa Karthikeyan, Janani Balaji, SriSri SatishKartik, Dannie Macrin and K.N.

Aruljothi

INTRODUCTION	289
STEM CELL FUNCTIONS	290
TYPES OF STEM CELLS	290
Stem Cells Based on Differentiation Potential	290
<i>Totipotent/Omnipotent Stem Cells</i>	291
<i>Pluripotent Stem Cells</i>	291
<i>Multipotent Stem Cells</i>	291
<i>Oligopotent Stem Cells</i>	291
<i>Unipotent Stem Cells</i>	292
Stem Cells Based on Origin	292
<i>Embryonic Stem Cells</i>	292
<i>Adult Stem Cells</i>	293
<i>Tissue-Resident Stem Cells</i>	293
<i>Induced-Pluripotent Stem Cells</i>	294
<i>Perinatal Stem Cells</i>	294
STEM CELL NICHE	294
Types of Niche	295
<i>Simple Niches</i>	295
<i>Complex Niches</i>	295
<i>Storage Niches</i>	295
EVOLUTION OF STEM CELLS	296
SIMILARITIES AND DIFFERENCES BETWEEN PLANT AND ANIMAL STEM CELLS	296
Similarities	296
Differences	296
INTRODUCTION TO STEM CELL THERAPEUTICS	297
Types of Stem Cells used in Therapeutics	297
<i>Embryonic Stem Cells</i>	298
<i>Adult Stem Cells</i>	298
STEPS IN STEM CELL THERAPY	298
Determination of Stem Cell Source	298
Specification of Cell Dosage	298
Administrative Methods	299
Stem Cell Manipulation for Effective Treatment	299
MAJOR BREAKTHROUGH IN STEM CELL THERAPY	299
MAJOR TYPES OF STEM CELL THERAPY	301
Somatic Cell Nuclear Transfer	301
Implementation of Stem Cell Therapy in Cancer Treatment	302

Risks Associated with Stem Cell Therapy	304
RESEARCH PROSPECTS OF STEM CELL THERAPY	305
PLANT STEM CELLS	306
Plant Tissue Culture	306
Steps in Plant Tissue Culture	307
<i>Pre-propagation</i>	307
<i>In vitro Cultivation</i>	307
<i>Culturing of Explants</i>	307
<i>Micropropagation</i>	307
<i>Hardening</i>	307
Challenges in Plant Tissue Culture	307
INTRODUCTION TO EPIGENETICS IN STEM CELLS	308
KEY EPIGENETIC MECHANISMS IN STEM CELLS	309
DNA Methylation	309
Histone Modification	310
Chromatin Remodeling	310
EPIGENETIC MECHANISMS IN MESENCHYMAL STEM CELLS (HUNTINGTON'S DISEASE)	311
EPIGENETIC CELL REPROGRAMMING IN IPSCS	311
Ectopic Expression of Transcription Factors	311
CRISPR–Cas9-Based Genome Editing for Reprogramming	312
EPIGENETIC THERAPY TARGETING BONE MARROW STEM CELLS	313
RECENT TRENDS	313
Stem Cells in Neurodegenerative Diseases	313
Therapeutic Potential of Dental Stem Cells	314
Cytokines Regulate the Fates of Hematopoietic Stem Cells	314
Bioengineered Scaffolds to Deliver Stem Cells for Wound Healing	315
ASSESSING THE RISKS AND FUTURE POTENTIAL IN STEM CELL REPROGRAMMING	315
CONCLUSION	316
CONSENT FOR PUBLICATION	316
CONFLICT OF INTEREST	317
ACKNOWLEDGEMENTS	317
REFERENCES	317
CHAPTER 15 CANCER STEM CELLS: CATALYSTS OF CANCER PROGRESSION	321
<i>Vedika Kartha, Saloni Semwal, Lakshmi Sai Varshini Yedavalli, Disha Kamath, S. Pooja, Dannie Macrin, Satish Ramalingam and K.N. Aruljothi</i>	
INTRODUCTION TO CANCER STEM CELLS	322
History and Evolution of Cancer Stem Cells	322
Current Concerns and Challenges	322
CHARACTERISTICS OF CANCER STEM CELLS	323
Interaction between Immune and Cancer Stem Cells	324
Intrinsic Features of Cancer Stem Cells	325
BIOLOGICAL PROPERTIES OF CANCER STEM CELLS	325
Marker Expression	325
<i>CD44</i>	325
<i>CD133</i>	326
<i>ALDH (Aldehyde Dehydrogenase)</i>	326
Microenvironment	326
<i>Extracellular Matrix</i>	327

<i>Immune Cell</i>	327
<i>Signaling Molecules</i>	327
TYPES OF CELLS SIGNALLING PATHWAYS RESPONSIBLE FOR CANCER STEM	
CELLS	328
Notch Pathway	328
Hedgehog Pathway	328
CANCER HETEROGENEITY	329
THE ROLE OF CANCER STEM CELLS IN TUMOR DEVELOPMENT AND	
METASTASIS	330
Mechanisms through which Cancer Stem Cells Mediate Tumor Progression and Metastasis	330
Cancer Stem Cells and Metastasis	330
Interaction with the Tumor Microenvironment (TME)	331
Cancer Stem Cells and Angiogenesis in Tumor Progression	332
Cancer Stem Cells in Metastatic Niche Formation	332
Therapeutic Implications of Targeting Cancer Stem Cells in Metastasis	333
METHODS OF CANCER STEM CELL IDENTIFICATION AND ISOLATION	333
Isolation using Cell Surface Markers	334
<i>Limitations</i>	334
Side Population (SP) Assay	335
<i>Limitations</i>	335
Aldehyde Dehydrogenase (ALDH) Assay	335
<i>Limitations</i>	335
Spheroid Formation Assay	336
<i>Limitations</i>	336
Combination Approaches	336
MECHANISM OF DRUG RESISTANCE AND METASTASIS	337
ATP-binding Cassette Transporter (ABC Transporter)	337
Apoptosis Avoidance	337
Numb Protein	337
STRATEGIES TO TARGET CANCER STEM CELLS	337
Target Cancer Stem Cells Markers	337
Target miRNA/LncRNA to Cancer Stem Cells	338
Cancer Stem Cells and Resistance to Therapies	338
Superior DNA Repair	338
Quiescence	338
CANCER STEM CELLS IN DIFFERENT CANCERS	339
Cancer Stem Cells in Breast Cancer	339
<i>New Approaches to Treatment</i>	340
Cancer Stem Cells in Skin Cancer	341
<i>Importance of Cancer Stem Cells in Skin Cancer</i>	341
<i>Methods for Identifying and Investigating Cancer Stem Cells</i>	342
<i>Novel Strategies for the Management of Cancer Stem Cells</i>	342
<i>Continuing and Preventive Care</i>	342
<i>Cancer Stem Cells in Novel Therapies</i>	343
Cancer Stem Cells Play an Important Role in Colorectal Cancer	344
<i>Methods for Identification and Study of Cancer Stem Cells</i>	344
<i>Novel Therapies Targeting Cancer Stem Cells</i>	344
<i>Current Preventive Treatment</i>	345
Cancer Stem Cells in Cervical Cancer	345
<i>Introduction and Importance</i>	345
<i>Characteristics of Cancer Stem Cells in Cervical Cancer</i>	346

<i>Mechanisms of Resistance to Therapy</i>	346
<i>Current Therapeutic Methods and Strategies</i>	347
Targeted Mechanisms against Cancer Stem Cells	347
<i>Targeting the Molecular Signaling Pathways</i>	347
<i>Targeting Cancer Stem Cell Markers</i>	348
<i>Targeting the Cancer Stem Cell Niche And The Quiescent State</i>	348
<i>Manipulation of miRNA Expression</i>	348
<i>Induction of Cancer Stem Cell Apoptosis</i>	348
<i>Induction of Cancer Stem Cell Differentiation</i>	348
<i>Cancer Stem Cells Targeting Therapy</i>	348
Therapeutic Implications of Cancer Stem Cells in Cancer Therapy	348
CONCLUSION	349
ACKNOWLEDGEMENT	349
REFERENCES	349
SUBJECT INDEX	354

FOREWORD

It is with great pleasure that I write this foreword for Dr. Kandasamy Nagarajan ArulJothi's remarkable book, *Cell Biology: Basics to Breakthroughs*. Dr. ArulJothi is a distinguished researcher and academician whose contributions to the fields of genetics, molecular biology, and precision medicine have significantly advanced our understanding of complex cellular processes. With over 70 scientific publications in renowned journals, his work spans diverse areas, including cancer biology, dyslipidemia, exosomal biomarkers, and stem cell therapy, making him a key figure in the field of biomedical research. His active participation in international collaborations, editorial board memberships, and ground-breaking research reflects his unwavering commitment to scientific excellence.

What makes this book particularly valuable is its dual focus on education and research. It serves as an essential resource not only for undergraduate and graduate students seeking to build a strong foundation in cell biology but also for researchers and educators looking to stay abreast of the latest scientific developments. The clear organization, coupled with detailed explanations, illustrative diagrams, and real-world applications, makes complex cellular processes accessible and engaging.

A unique highlight of *Cell Biology: Basics to Breakthroughs* is the scientific contributions from world-renowned experts in the field, who bring diverse insights and expertise. The book features Dr. Krishnan Anand from the University of the Free State, South Africa, whose work in cancer biology and molecular pathology adds depth to the chapters on cancer stem cells and therapeutic interventions. Dr. Satish Ramalingam from SRM University, a distinguished expert in molecular diagnostics and translational medicine, provides valuable perspectives on cell signaling and emerging therapeutic targets. Additionally, Dr. Prakash Gangadharan from the School of Medicine, Kyungpook National University, Republic of Korea, offers his expertise in cellular mechanisms of disease and regenerative medicine, enriching the book with clinically relevant insights. Their collaborative editorial contributions enhance the scientific rigor and global relevance of this volume, making it a truly comprehensive resource for cell biology enthusiasts and professionals alike.

Dr. ArulJothi's ability to integrate basic principles with translational insights makes *Cell Biology: Basics to Breakthroughs* a timely and impactful contribution to the field. By highlighting the clinical and therapeutic relevance of cell biology, this book has the potential to inspire future researchers and contribute to the advancement of biomedical science. I am confident that this book will serve as an invaluable reference for students, scientists, and educators alike, offering profound insights into the intricate world of cells and their role in health, disease, and biotechnology.

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PREFACE

A remarkable trip to the fundamental building blocks of life, cell biology reveals the complicated and crucial mechanisms that regulate the survival, development, and evolution of organisms. This area of study is an incredible adventure into the fundamental building blocks of life. The purpose of this book, titled *Cell Biology: Basics to Breakthroughs*, is to provide readers with a clear and thorough overview of both the fundamentals of cell biology and the most recent advancements in the field. Students, researchers, and anybody else who has an interest in the inner workings of life are the target audience for this kind of instruction since it is designed to gradually expand knowledge, beginning with fundamental principles and progressing to cutting-edge study areas.

In the first few chapters of this book, the authors discuss the origins of life as well as the structural and functional differences that exist between prokaryotic molecules and eukaryotic cells. By reading these chapters, you will gain detailed knowledge that lays the foundation for comprehending the fundamental ideas that underlie cellular biology generally.

After that, we will dig into the primary organelles that are responsible for defining the architecture and function of the cell. In each chapter, an essential component is discussed, including the nucleus, the endoplasmic reticulum, the Golgi apparatus, ribosomes, lysosomes, vacuoles, and plasma membranes. As the reader progresses through these chapters, they will get a comprehensive understanding of how the complex machinery of the cell works in harmony to maintain life forms. After that, we move on to semi-autonomous organelles like mitochondria and chloroplasts, focusing on the distinct functions that these organelles play in the generation of energy and the activities of the metabolic system. The topic of discussion continues with the cytoskeletal components and molecular transport, which are responsible for the conformation, structure, and mobility of the cells within the cell.

It is essential to modern biology to have a solid understanding of cellular communication, and cell signalling and transduction are the processes that explain the intricate ways in which cells detect and react to their surroundings. Consequently, this naturally leads to the regulation of the cell cycle and the processes that regulate cell division, which is then followed by a discussion of cell death, which is an essential step for preserving homeostasis. The clinical and therapeutic aspects of the subject matter are brought into emphasis in the final chapters of the book. It is the study of cancer biology that investigates the uncontrolled development of cells and the harmful effects of this expansion. The final chapters provide an in-depth examination of the intriguing world of stem cells and cancer stem cells, discussing the therapeutic potential of these cells as well as their increasing significance in contemporary medicine.

By the time you reach the conclusion of this book, you will not only gain a profound understanding of cell biology, but you will also have insights into the most recent achievements that are affecting research and therapy. Our intention is that this book will serve as both a resource and a source of inspiration for future investigations in the field of biology.

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This book represents a significant journey of learning, reflection, and progress, made possible through the invaluable support, guidance, and encouragement of numerous individuals. I wish to convey my sincere appreciation to God, my family, and my esteemed research mentors and colleagues.

I want to express my sincere gratitude to my research group, special kudos to Mr. K Kumaran, Ms. Kruthika, Ms. Sanjana D, Ms. Raksa, Ms. Srisri SK, Mr. Danyal Reyaz, Ms. Sayantani C, and Mr. Sasitharan T for their steadfast support and confidence in me. Their love, patience, and understanding during this journey have truly been invaluable to me. I appreciate your support in pursuing my dreams and the understanding you've shown in allowing me the time and space to do so.

I sincerely appreciate the invaluable editorial insights and contributions provided by my beloved associates, Dr. Satish Ramalingam, Dr. Krishnan Anand, Dr. Prakash Gangadaran, Dr. Kirubakaran Rangasamy, and Dr. Dannie Macrin. I truly appreciate their insightful suggestions, which have significantly enhanced my ideas and improved the book beyond what I could have accomplished alone.

I extend my deepest gratitude to my mentor, Prof. Devi Ariketh, whose mentorship, insightful guidance, and unwavering support were instrumental in shaping my research and personal growth.

My heartfelt thanks to my mentor, Prof. Sharon Prince, Professor and Head of the Cell Biology Division at the University of Cape Town, whose guidance and inspiration have profoundly shaped my passion and expertise in cell biology.

Finally, I would like to express my heartfelt gratitude to the Department of Genetic Engineering and SRM Institute of Science and Technology for their unwavering support and invaluable resources, which have been instrumental in the completion of this book.

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CHAPTER 1

Foundations of Life: Cells and Origin

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Abstract: Various theories explain life's origin, including the Oparin-Haldane hypothesis, which suggests that life originated from simple organic molecules in Earth's early reducing atmosphere. This was also supported by the Miller-Urey experiment. The cell is the most fundamental unit of an organism. The cell theory states that all living organisms are composed of cells; an organism's basic unit is a cell, and cells arise from pre-existing cells. A basic cell consists of a nucleus, cytoplasm, cell membrane, and cell organelles. The cell organelles are suspended in the cytoplasm. Prokaryotic cells have an undefined region composed of genetic material called the nucleoid and are devoid of a membrane, unlike eukaryotes. Organelles present in all Eukaryotic cells are the Endoplasmic Reticulum, Ribosomes, Golgi Apparatus, Mitochondria, Plastids, and Vacuoles. Each organelle is specialized to function in a certain way, thereby regulating the cell's metabolism. There is a distinct difference between animal and plant cells. Some constituents are specialized for the plant cell, such as the Cell wall, Vacuoles, and the Plastids. Some are specialized for animal cells, such as Centrioles, lysosomes, Cilia, and Flagella. Cells are effectively detected, viewed, and characterized by numerous tools. The microscope plays an integral role in the world of Cell Biology. Since the invention of the standard microscope, there have been many variations to it, enhancing our ability to view microscopic structures.

Keywords: Abiogenesis, Cell organelles, Eukaryotes, Living being, Prokaryotes.

INTRODUCTION**Theory of Spontaneous Generation**

An ancient theory proposed around 350 BC was widely believed until the seventeenth century. Greek Philosopher Aristotle articulated that living things could arise from non-living components. He did so by raising striking questions,

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such as how new fish are introduced in a freshly formed pond or frogs' unexpected appearance on the Nile River's banks [1]. He proposed that an organism can arise from non-living material if the material has *pneuma* or 'vital heat'.

For a long time, it was believed that living organisms could spontaneously arise out of their niche. It wasn't until the late 17th century that Francesco Redi, an Italian Scientist, refuted the Theory of Spontaneous Generation by providing solid proof that maggots did not appear spontaneously on a slab of rotten meat stored for days [2]. He disproved this by storing one slab in an airtight container, another in the open, and the other covered with gauze. He had observed that only that maggot arose on the uncovered meat. He later concluded maggots did not arise from the beef in the airtight contained. However, he noticed maggots on the gauze and concluded that the maggots were offspring of common flies.

Furthermore, Louis Pasteur (1859) continued to disprove the theory by boiling meat broth in a swan-necked flask. He aimed to prove that the downward curve of the flask prevents the particles from reaching the broth, hence hindering growth. When the flask was overturned, the particles could reach the broth much better, and it clouded immediately. The 'Law of Biogenesis', meaning life arises from previously existing life, had a good conclusive result, but not many favored Pasteur's findings [3].

It was not until John Tyndall's experiment in 1876 that Pasteur's conclusions were supported. Tyndall repeated Pasteur's experiments only to observe that some boiled growth media had remained sterile while others did not, despite boiling for a long time. He discovered that endospores (heat-resistant species that can develop into bacteria under favorable environmental conditions) could grow on a nutrient medium, posing a hindrance to arriving at an accurate conclusion [4]. Tyndall thus devised a sterile medium where even endospores could not survive, which required a series of mechanisms and steps to ensure sterility.

Abiogenesis for the Origin of Life

The notion that living organisms arose from non-living matter on Earth more than 3 billion years ago forms the backbone concept of abiogenesis. Russian biochemist Aleksandr Oparin and British scientist J.B.S. Haldane, in the 1920s, proposed individual ideas that formed a theory, and the basic outline of it is that non-living matter, with the help of an external source of energy, for example, UV radiation, can form organic molecules [5].

Oparin believed Coacervates were the precursors to the basic cell, and they may have been one of the very first entities to exhibit life-like properties, such as

growth and reproduction. Coacervates are droplet-like structures formed during liquid-liquid phase separation [6]. Oparin, backed by many other prominent scientists, firmly believes that Coacervates are the precursors of cells.

Haldane believed that simple inorganic materials formed into more complex molecules in the presence of external energy sources, such as UV radiation, over time. The combination of both notions is what laid an essential foundation for abiogenesis.

The Miller-Urey experiment performed in 1953 by American scientists Harold C. Urey and Stanley Miller provided solid proof for the Oparin-Haldane hypothesis [7]. The scientists sealed Hydrogen, Methane, Ammonia, and water in a glass flask and generated an electrical discharge, simulating lightning. After a week of discharging electrical sparks, it was observed that the water had turned reddish and turbid, and yellow-brown deposits were on the electrodes inside the apparatus. This indicated the synthesis of amino acids – the building blocks of protein. The groundbreaking experiment and Oparin-Haldane theory paved the way for research focused on Astrobiology- the study of life in the universe.

Cell and its Contents

Following the invention of the microscope in the 16th century, Robert Hooke, in the year 1665, observed small compartments or honeycomb-like structures and named them cells, as the cellulose walls in the cork he observed reminded him of rooms or monasteries usually occupied by monks (cellula) (Fig. 1). The Latin word ‘cellula’ means storeroom or chamber. All of these were published in his work *Micrographia* [8]. These small compartments would become all living organisms' basic, functional, and structural units.

In 1674, Anton Van Leeuwenhoek seemingly observed protists and, years later, bacteria, which he termed the coin ‘animalcules,’ meaning microscopic animals [9]. This was the first instance a living cell had been discovered. French Chemist Françoise Raspail laid an essential foundation for the Cell Theory – that cells arise from pre-existing cells, which he hypothesized from witnessing binary fission, wherein a single cell divides into two daughter cells [10].

CHAPTER 2

Prokaryotes vs. Eukaryotes: Comparative Structural and Functional Insights

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Abstract: Organisms are fundamentally divided into prokaryotes and eukaryotes, except for viruses. Further classifications are depicted in various classifications, among which the most recent one is Whittaker's classification. Prokaryotes, the primitive organisms, gave rise to eukaryotes, and this transformation led to simple single-celled organisms evolving into colonies and multicellular cells. Prokaryotic cells, which include bacteria and archaea, exhibit diverse metabolic processes such as autotrophy and heterotrophy. In contrast, eukaryotic cells are characterized by the presence of a well-defined nucleus and membrane-bound organelles, unlike prokaryotic cells. In contrast, eukaryotes developed around 2 billion years ago and possess complex cellular structures, including a well-defined nucleus and various organelles, such as mitochondria and chloroplasts. The transition from unicellular to multicellular life is a significant evolutionary milestone that involves various adaptations and mechanisms, including cell-to-cell communication, adhesion, and coordinated growth. Genetic conservation and epigenetic mechanisms play pivotal roles in the development of multicellular structures, as demonstrated in organisms like fungi and metazoans. Eukaryotic cells, such as those from yeast and mammalian sources, are pivotal in biotechnological applications, including the production of recombinant proteins and gene therapy. Their ability to properly fold and process proteins allows for the creation of functional biopharmaceuticals and vaccines that simulate pathogen structures to invoke robust immune responses. Notable eukaryotic microorganisms like algae and fungi are also increasingly recognized for their potential in sustainable biofuel production. Since genes serve as the backbone for almost all cells, they can be manipulated to be more user-friendly.

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Keywords: Biofuel production, Characteristics, Eukaryotes, Evolution, Evolutionary biology, Gene therapy, Genes, Multicellularity, Prokaryotes, Recombinant proteins, Whittaker's classification.

INTRODUCTION TO PROKARYOTES

With a simpler cell structure than eukaryotes, prokaryotes are unicellular creatures devoid of membrane-bound organelles or a nucleus; their genetic material is arranged in a single circular DNA molecule that floats freely inside the cell. Common examples of prokaryotes, bacteria, and archaea lack compartmentalization, which helps them to adapt and flourish in many habitats, as shown in Fig. 1. Among the first living entities on Earth, prokaryotes have greatly affected metabolic reactions, forming the ecosystems of the planet [1]. Originally discovered in the late 17th century with the development of the microscope, Dutch scientist Antonie van Leeuwenhoek saw “animalcules”—microbes—in 1676, hence establishing the basis for microbiology [2]. While Carl Woese's genetic categorization in the 1970s separated Archaea from Bacteria, redefining our knowledge of prokaryotes, advances in staining and microscopy techniques helped scientists like Louis Pasteur and Robert Koch identify bacteria as agents of illness.

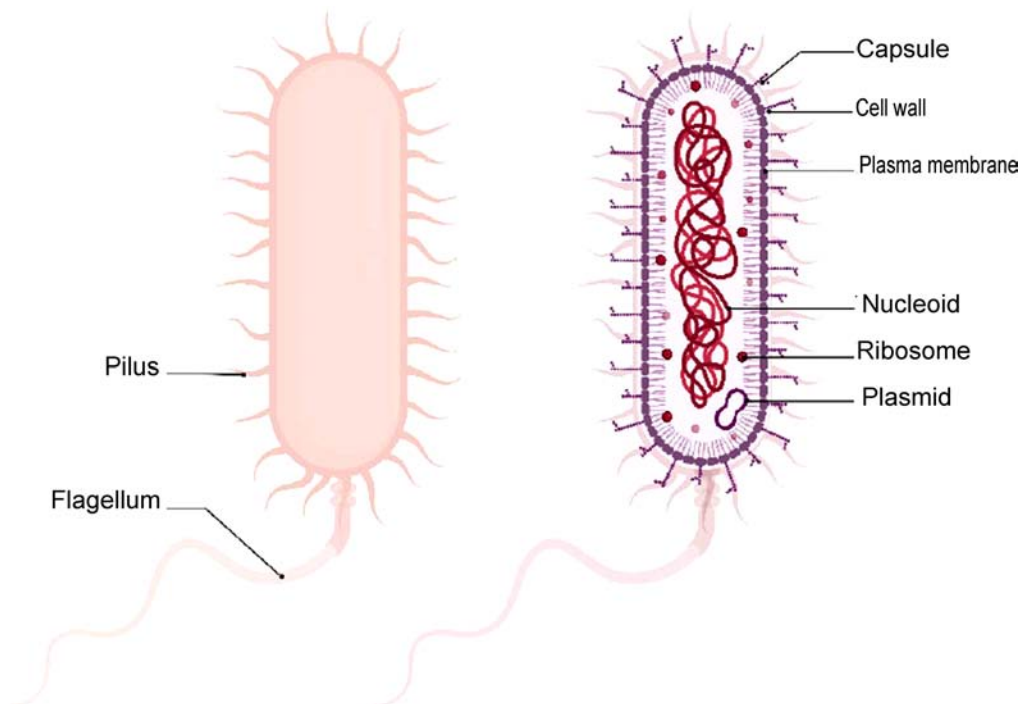


Fig. (1). Bacterium, a prokaryote

Crucially for plant development, prokaryotes drive biogeochemical cycles that help break down organic matter and recycle nutrients and nitrogen fixation, supporting ecosystems. Beyond their function in the surroundings, prokaryotes are essential for human life since they help digestion and enable waste breakdown and fermenting [3]. With their research opening the path for industrial uses, bioremediation, and genetic engineering, they also synthesize vitamins, antibiotics, and other chemicals, thereby stressing the need for prokaryotes to maintain the environment and improve health and quality of life.

Prokaryotic Diversity

Prokaryotes exhibit different forms, structures, metabolic pathways, and ecological responsibilities among their tremendous and varied diversity. Their adaptability lets them live practically anywhere on Earth, from human bodies to deep waters. Their taxonomy, form, metabolic strategies, and ecological niches are examined here.

Classification of Prokaryotes

Separated by genetic, biochemical, and physiological traits, bacteria and Archaea represent the two main domains of prokaryotes. Among the most well-known prokaryotes—from helpful gut microorganisms to disease-causing pathogens—are bacteria [4]. Living in almost every environment on Earth, they are pretty varied and have cell walls typically made of peptidoglycan, a unique polymer providing structural support. Previously believed to be bacteria, Carl Woese identified archaea as a discrete domain based on their distinct genetic and metabolic properties. Like bacteria, Archaea can survive in hostile conditions because their cell membranes consist of ether-linked lipids rather than ester-linked lipids. Usually, extremophiles survive in hot springs, salt flats, and deep-sea vents. Archaea can also be found in more common environments like the human intestine [5].

Morphological Diversity

The range of forms, sizes, and configurations that prokaryotes exhibit enhances their adaptability and usefulness in many contexts. Among the most often occurring forms are cocci (spherical-shaped bacteria such as *Streptococcus*), bacilli (rod-shaped cells like *Escherichia coli*), *spirilla* (spiral-shaped cells like *Spirillum*), and vibrios (comma-shaped cells like *Vibrio cholerae*). Usually small, ranging from 0.2 to 5 micrometers, prokaryotic cells can vary in size depending on their environment and genes. Division patterns affect their cellular structures as well; they produce linear chains (as seen in streptococci), grape-like clusters

Nucleus: the Control Centre of the Cell

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Abstract: The nucleus is a crucial organelle in eukaryotic cells, serving as the primary center for genetic material and essential cellular processes like replication, transcription, and gene expression. The nucleus, surrounded by a double-membrane nuclear envelope, includes critical elements like chromatin, the nucleolus, and nuclear pore complexes, each integral to functions such as gene expression, RNA processing, and ribosome assembly. This document offers a comprehensive examination of nuclear structure and its functional dynamics, highlighting the importance of Cajal bodies in RNA metabolism and the influence of nuclear organization on gene regulation. Furthermore, it underscores the consequences of nuclear irregularities in several illnesses, such as laminopathies and chromatin remodeling disorders, which may have significant health ramifications. As research advances, novel treatment methods aimed at nuclear functions, including gene editing technologies and techniques for treating laminopathies, are becoming increasingly prominent. These observations highlight the nucleus's essential function in preserving cellular integrity and operation and its significance in health and disease management.

Keywords: Cajal bodies, CRISPR-Cas9, DNA replication, Eukaryotic cells, Gene expression, Genetic material, Hutchison-Gilford Progeria Syndrome, laminopathies, nuclear matrix, nuclear pore complexes, RNA processing, Spinal muscular atrophy, Treacher Collins Syndrome.

INTRODUCTION

The nucleus is a membrane-bound organelle that acts as the command center of eukaryotic cells, ensuring genetic information integrity and controlling cellular activity. It is usually the most visible organelle in the cell, occupying a large percentage of the cellular volume. Gene expression, DNA replication, and cell division occur exclusively because of the eukaryotic cell's nucleus. In the prokaryotes, a nucleoid is present instead of a nucleus. The genetic material known as DNA (deoxyribonucleic acid) will be present as chromatin, a mixture of

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proteins and DNA that is arranged into chromosomes during cell division. The DNA that codes for everything from protein synthesis to cell growth and reproduction is found in the nucleus in eukaryotes / nucleoid in prokaryotes and includes all the instructions for the cell to function. Despite the more straightforward organization of genetic material, prokaryotes effectively regulate gene expression and adapt rapidly to environmental changes. Their streamlined genome organization and mechanisms, like operons (clusters of genes under a single promoter), enable efficient and coordinated gene expression. This efficiency contributes to their high adaptability and survival across diverse environments. At the same time, eukaryotic cells have a highly organized nucleus with complex DNA packaging that facilitates intricate regulation of gene expression and cellular specialization. In contrast, prokaryotic cells lacking a defined nucleus possess a more straightforward yet efficient system of DNA organization that supports rapid growth and adaptability. The presence of a membrane-bound nucleus is the characteristic that sets eukaryotic cells apart from prokaryotic cells, which have genetic material distributed throughout the cytoplasm and no clearly defined nucleus [1].

History

Over the decades, the discovery and comprehension of the nucleus have undergone substantial changes, playing a pivotal role in advancing cell biology. Robert Brown, a Scottish botanist, coined the term “nucleus” for the first time in a biological context in 1831. Brown saw a tiny, thick, spherical structure inside the cells of orchids and other plants while investigating plant cells under a microscope. He called this structure the “nucleus,” which is Latin for “kernel” or “core.” Though Brown did not wholly comprehend its purpose then, he recognized it as a recurring and essential aspect of the cell. In the years that followed, many scientists verified Brown's findings and started to make assumptions regarding the function of the nucleus. Matthias Schleiden and Theodor Schwann contributed to the cell theory in the middle of the 19th century. This theory postulated that all living things comprise cells, with the nucleus considered a crucial component [2].

Expanding on this idea, Rudolf Virchow proposed in 1855 that all cells originate from pre-existing cells, highlighting the significance of the nucleus in cell division and inheritance. Improvements in staining methods and microscopy during the late 19th century gave scientists a better understanding of the nucleus. Walther Flemming's 1880s discovery of chromosomes inside the nucleus demonstrated their function in mitosis or cell division. Later, August Weismann put out the theory that the nucleus contained genetic material, which impacted the field of genetics. Thomas Hunt Morgan and his associates' work showed that

genes found on chromosomes inside the nucleus were the units of heredity, significantly advancing our knowledge of the function of the nucleus in the early 20th century. This discovery confirmed the nucleus's function in genetic information storage and inheritance [3]. The nucleus's function as the cell's command center was further demonstrated in 1953 with the discovery of the DNA structure by James Watson and Francis Crick. It was discovered that DNA, which is found in the nucleus of eukaryotic cells, is the molecule in charge of storing and transferring genetic information. This discovery made modern molecular biology possible by establishing the nucleus as the primary locus of gene control, replication, and expression [4].

STRUCTURE AND FUNCTION

Structure of Nucleus in Eukaryotes

The nucleus comprises several parts that form a highly ordered structure supporting its role as the cell's command center. The nuclear envelope is a two-layered barrier that surrounds the cell nucleus and keeps it separate from the cytoplasm. This envelope consists of an inner and outer membrane, both lipid bilayers. The outer membrane is continuous with the endoplasmic reticulum, facilitating the transport of materials between the nucleus and cytoplasm. Nuclear pores are embedded within the nuclear envelope, which regulate the passage of molecules such as nucleic acids and proteins into and out of the nucleus. A gel-like substance fills the nuclear space inside the nucleoplasm, providing structural support and a medium for suspending various nuclear components. Among these components, chromatin, a complex of DNA and proteins, plays a crucial role in packaging genetic material and regulating gene expression. Another vital structure within the nucleoplasm is the nucleolus, a dense, spherical body primarily responsible for producing and assembling ribosomes (Fig. 1) [1].

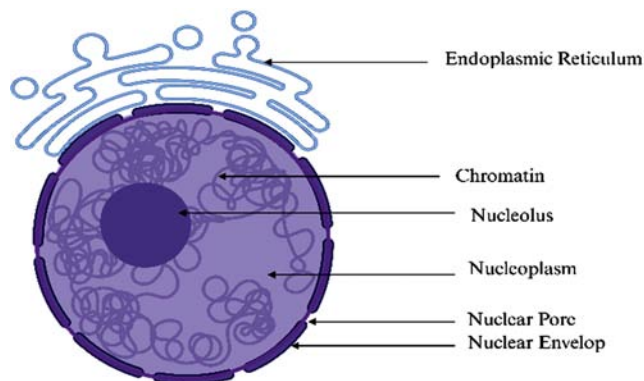


Fig. (1). Structure of the nucleus in eukaryotes.

CHAPTER 4

Endoplasmic Reticulum: The Cellular Factory

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Abstract: The endoplasmic reticulum is an organelle that performs dynamic functions in many critical cellular processes. This chapter discusses the evolutionary aspects of the Smooth ER (SER), the tasks of RER, their involvement in cellular stress responses, and implications for human health and disease. Complexities of ER stress and their relationship to conditions such as ulcerative colitis and type 2 diabetes will be discussed. The chapter discusses the therapeutic potential of targeting ER stress pathways as therapy for UC. We also discuss the relationship between the ER and autophagy, a cellular degradation and recycling process. Discussion is taken upon the role of the mitochondria-associated ER membrane (MAM) in calcium signaling and its implications for various cellular processes. This chapter summarizes the roles of ER, trying to unveil the complexity of the ER functions and the importance of the ER in both human health and disease.

Keywords: Apoptosis, Autophagy, Endoplasmic Reticulum, ER Stress, Glycosylation, Homeostasis, Lipid Synthesis, Mitochondria, Mitochondria-Associated ER Membrane (MAM), Protein Folding, Quality Control, Smooth ER, Type 2 Diabetes, Ulcerative Colitis.

INTRODUCTION

The endoplasmic reticulum encompasses a membrane-bound organelle with the framework of a network of sacs and tubules. It is continuous with the nuclear envelope. The ER is vital in shipping the synthesized proteins to the Golgi apparatus for further processing. Apart from synthesizing proteins, the ER synthesizes the lipids that essentially serve as the membrane in all organisms. The endoplasmic reticulum can be classified into two types based on the compounds they synthesize: Smooth Endoplasmic Reticulum (SER) and Rough Endoplasmic

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Reticulum (RER). The Rough ER has attached ribosomes that help in protein synthesis. Smooth ER lacks ribosomes, which are key drivers in protein synthesis. Therefore, the SER is involved in the production of lipids, and the RER plays a role in Protein synthesis. Apart from synthesizing proteins and lipids, the ER plays other vital roles in transporting proteins and lipids, storing Ca^{2+} ions, steroid production, stress regulation mechanisms (endoplasmic reticulum stress), and protein folding machinery [1]. Fig. (1), below, denotes the structural orientation of the endoplasmic reticulum. Rough ER has ribosomes on its surface, whereas smooth endoplasmic reticulum lacks ribosomes.

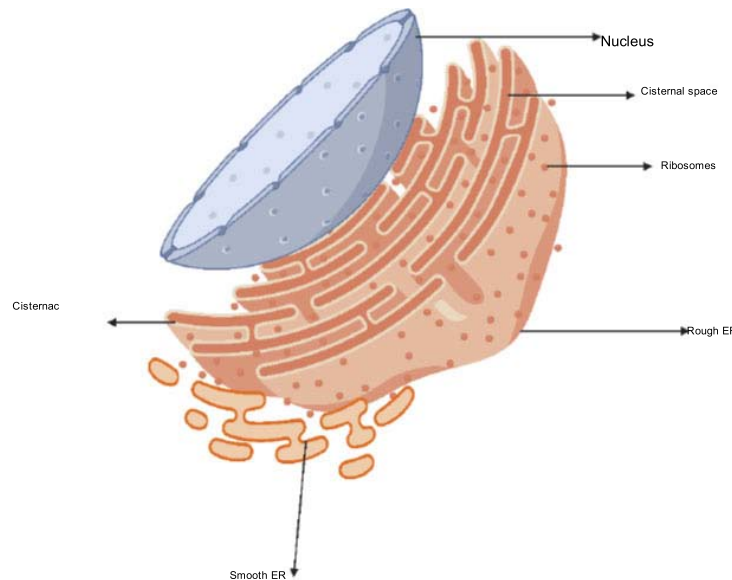


Fig. (1). Structural orientation of endoplasmic reticulum.

FUNCTIONS OF RER

The protein translocation in the RER occurs in two ways,

1. Where the protein, while being synthesized by the ribosomes, undergoes the translocation and translation simultaneously in ER, called the co-translational translocation.
2. The proteins synthesized by the free ribosomes are translocated into the ER using the SEC pathway.

The ER synthesizes three significant proteins: secretory proteins, integral membrane proteins, and Luminal resident proteins. The secretory proteins are involved in folding proteins and delivering specific proteins. The integral

membrane proteins are derived from the ER, and almost 30% of the entire human genome codes for the integral membrane proteins. The last category of proteins is called the luminal resident proteins of the ER, Golgi, Nuclear envelope, and Lysosomes. Therefore, RER plays a significant role in the biosynthesis of proteins. An example of the integral membrane protein is Single Carboxy-terminal TM spans (tail-anchored membrane proteins). Most organelles' protein requisite are satisfied by the protein synthesis at the RER. Therefore, molecules for protein synthesis must reach the ER for processing, followed by proper folding. The protein folding machinery folds the protein synthesized by the ribosomes at the RER, called the chaperones. Folding is essential for building a fully structural and functional unit of protein. Therefore, proteins reach a thermodynamically stable state. Also, proteins are not independent in their functioning. Their interactions can range from short and temporary within their functional group to stable oligomers in biological synthesis. The majority of proteins in the secretory pathway include oligomers. Therefore, the proteins must be folded at the late translational steps [2].

The Process of Protein Folding Associated with Molecular Chaperones

The chaperon-assisted protein folding is followed by the assembly of multi-subunit proteins, formation of the disulfide bond, and N-glycosylation (initial stages); in the case of the proteins of the plasma membrane, glycolipid anchors are added to the proteins. The Luminal proteins in the ER play the primary role in assisting the folding of proteins and assembly of the polypeptide units that are newly translocated. The molecular chaperones, being proteins, act on a non-native protein to stabilize or bring about its native conformation. These molecular chaperones do not appear in the final three-dimensional confirmation of the proteins. The major ER resident chaperones are Hsp70 Bip, lectin chaperones like calnexin, and calreticulin (folding glycoproteins). Bip releases the abnormally folded proteins, which are then removed from the ER by the Endoplasmic reticulum-associated degradation process [3]. In the process of N-glycosylation, asparagine residues of proteins are glycosylated that are present within the ER. By this time, the translation is still in process (co-translational). In the first step, the oligosaccharides with 14 sugar residues are added to the acceptor molecules of protein, that is, the asparagine residue. The lipid carrier dolichol is the site for synthesizing the oligosaccharide anchored to the ER membrane. From here, the oligosaccharide unit is transferred to the asparagine residues by an enzyme associated with a membrane called the oligosaccharide transferase. Three glucose units are removed from the long oligosaccharide, and the protein remains in the endoplasmic reticulum. As discussed earlier, the protein from the ER has to reach the Golgi apparatus for further modification, packing, and shipping. Glycosylation is significant as it provides signals to promote protein folding, followed by

CHAPTER 5

Golgi Apparatus: Shaping and Shipping Cellular Proteins

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Abstract: The Golgi body is an organelle in charge of altering proteins and lipids after translation and getting them ready for transport to spots within and outside the cell. This part of our study investigates its makeup, focusing on three key sections— cis side, middle section, and trans side, and how they contribute to sugar attachment to proteins, lipid processing, and protein arrangement. Furthermore, the chapter discusses how it plays a role in cell functions like secretion and creating lysosomes, and its impact on diseases like neurodegenerative conditions, infections, and cancer. It also covers progress in treatments and research related to the Golgi apparatus, including its involvement in the dynamics of COVID-19.

Keywords: Cancer, Cisternae, COVID-19, Golgi Apparatus, Lysosome Formation, Neurodegenerative Diseases, Phosphorylation, Post-Translational Modification, Protein Sorting, RAB GTPase, Sphingomyelin, TGN (Trans Golgi Network), A-Synuclein.

INTRODUCTION

The Golgi apparatus is a fundamental organelle and a key feature of eukaryotic cells that distinguishes them from prokaryotes. It consists of membrane-bound vesicles attached to stacked, flattened sacs known as cisternae. It helps in the post-translational modification and sorting of proteins and lipids. It packages these modified molecules into vesicles, which are then transported to specific locations inside and outside the cell through endocytic and exocytic pathways. As a result, it is vital for preserving cellular homeostasis and facilitating proper cell functions, making it crucial for intracellular transport and protein secretion. The Golgi apparatus can be traced back to the Last Eukaryotic Cell Ancestor (LECA), implying that the presence of the Golgi is nearly two billion years old, originating around the time of the divergence of major eukaryotic lineages [1]. The endosym-

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biotic theory has been turned down, and the trending view is that the Golgi and other organelles of the endomembrane system evolved through autogenous means, meaning that they originated from inside the cell and not from endosymbiosis. Functional diversity exists in the different eukaryotes, from the ring Golgi in the malaria parasite to the mobile Golgi in the plants. This merely underlines the functional plasticity of the Golgi and its adaptation to various ecological niches.

STRUCTURE OF GOLGI APPARATUS

It is worth noting that the early description of the Golgi apparatus was “apparato reticulare interno”, meaning the internal reticular apparatus was based on its morphology. It is a membrane-bound organelle, made of a series of cisternae. These stacks often appear as a series of flattened pancakes or discs. The size of the Golgi stacks can differ, with each stack containing a variable number of cisternae. These stacks are encircled by a network of vesicles and tubules, which are essential for the transport of materials to and from the Golgi apparatus.

The Golgi apparatus has three compartments, called “cis”, “medial,” and “trans”. The cis compartment is the cisternae near the Endoplasmic Reticulum (ER), the medial compartment is the central layers of cisternae, and the trans compartment is the cisternae farthest from the ER. There are two Golgi networks, which are the Cis Golgi network and the Trans Golgi network; these comprise the outermost cisternae located at the cis and trans faces of the Golgi, respectively. These networks are responsible for sorting the proteins and the lipids that are received at the cis face from the ER and released at the trans face by the Golgi (Fig. 1).

CIS Face

The cis face in the Golgi apparatus is closest to the ER. It is located to facilitate the transfer of the newly synthesized protein and lipids from the ER. Its primary function is to receive materials (protein and lipid) that need to be processed in the Golgi apparatus. It also initiates modification of the incoming molecules, like glycosylation, sulfation, and phosphorylation. It recognizes the specific signals on the incoming molecules, which decides its destination within the Golgi stacks, thus it is the “receiving a sorting hub” of the organelle. The Golgi apparatus consists of stacked cisternae linked by tubular structures, forming a network referred to as the Golgi ribbon. At the entrance of the Golgi, known as the cis-Golgi, Vesicular Tubular Clusters (VTCs) act as a transitional area between the endoplasmic reticulum and the stacked cisternae [2].

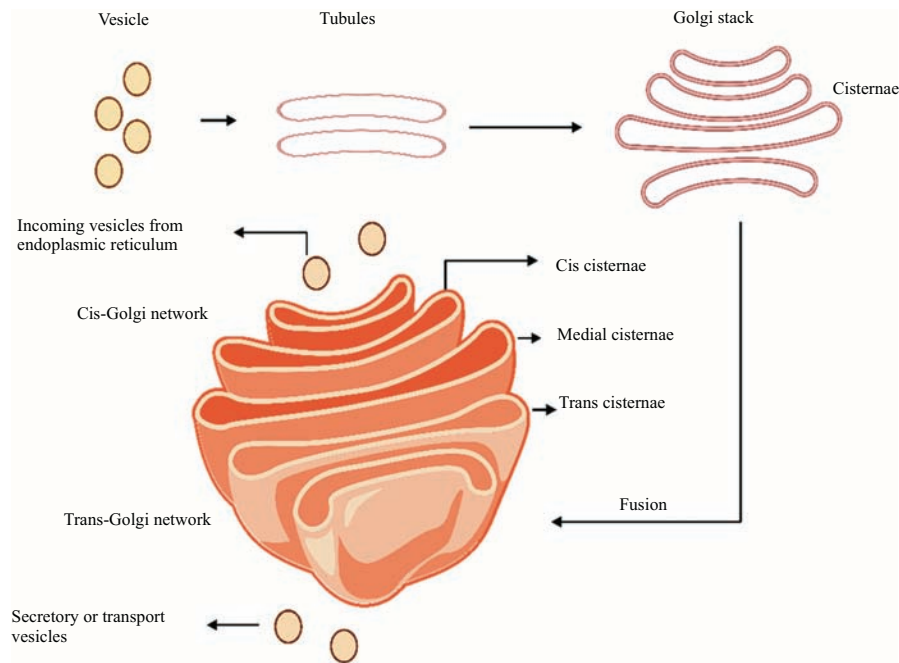


Fig. (1). The Golgi apparatus is composed of stacked cisternae interconnected by tubules, receiving vesicles from the endoplasmic reticulum and sorting proteins for their final destinations.

MEDIAL Face

The medial face is positioned between the cis face and the trans face within the Golgi stacks. The primary role of the medial face is to modify proteins and lipids. It is at this stage that many post-translational modifications occur, including glycosylation (addition of sugar molecules), sulfation, and phosphorylation, which are crucial for the functional maturation of proteins and the alteration of lipid molecules. The medial cisternae contain various enzymes that facilitate these modifications. It also helps in pH regulation within the cisternae, which is crucial for the functioning of the enzymes involved in various modifications. The medial face is responsible for further processing materials destined for secretory vesicles that will carry proteins and lipids to the cell membrane or for the formation of lysosomes, which contain enzymes for breaking down cellular waste.

TRANS Face

The trans face is positioned at the opposite end of the Golgi apparatus from the cis face and is called the “shipping” face, as it is responsible for directing modified molecules to their final destinations within or outside the cell. The primary role of the trans face is to sort out the proteins and lipids that have undergone

CHAPTER 6

Ribosomes: Engines of Protein Synthesis**Sanjana Dhayalan¹, Shalini Roy¹, K. Kumaran¹ and K.N. Aruljothi^{1,*}**¹ *Department of Genetic Engineering, SRM Institute of Science and Technology, Kattankulathur, Chengalpattu, Tamil Nadu, India*

Abstract: Ribosomes are essential macromolecular complexes that translate genetic information from mRNA into proteins. Composed of rRNA and ribosomal proteins, they consist of two subunits, large and small, in both prokaryotic (70S) and eukaryotic (80S) cells, with structural differences reflecting their evolutionary adaptations. Ribosome biogenesis (Ribi) is a tightly regulated process initiated by rDNA transcription and subunit assembly in eukaryotes and involving factors like NusA and NusG in prokaryotes. Ribosomes drive protein synthesis through complex processes of initiation, elongation, termination, and recycling, facilitated by GTPases and elongation factors, ribosomopathies cause diseases such as Diamond-Blackfan Anemia and Shwachman-Diamond Syndrome, linked to mutations in ribosomal proteins. Ribosomal dysfunction can lead to both hypo- and hyper-proliferation, increasing cancer risk. Recent advances in targeting ribosome-related pathways, such as mTORC1 inhibition, and techniques like ribosome profiling have provided insights into diseases like leukemia and medulloblastoma, revealing non-canonical ORF translation and novel therapeutic targets. These findings highlight the therapeutic potential of modulating ribosome function in disease treatment.

Keywords: ABCE1, Biogenesis, Cancer, Elongation, Eukaryotes, Fingerprinting, GTPases, Prokaryotes, Protein Synthesis, Ribosomal Subunits, Ribosome Profiling, Ribosomopathies, rRNA, Termination, tRNA.

INTRODUCTION

Ribosomes are essential cellular organelles found in all living organisms, serving as the molecular machinery responsible for protein synthesis, a fundamental biological process. These intricate macromolecular complexes comprise ribosomal RNA (rRNA) and ribosomal proteins; they are essential for converting genetic information from messenger RNA (mRNA) into polypeptide chains, which fold into valuable proteins. These indispensable organelles play a central role in synthesizing proteins, which are vital for the structure and function of all

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living organisms. Their intricate composition and sophisticated mechanisms underscore the complexity of cellular processes and the evolutionary significance of protein synthesis. The ribosomal function provides insight into fundamental biological processes and has implications for medical research, particularly in developing antibiotics that target bacterial ribosomes without affecting eukaryotic cells. Ribosomes are categorized into two primary types based on cellular organization: prokaryotic and eukaryotic ribosomes. Despite their differences, both types share a similar functional architecture. The structural differences between prokaryotic and eukaryotic ribosomes are significant, particularly in the size and composition of the rRNA and protein components [1].

Ribosome in Prokaryotes and Eukaryotes

In eukaryotic cells, ribosomes include two distinct subunits: the large ribosomal subunit (60S) and the small ribosomal subunit (40S). The 60S subunit comprises three rRNA molecules: 5S, 5.8S, 25S/28S rRNA, and 46 associated ribosomal proteins. The small 40S subunit contains a single rRNA molecule, the 18S rRNA, associated with 32 ribosomal proteins. The highly regulated process of ribosomal component assembly takes place in the nucleolus, where mature ribosomal subunits are formed by the combination of ribosomal proteins and rRNA transcription [2]. Prokaryotic ribosomes in organisms such as bacteria are slightly smaller than their eukaryotic counterparts. They consist of a 50S subunit and a 30S subunit. 34 proteins and 5S and 23S rRNA are found in the 50S subunit, whereas 21 proteins and 16S rRNA are found in the 30S subunit.

The protein synthesis, or translation, involves several key steps and the coordinated action of both ribosomal subunits [3]. The small ribosomal subunit first attaches itself to the mRNA molecule and searches for the start codon. The assembly of the ribosomal subunits—40S and 60S in eukaryotes, 30S and 50S in prokaryotes—around the mRNA molecule is the first step in the translation process. In eukaryotes, the initiator tRNA, which carries methionine (Met-tRNAⁱ), and the small 40S ribosomal subunit work together to scan the messenger RNA (mRNA) for the start codon (AUG). The big 60S subunit joins the complex upon identification of the start codon, completing the formation of the 80S ribosome.

The Shine-Dalgarno sequence on the mRNA is bound by the 30S subunit in prokaryotes, and the 50S subunit then joins to form the 70S ribosome. The large and tiny subunits join forces upon identification of the start codon to produce a functional ribosome that can synthesize proteins.

During translation, the ribosome facilitates the decoding of the mRNA sequence into a corresponding amino acid sequence. The primary function of the small

subunit is to decode the messenger RNA (mRNA) and guarantee that the appropriate transfer RNA (tRNA) molecules, each containing a distinct amino acid, are introduced into the ribosomal A (aminoacyl) site. Through its peptidyl transferase activity, the larger subunit facilitates the creation of peptide bonds between neighbouring amino acids once the proper tRNA has been positioned. The elongation of the developing polypeptide chain depends on this enzyme action. The tRNA molecules are successively added as the ribosome translocates along the mRNA. When the expanding polypeptide chain reaches a stop codon, it is released from the ribosome, indicating that translation has stopped. The completed polypeptide then undergoes folding and post-translational modifications to achieve its final functional conformation.

In eukaryotic cells, ribosomes are crucial for protein synthesis and can be found in several distinct locations, each serving specific functions. They exist in two primary forms: freely floating in the cytoplasm, referred to as cytoribosomes, or attached to the outer surface of the Rough Endoplasmic Reticulum (RER), thereby forming the rough ER. This association with the RER is significant as it facilitates the co-translational translocation of nascent polypeptide chains into the lumen of the endoplasmic reticulum. Once inside the ER, these polypeptides undergo further modifications, such as glycosylation and folding, before being trafficked to their final destinations within or outside the cell. Additionally, ribosomes are present in the stroma of plastids, such as chloroplasts, and within the mitochondrial matrix. In these organelles, ribosomes synthesize proteins encoded by their respective organellar genomes, which are distinct from nuclear DNA. This unique feature underscores the endosymbiotic theory, suggesting that mitochondria and plastids originated from ancestral prokaryotic cells. In contrast, prokaryotic cells, which lack membrane-bound organelles, exhibit a simpler ribosomal distribution. Ribosomes in prokaryotes are freely distributed throughout the cytoplasm. These ribosomes, typically smaller than their eukaryotic counterparts, are involved in translating mRNA into proteins directly within the cytoplasmic environment. This structural simplicity allows for rapid protein synthesis, which is essential for the adaptability and survival of prokaryotic organisms in diverse environments. The localization of ribosomes in eukaryotic and prokaryotic cells reflects their essential roles in protein synthesis, with distinct adaptations that facilitate cellular functions and processes.

Origin and Evolution of the Ribosome

The origins of ribosomes can be traced back to the early Earth, approximately 4 billion years ago, when the first molecules of life began to form. It is widely accepted that simple RNA molecules, capable of catalyzing chemical reactions, played a crucial role in the emergence of life. This concept is often referred to as

Lysosomes: The Cell's Digestive System

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Abstract: Lysosomes serve as essential organelles in eukaryotic cells, facilitating the recycling and degradation of cellular waste through the action of hydrolytic enzymes. This acidic organelle engages with various vesicles, such as phagosomes and endosomes, to dismantle biomolecules encompassing proteins, lipids, and nucleic acids. Recent studies have illuminated the function of lysosomes in autophagy, a process wherein they contribute to the degradation of cellular constituents in response to stress or periods of starvation. Three main types of autophagic processes, macroautophagy, microautophagy, and chaperone-mediated autophagy, use central mechanisms that help the trafficking of their selective cargo. The activity of lysosomes is connected with several diseases: lysosomal storage diseases, Parkinson's disease, and muscular dystrophy are generally caused by either a lack of efficiency of the autophagosome and lysosome fusion or impairment of lysosome digestion. Mechanisms of lysosomal regeneration involve Autophagic Lysosome Reformation (ALR) and Endocytic Lysosome Reformation (ELR), with the presence of essential functions to maintain lysosomal function. Further knowledge about such processes may allow for the creation of therapies for neurodegenerative and muscular disorders characterized by a significant contribution of lysosomal dysfunction.

Keywords: Acidification, Autophagosome, Autophagy, Endocytosis, Endosomes, Hydrolytic Enzymes, Lysosomal Dysfunction, Lysosomes, Lysosomal Membrane Proteins, Neurodegeneration, Phagocytosis, SNARE Proteins..

INTRODUCTION

Commonly known as suicide bags, they are cell organelles in Eukaryotic cells that help break down and recycle waste products from the cell. Lysosomes have hydrolytic enzymes that help break down biomolecules. Some of those hydrolytic

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enzymes include Proteases, Lipases, Nucleases, and Glycosidases. Due to these enzymes, the lysosome has an acidic pH of about 5, and the pH of the cytoplasm is neutral (7.2 to 7.8); the lysosome's membrane separates both. We currently understand lysosomes as a very varied group of organelles that vary in location, acidity, functions, and the hybrid organelles they create through fusion/fission and “kiss and run” processes. Endolysosomes, phagolysosomes, autolysosomes, and other terms are used to identify different lysosomal compartments. For example, endolysosomes are intermediate compartments formed during the fusion of endosomes and lysosomes, and they are typically acidic.

Phagolysosomes form when phagosomes fuse with lysosomes to degrade engulfed particles. Autolysosomes result from the fusion of autophagosomes with lysosomes during autophagy. There's also a distinction between acidic lysosomes (like endolysosomes, which actively degrade material) and non-acidic compartments (such as terminal lysosomes), which have completed their digestive function [1, 2].

Discovery of Lysosomes

During the 1950s, de Duve *et al.* from the University of Louvain in Belgium wanted to know how insulin acts in liver cells. They wanted to find where the glucose-6-phosphatase enzyme resides, which regulates blood sugar. They added rat liver debris to distilled water and centrifuged it. They found the enzyme with high activity, but when they purified the enzyme from the cell extracts, they could not precipitate it; however, they could not dissolve it again [3]. The strategy they used was a more advanced differential centrifugation that separates cell constituents based on size and density. They lysed the liver cells, fractionated them in a sucrose medium, and assayed glucose-6-phosphatase activity in the microsomal fraction. Differential centrifugation led them to the conclusion that the activity of the control enzyme, acid phosphatase, was only 10% of what it should have been. One of the scientists kept some of the fractioned samples in the fridge for five days (Fig. 1). On the other hand, when assaying the phosphatase activity, it was found to be equal to that which was expected. De Duve and his co-workers then hypothesized that a membrane could limit the access of an enzyme to its substrate, and resting the samples would give the enzymes time to become accessible. To these membrane-bound sacs containing acid phosphatase along with other lytic enzymes, de Duve gave the name “lysosomes” [4]. An acceptance of the discovery of lysosomes came when Christian de Duve was awarded the Nobel Prize in Physiology or Medicine in 1974. That same year, Alex Novikoff of the University of Vermont paid a visit to de Duve's laboratory. As an expert microscopist, Novikoff took the first electron micrographs of this newly discovered organelle from partially purified samples of the lysosome. The vital

role of lysosomes, for instance, was discovered by the work of Werner Strauss and his laboratory [2]. Strauss sought to determine how cells can draw molecules from outside their boundary wall through a so-called endocytosis process. He tagged proteins, followed their movement within cells, and found that protein fragments are engulfed in lysosomes. He concluded that lysosomes break down proteins.

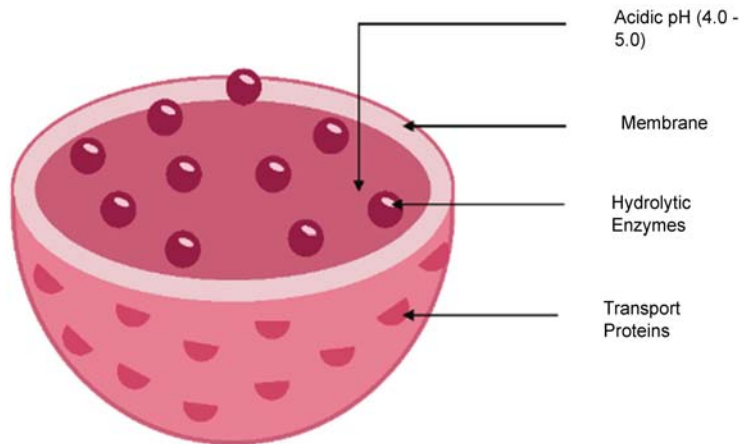


Fig. (1). Lysosome structure showing its membrane, digestive enzymes, and organelles involved in cellular degradation and recycling.

In other experiments, Zanvil Cohn incubated radiolabeled bacteria with macrophages and found that the different components of bacteria, including lipids, carbohydrates, and amino acids, were concentrated in lysosomes. Cohn deduced that lysosomes are the digestive organs of the cell, disposing of materials not produced by the cell itself and cellular products. Thus, lysosomes can be regarded as recycling stations that aid in cellular waste disposal and recycling of material ends [5].

Evolution of Lysosomes

Lysosomes are formed when vesicles from the Golgi Complex fuse with endosomes. Endocytosis is a process in which a section of the plasma membrane pinches off to form vesicles known as endosomes (Fig. 2) [5].

Vacuoles: The Storage Vaults of the Cell

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Abstract: Vacuoles are cell structures enclosed in membranes that play roles in various cellular functions within plant cells, some animal cells, and fungi cells. This section delves into the array of duties that vacuoles perform, like storing water and nutrients and aiding in maintaining turgor pressure, pH balance, and detoxification within cells. It also discusses the variations seen among organisms with a focus on specific vacuole types, including the central vacuole found in plants, the autophagic variety seen in animals, and the versatile fungal vacuoles. Furthermore, the chapter explores how vacuoles play a role in autophagy and defense mechanisms and the importance of vacuole-specific proteins such as Tonoplast Intrinsic Proteins (TIPs). It also covers progress in vacuole research, including their potential for use in drug delivery systems, enhancing plants' ability to withstand stress.

Keywords: Absciscic Acid, Autophagy, Cellular Detoxification, Glycerol, Fungal Vacuole, Ion Regulation, Membrane, Nutrients, Osmoregulation, Plant Vacuole, Tonoplast, Tonoplast Intrinsic Proteins (TIPs), Turgor Pressure, Vacuolar pH..

INTRODUCTION

Vacuoles are membrane-bound organelles found primarily in plant and fungal cells and some animal cells. In animal cells, vacuoles are relatively small, typically ranging from 50 to 500 nanometers in size. In contrast, plant cells contain a large central vacuole that can occupy between 30% and 90% of the cell's total volume, depending on the cell type and developmental stage. They act as storage compartments that can hold a variety of substances, including water, ions, nutrients, waste products, and pigments.

Vacuoles are essential for various cellular functions. They store water, nutrients, and waste products. Their ability to regulate water content helps maintain cell shape and rigidity, contributing to turgor pressure, the internal pressure exerted by

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the vacuole in/within the cell. Additionally, vacuoles are crucial for maintaining the pH of the cell's internal environment through proton pumps and ion exchange, ensuring optimal conditions for cellular processes like enzyme activity and metabolism. They also serve as detoxification centers, storing harmful substances. Furthermore, vacuoles are responsible for pigmentation by storing pigments like anthocyanins and carotenoids that provide color to flowers, fruits, and leaves. Lastly, they play a role in autophagy, a cellular process that involves breaking down cellular components.

The discovery of vacuoles dates back to the early days of microscopy. While Anton van Leeuwenhoek is often credited in the late 17th century for his observations of single-celled organisms, earlier scientists likely observed vacuoles using basic microscopes. As microscopy techniques advanced, scientists began to appreciate the diversity and significance of vacuoles in different cell types. While the term “vacuole” comes from the Latin word “vacuoles,” which means “empty,” the organelle is not truly empty space but instead contains a variety of substances. Research on vacuoles remains a vital area in cell biology, with scientists investigating their roles in plant growth, development, and responses to stress.

Types of Vacuoles

Plant Vacuoles

In plant cells, the central vacuole is the most prominent type of vacuole. This large, membrane-bound organelle takes up a significant portion of the cell's volume. The central vacuole is surrounded by a membrane called tonoplast, which is also known as the vacuolar membrane. The central vacuole serves various functions, including:

- **Storage:** It stores water, nutrients, ions, toxins, and waste products.
- **Turgor pressure:** The pressure exerted by the fluid inside the vacuole maintains the cell's rigidity and the cell's structure.
- **pH regulation:** It regulates the pH of the cell's internal environment.
- **Detoxification:** It stores harmful substances.
- **Pigmentation:** It contains pigments that provide colour to flowers, fruits, and leaves.
- **Autophagy:** It breaks down cellular components.

There are primarily two types of plant vacuoles.

- **Lytic Vacuole (LV):** LVs are developed early during embryogenesis. These vacuoles help in breaking down unwanted cellular substances using hydrolase

enzyme. Therefore, hydrolase enzyme are very crucial in the degradation and recycling of cellular substances [1].

- **Protein Storage Vacuole (PSV):** PSVs are formed after LVs during embryogenesis. In addition, in maturing seeds, LVs are transformed to PSVs, but during germination, this process is reversed. These vacuoles are used to store nutrients and proteins, which may be needed at different stages of growth or under stress conditions. It's possible for protein storage vacuoles and lytic vacuoles to combine to form a large central vacuole [1].

Animal Vacuoles

Animal cells typically lack the large, central vacuoles that are characteristic of plant cells. Nevertheless, they possess various smaller vacuoles that fulfill specific roles. These vacuoles are generally less prominent than the central vacuoles found in plants.

Different Types of Vacuoles Present in Animals

Food Vacuoles - These are membrane-bound structures that form when cells take in food particles through a process called phagocytosis. They merge with lysosomes, which contain enzymes for digestion, to break down the ingested food.

Autophagic Vacuoles - These are created when cells need to recycle their own materials. They engulf damaged organelles or cellular waste and transport them to lysosomes for breakdown.

Secretory Vacuoles - These vacuoles are responsible for storing and releasing substances like hormones or neurotransmitters.

Contractile Vacuoles - Although more commonly found in protists, these vacuoles can sometimes be present in animal cells, particularly those in hypotonic environments. They play a role in osmoregulation by maintaining the cell's water balance and expelling excess water.

Fungal Vacuoles

Fungal vacuoles are organelles that carry out a variety of essential functions for the growth, differentiation, symbiosis, and pathogenesis of fungi. These acidic storage compartments are involved in protein degradation, maintaining cellular homeostasis, membrane trafficking, signalling processes, and nutrient storage. They play a role in glycoprotein turnover and hydrolysis, store phosphate, calcium, and amino acids, and help regulate pH, ion homeostasis, osmotic balance, and cytoplasmic detoxification. Additionally, they are crucial for

CHAPTER 9

Plasma Membrane: Gateway and Sentinel of Cellular Exchange

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Abstract: The plasma membrane is a critical cellular structure that acts as a selective barrier, controlling the movement of molecules in and out of the cell while enabling communication with the external environment. This chapter provides an in-depth exploration of the plasma membrane, beginning with its structure and the fluid-mosaic model, which describes the dynamic organization of lipids, proteins, and carbohydrates. It then examines the various functions of the membrane, including maintaining cellular integrity, supporting signal transduction, and regulating the movement of substances through mechanisms like passive diffusion, facilitated diffusion, active transport, and group translocation. Key transport proteins such as glucose transporters and ion channels are discussed in detail, highlighting their roles in maintaining cellular homeostasis. The chapter also addresses the pathological significance of plasma membrane dysfunction, linking abnormalities to diseases such as hypertension, sphingolipid-related disorders, and neuronal vulnerabilities. Further, it explores the role of the plasma membrane in neurotrophic signaling and insulin regulation. Finally, recent advances in therapeutic approaches, including CAR-T cell therapy and liposomal drug delivery systems, are examined for their potential in disease treatment.

Keywords: Active Transport, Aquaporin, Bilayer, Diffusion, Endocytosis, Exocytosis, Facilitated Diffusion, Insulin, Lipid Raft, Membrane Potential, Neurotransmitter, Phospholipid, Potassium Channel, Sphingolipidoses..

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INTRODUCTION

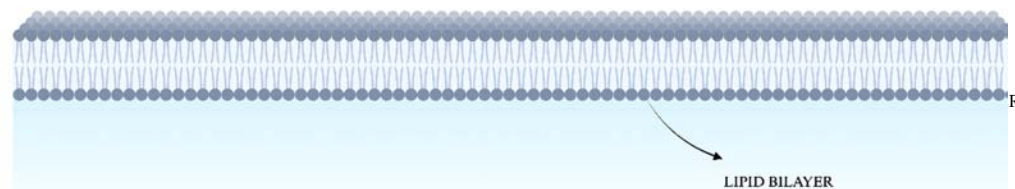
The plasma membrane is a lipid bilayer with embedded proteins that control the movement of substances in and out of the cell while enabling interactions. Organellar membranes have a similar structure but differ in composition and function based on the organelle's role. Membrane-bound organelles include the mitochondria, endoplasmic reticulum, Golgi apparatus, lysosomes, and peroxisomes. Non-membranous organelles include ribosomes, centrioles, cilia, and flagella [1].

STRUCTURE

Gorter and Grendel Membrane Theory (1920)

This theory proposes that the plasma membrane is made up of a lipid bilayer. An experimental investigation on erythrocytes from different sources was carried out, and the experimental results supported their hypothesis. Gorter and Grendel failed to explain the membrane functions and could not explain the other membrane components other than lipids. This model set the stage for later models by highlighting the importance of lipid organization (Fig. 1) [2].

GORTER AND GRENDL MEMBRANE MODEL



DAVSON-DANIELLI MODEL

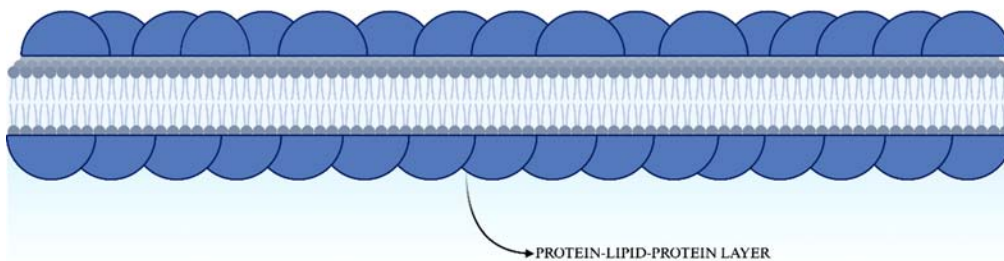


Fig. (1). Depiction of the difference between the gorter and Grendel membrane model and the Davson-Danielli model.

Pauci-molecular Theory

In 1935, Davson and Danielli came up with a model for plasma membrane called the Pauci-molecular theory. According to this theory, the membranes have a central “lipoid” region, which in general is surrounded by monolayers of lipids and is covered by protein sheets. This model is like a sandwich model made of protein-lipid-protein. In the year 1959, J. David Robertson proposed that all the cellular membranes possess a similar underlying structure, the unit membrane. He used extensive metal staining for the visibility of plasma membranes under an electron microscope. When observed under a microscope, the cell membrane exhibits a trilaminar appearance, featuring a lighter inner region and two darker outer regions. Robertson noted that the membrane consists of a lipid bilayer with sheets of mucoproteins on both sides, which supports the pauci-molecular theory [3]. The pauci-molecular theory proposed that all the membranes have equivalent structure, the same thickness, and the same lipid-protein ratio in the membrane surface. The Pauci-molecular theory couldn't explain the presence of membrane proteins outside the membrane or the active transport of macromolecules. It also failed to account for the dynamic nature and heterogeneity of membrane structures. Until 1972, the most accepted plasma membrane model was the pauci-molecular theory. After 1972, the scenario completely changed [4].

Fluid Mosaic Model (1972)

S. Jonathan Singer and Garth Nicolson proposed the fluid-mosaic model. This theory says that the plasma membrane is made up of a lipid bilayer where proteins are embedded in it. The phospholipid bilayer illustrates the effects of hydrophilic and hydrophobic interactions in lipids. The non-polar fatty acid chains of the phospholipid stand away from being in contact with water by increasing the hydrophobic interaction. The ionic groups are in direct contact with the aqueous phase, thereby giving rise to hydrophilic interactions. The hydrophilic and hydrophobic interactions are crucial for maintaining the membrane's integrity and functionality, as they help create a stable barrier that controls the movement of substances in and out of the cell. The dipole-dipole interaction between ionic pairs of phosphatidylcholines (zwitterionic phospholipid) is the reason for the stable lipid bilayer structure. According to this theory, there are two types of membrane proteins-peripheral and integral (Fig. 2).

CHAPTER 10

Mitochondria and Chloroplasts: Evolutionary Engines of the Cell

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Abstract: Mitochondria and chloroplasts are essential organelles driving energy production and metabolism in eukaryotic cells. Mitochondria, known as the cell's "powerhouses," generate ATP through oxidative phosphorylation, utilizing nutrients to fuel cellular processes. According to the endosymbiotic theory, mitochondria originated from free-living bacteria, and their distinct mitochondrial DNA (mtDNA) is maternally inherited. Dysfunction in mitochondria—often due to accumulation of mtDNA mutations—is associated with neurodegenerative diseases, metabolic syndromes, and cancer, with recent studies revealing their roles in aging and disease. These insights are advancing therapies to improve mitochondrial health, and genome editing now holds the potential for correcting pathogenic mtDNA mutations. Chloroplasts in plant cells are essential for photosynthesis, synthesizing carbohydrates, amino acids, fatty acids, and membrane lipids. Chloroplasts are sensitive to temperature stress, and their malfunction can produce Reactive Oxygen Species (ROS), which may be fatal to plant cells. This sensitivity makes chloroplast health crucial for plant resilience, especially under climate stress. Future research in mitochondrial and chloroplast should concentrate on offering potential treatments for human diseases and developing stress-resistant crops. Genome editing technologies could address mitochondrial and chloroplast dysfunction, creating innovative therapies and sustainable agricultural practices to address health and environmental challenges.

Keywords: Cellular Respiration, Endosymbiotic Theory, Energy Production, Electron Transport Chain, Krebs Cycle, Mitochondria-Associated Membranes (MAMs), Mitochondrial DNA (mtDNA), Oxidative Phosphorylation, Reactive Oxygen Species (ROS), Uniparental Inheritance..

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MITOCHONDRIA

Introduction

It was close to a billion years since cellular respiration had become a vital part of the cell when science discovered the pan portion of the cell during the middle of the nineteenth century. Based on light microscopy, the first diagrams of cellular structure showed the existence of granular structures known as mitochondria in the cell's cytoplasm. The first observation of the presence of mitochondria in muscle cells was made during the 1850s by the anatomist Kölliker (1856). However, in the 1890s, Altman devised a fantastic proposal: the granules, which Deichmann called “bioclasts,” were symbiotes. This idea was expanded on by Mereschkowsky in 1905.

The same idea concerning chloroplasts was first proposed by C. Simons in 1883, before Schimper. The term “Mitochondrion” was first used to explain data by Benda in 1898. The name is derived from the Greek terms: “mitos” in thread, referring to an activity in the mitotic field, and “chondros” in grain, referring to an activity associated with spermatogenesis. As mitochondria are the primary site of energy conversion in the cell, or where “cellular respiration” takes place, they are also known as the organelles that produce blood inside the cell. However, the causes of such a phenomenon were not studied until the 1950s, before the genesis of mitochondria was considered. Very early in the 1950s, Mitchell (1952) and Ephrussi (1950) reported post-Mendelian genetic variation to control the forces determining the division of yeast mitochondria. A little while later, McLean *et al.* (1958) reported that mitochondria can synthesize polypeptides. Luck and Reich (1964), Nass, early in the nineteen sixties, since many groups. Their significant contributions regarding the mitochondria and their invention of the mitochondrion were related. However, the hypothesis on the endosymbiotic origin of mitochondria, which had existed, started to attract new and enthusiastic followers in the 1960s with the introduction of new biochemical methods. It would be superfluous to emphasize that the discovery of the DNA contained within the organelles and the non-classical inheritance of organelles were the two components that contributed to the resurgence of interest.

Once Margulis had rephrased the endosymbiotic theory into endosymbiosis (1970), it became recognized how mitochondria have evolved into the cell. It is accepted that the mitochondrion originated from a free-living bacterium. However, there is still an argument regarding how this endosymbiosis came about and which living things this process encompassed. There are about a hundred years between the first reports of mitochondria and the modern understanding of the structure, function, inheritance, and origin of mitochondria. Since 2003, the

work of Tsang and Lemire has made it possible to identify the mitochondrial genomes of more than 250 species by examining the effects of mutations in several mitochondrial genes.

Origin and Evolution of Mitochondria

Mitochondria in eukaryotes are suspected of having an endosymbiotic origin. In the 1960s, Margulis Marshaled confirmed this hypothesis based on two factors, *i.e.*, the presence of DNA and a separate translational system distinct from one in the cytosol. Further elaborate studies of mitochondrial genomes and genes proclaim that mitochondria have a bacterial origin. However, another theory explained the mitochondrial origin, *i.e.*, the hydrogen hypothesis given by Martin and Muller in 1998. According to this, the origin of heterotrophic organelles and other eukaryotes was identical.

The symbiotic origin is based on two subjects: the “Archezoan Scenario” and the “Symbiogenesis Scenario.” The former is based on endosymbiosis and the latter on the hydrogen hypothesis. The archean scenario explains the presence of a proto-mitochondrial endosymbiont in mitochondrial eukaryotes, and the symbiogenesis scenario explains the hold of an alpha proteobacterium in an archaeobacterial cell [1, 2].

Mitochondrial Inheritance in Eukaryotes

The mitochondrion is usually inherited from a single parent (maternal inheritance). This is responsible for the offspring’s homoplasmy (where all mtDNA are genetically identical)—the disruption of MT homoplasmy results in mitochondrial dysfunction and cancer in human beings. There are two methods for ensuring mtDNA homoplasmy: (i) Strict Maternal Inheritance (SMI) and (ii) the bottleneck mechanism. A strict maternal inheritance pattern is observed in the shell coiling of snails (*Limnaea stagnalis*). The three stages that control the bottleneck mechanism are:

- (i) A bottleneck that occurs during oogenesis expresses mutations that allow purification before the resumption of replication during the oocyte's maturity, lowering the copy number of mtgenes.
- (ii) Selective replication during oogenesis led to the establishment of a genetic bottleneck.
- (iii) Another one forms during embryogenesis.

Recent reviews provide a glimpse into the status of maternal inheritance research (reviews are available in Jokinen and Battersby 2013 and Mishra and Chan 2014).

CHAPTER 11

Cytoskeleton: The Cell's Backbone and Highway

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Abstract: The cytoskeleton is an essential framework in eukaryotic cells, providing structural support, maintaining shape, and facilitating intracellular transport and movement. This chapter explores the key components of the cytoskeleton: microfilaments, intermediate filaments, and microtubules, discussing their structures, assembly mechanisms, and roles in cellular processes. Microfilaments, composed of actin, contribute to cellular shape, movement, and muscle contraction through mechanisms like treadmilling and interaction with myosin. Intermediate filaments, including keratin and vimentin, provide mechanical strength and support cellular integrity under stress. Microtubules, composed of tubulin, are involved in mitosis, intracellular transport, and the maintenance of cell polarity. Additionally, this chapter delves into the role of motor proteins like kinesin and dynein in facilitating molecular transport along microtubules and how cytoskeletal dynamics are crucial for both healthy cellular function and pathological conditions like cancer and neurodegenerative diseases.

Keywords: Actin, Cancer, Cell Motility, Dynein, Intermediate filaments, Keratin, Kinesin, Locomotion, Microfilaments, Microtubules, Muscle, Neurodegenerative Diseases.

INTRODUCTION

In addition to the cytoplasm, plasma membrane, and organelles, eukaryotic cells have something like a hidden superhero called the cytoskeleton. This internal framework comprises tiny fiber proteins, which we call cytoskeletal fibers. The cytoskeletal fibers play a critical role in maintaining cellular integrity by preserving cell shape and providing mechanical support. The plasma membrane functions as a selective barrier that protects the intracellular environment. Within

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the cytoplasm, distinct organelles carry out specialized biochemical processes: mitochondria facilitate ATP production through cellular respiration, the Golgi apparatus is responsible for protein modification and trafficking, and the endoplasmic reticulum is involved in protein synthesis and intracellular storage. Now, let us talk about the cytoskeleton. Think of it as the house's skeleton. Just like our bones give our bodies structure, these cytoskeletal fibers give the cell its shape and help it stay strong. They're like the support beams in our house. But the cytoskeleton does more than that. It's like the cell's personal trainer. It helps the cell move around, divide into two when it needs to make a copy of itself, and even helps things move around inside the cell. So, in simple terms, the cytoskeleton is like the cell's own hero, working behind the scenes to keep everything in order, maintain shape, and ensure the cell can do everything it needs to do [1].

Types of Cytoskeletal Fibers

The cytoskeleton is a complex network of fibers that provides structural support, shape, and organization to cells. It plays a critical role in various cellular processes, including movement, division, and intracellular transport. This dynamic system is made up of different types of protein-based filaments, each contributing to specific cellular functions (Table 1).

Table 1. Types of protein present in the cytoskeleton, along with the protein size.

Type	Protein	Site	Size (kDa)
I	Acidic Keratins	Epithelial Cells	40-60
II	Neutral or Basic Keratins	Epithelial cells	50-70
III	Vimentin	Fibroblasts, white blood cells, and other cell types	54
	Desmin	Muscle cells	53
	Glial fibrillary acidic proteins	Glial cells	51
	Peripherin	Peripheral neurons	51
IV	Neurofilament proteins		67
	NF-L		150
	NF-M	Neurons	200
	NF-H	Stem cells	66
	α -Nexin		200
	Nestin		
V	Nuclear lamins	Nuclear lamina of all cell types	60-75

The three types are:

- Microfilaments
- Intermediate filaments
- Microtubules

MICROFILAMENTS

Structures that were a part of the cytoskeleton that supported the flexibility and increased the strength of the cell were found in 1974 by B. Paleviz *et al.* They are microfilaments, as we now refer to them.

Structure

Microfilaments are little rod-shaped structures approximately 6 nm in diameter, found inside the cytoskeleton of eukaryotic cells. Microfilaments are constituted by the aggregation of actin monomers. It comprises two strands of actin protein subunits coiled in a helical formation, hence referred to as actin filaments. The actin subunits that constitute microfilaments are globular actin (G-actin), which, upon polymerization, are referred to as filamentous actin (F-actin). The actin subunits assemble directionally to produce microfilaments. Subunits possess a “top” and a “bottom,” with the top of one subunit consistently interacting with the bottom of another. The terminal subunit of a filament is referred to as the negative (-) end, while the opposing end, which experiences greater polymerization, is designated as the plus (+) end. Microfilaments possess polarity, although this pertains solely to their directionality and is unrelated to electrical charge [2] (Fig. 1).

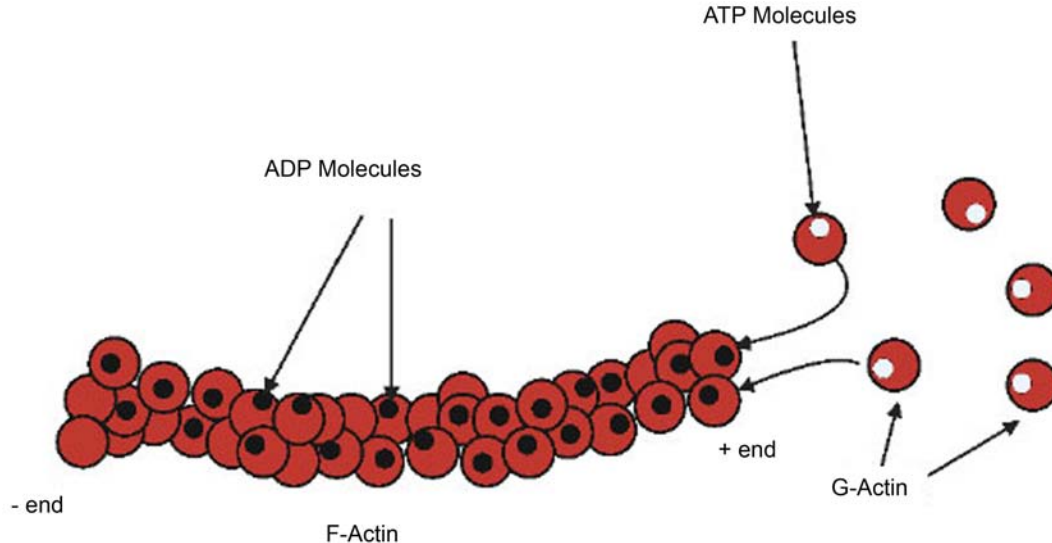


Fig. (1). Structure and assembly of microfilaments.

CHAPTER 12

Signal Transduction Pathways Orchestrate Cellular Communication: A Narrative Review

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Abstract: Cell signaling and signal transduction coordinate cellular communication and the execution of diverse biological functions. This review synthesizes how cells sense and integrate external and internal cues through core pathways, with emphasis on G Protein–Coupled Receptor (GPCR) and Mitogen-Activated Protein Kinase (MAPK) cascades that govern growth, differentiation, and apoptosis. We outline the roles of second messengers, protein kinases, and transcription factors in propagating and tuning signals, and describe regulatory mechanisms that maintain fidelity and specificity. We also summarize how pathway dysregulation underlies disease, including cancer, and discuss the therapeutic implications of targeting signaling networks.

Keywords: Cell signaling, Signal transduction, GPCR, MAPK, Second messengers, Protein kinases, Transcription factors, Apoptosis.

INTRODUCTION

Cell signaling enables cells to communicate with each other and with their environment to sustain physiological function. This communication network is essential for organismal growth, adaptation, and survival in dynamic contexts [1]. Signal transduction converts extracellular and intracellular cues into specific

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cellular responses, ensuring appropriate reactions to stimuli associated with immune defense, growth, differentiation, and other core processes [2].

Signals include hormones, growth factors, neurotransmitters, and environmental inputs such as temperature and light [3]. Cells operate within integrated networks that support organismal health. Cell signaling coordinates growth and development, maintains homeostasis, mediates immune responses, and ensures proper nervous system function [4].

Multicellular organisms develop from a single fertilized egg through cell division, differentiation, and controlled tissue formation, all tightly regulated by signaling pathways [5]. During embryogenesis, signaling molecules guide cell positioning and organogenesis by activating gene programs for specialization [6]. To preserve a stable internal environment, cells continuously sense changes in temperature, nutrient levels, and pH and adjust their activity to restore equilibrium [7]. Hormones such as insulin and glucagon exemplify signaling molecules that regulate metabolism and energy balance [8].

The immune system relies on signaling to detect and eliminate pathogens. Immune cells such as T cells and B cells use signaling to recognize invaders, activate other immune cells, and coordinate an effective defense through cell–cell communication [9]. Neurons communicate *via* electrical and chemical signals to transmit information across extensive networks; neurotransmitters enable rapid, coordinated responses underlying movement, sensation, and cognition [10].

TYPES OF CELL SIGNALING

Cell signaling employs distinct modes of communication defined by distance and delivery. The four principal forms are autocrine, paracrine, endocrine, and juxtacrine signaling.

Autocrine Signaling

In autocrine signaling, a cell releases signaling molecules that bind receptors on its own surface. This feedback modulates the cell's activity and is especially important in development and immune responses [11]. Cancer cells often exploit autocrine loops to drive their growth and survival.

Paracrine Signaling

In paracrine signaling, cells secrete molecules that act on nearby cells, mediating local communication [12]. During tissue development and repair, growth factors stimulate adjacent cells to proliferate and divide; paracrine cues similarly coordinate wound healing [13].

Endocrine Signaling

Endocrine signaling involves hormone release into the bloodstream, allowing signals to reach distant targets. This long-range communication coordinates metabolism, development, and reproduction across organ systems; endocrine glands such as the pituitary, thyroid, and pancreas orchestrate these responses [14].

Juxtacrine Signaling

In juxtacrine signaling, cells communicate through direct physical contact. Membrane-bound ligands on one cell engage receptors on an adjacent cell, triggering local responses. During embryonic development and tissue differentiation, where precise cell-to-cell interactions build ordered structures, juxtacrine signaling is essential [15].

Role of Signal Transduction in Cellular Communication

After a signal reaches its target cell, it must be converted into a specific intracellular response—a process termed signal transduction. Signal transduction comprises sequential molecular events that amplify and relay information from the cell surface to the intracellular machinery that produces the appropriate response. Major components include:

Receptors

Receptors reside on the cell surface or within the cell and bind specific signaling molecules; they are the first points of contact for external cues [16]. Two principal classes of cell-surface receptors mediate signal transduction: G Protein-Coupled Receptors (GPCRs) and Receptor Tyrosine Kinases (RTKs) [17]. Ligand binding induces a conformational change that initiates the intracellular signaling cascade [18].

Signaling Molecules

Signaling molecules—such as hormones, growth factors, and neurotransmitters—are released by a cell and bind receptors on target cells to initiate transduction [19]. Insulin, for example, binds its receptor to regulate glucose uptake and metabolism [20].

Signal Transduction Proteins

Once receptors are activated, intracellular signaling proteins propagate and amplify the signal, often through cascades in which one protein activates the next

CHAPTER 13

Cell Death: Mechanisms and Mysteries beyond Apoptosis

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Abstract: This book chapter on cell death explores the mechanisms, significance, and implications of various forms of cellular demise in health and disease. It delves into classic pathways like apoptosis, which ensures cellular homeostasis, immunity, and necrosis, traditionally viewed as accidental cell death but now recognized for its role in inflammation and tissue damage. Emerging forms of cell death, such as autophagy, pyroptosis, ferroptosis, and NETosis, highlight the complexity of cellular life and death decisions. The chapter underscores how each form of death is tightly regulated by specific signaling pathways and proteins, contributing to tissue development, immune responses, and the progression of diseases like cancer and neurodegenerative disorders. A particular focus is placed on the molecular crosstalk between these pathways and their potential as therapeutic targets. The historical context of cell death research—from identifying apoptosis to modern-day advancements in targeted therapies and gene-editing technologies—provides a comprehensive view of how our understanding of cellular death has evolved.

Keywords: Apoptosis, Autophagy, Cancer, Disorders, Ferroptosis, Immune responses, Necrosis, NETosis, Neurodegenerative, Pyroptosis, Signalling pathways, Therapeutic targets.

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INTRODUCTION

In the intricate tapestry of life, cell death stands as a cornerstone, shaping the very essence of biological existence. It is a process as ancient as life itself, a fundamental mechanism intricately woven into the fabric of development, homeostasis, and disease. Cell death is not a mere cessation of cellular activity; instead, it is a meticulously regulated, multifaceted series of events that culminate in the demise of a living entity. Beyond its apparent simplicity lies a world of complexity, where cellular components intricately communicate and orchestrate a symphony of self-destruction. The study of cell death has transcended the realms of mere biological inquiry; it has become a frontier where life sciences, genetics, and medicine converge. At its core, cell death represents nature's ultimate balancing act. It is the balance of nature, which calls for everything that is ever born to meet its end, its death. Every human, every microbe has a certain life span, after which death is inevitable. This concept, at the most cellular level, is more frequent and much more complex. There are countless cells in our body dying every day, and countless still that are newly generated. It is a process of profound importance, eliminating cells that have outlived their usefulness, have suffered irreparable damage, or have succumbed to genetic aberrations. This process is indispensable in sculpting and developing organisms and ensuring the proper formation of tissues, organs, and intricate structures [1]. Additionally, cell death acts as a guardian, protecting the organism from potential threats. It is the body's natural response to infections, injuries, and internal dysregulation that underlie various diseases, preventing the spread of damage to neighbouring cells and tissues. Yet, the significance of cell death extends far beyond its role as a biological safeguard. It is a process deeply entwined with the very essence of evolution. The ability to control cell death has played a pivotal role in the emergence of complex life forms, allowing for the development of intricate biological systems. In essence, the study of cell death is a study of life itself—a journey into the heart of existence, exploring the delicate balance between creation and destruction, growth and decay [2, 3].

In this chapter, we embark on a comprehensive exploration of cell death. We delve into the mechanisms that govern this intricate process, uncovering the molecular intricacies that define its various forms. From the classic pathways of apoptosis and necrosis to the emerging realms of autophagy, pyroptosis, and ferroptosis, we unravel the mysteries that underlie each type of cellular demise. Moreover, we examine the physiological significance of cell death, its pivotal role in development, tissue maintenance, and immunity, as well as its implications in diseases ranging from cancer to neurodegenerative disorders. As we venture deeper into the labyrinth of cell death, we also cast our gaze towards the future. Recent advancements in the field have opened new avenues of research,

promising innovative therapies and treatments. From targeted therapies tailored to specific cell death pathways to the revolutionary potentials of gene-editing technologies, the landscape of cell death research is constantly evolving [4].

The Historical Tapestry of Cell Death Research

The annals of cell death research are rich with the endeavours of pioneering scientists whose meticulous observations and ingenious experiments have illuminated the enigmatic pathways of cellular demise. Our journey through time takes us back to the 19th century, when rudimentary microscopes provided the first glimpses into the microscopic world. Eminent scientists like Rudolf Virchow and Carl Vogt observed cellular structures and noted instances where cells seemed to disintegrate, laying the foundation for the intriguing field of cell death studies. In the early 20th century, the term “apoptosis” made its debut in scientific discourse, thanks to the work of Walther Flemming. This term, embodying the Greek notion of a “falling off” or “dropping,” encapsulated the elegant yet complex process of programmed cell death. Flemming's observations sparked curiosity, fuelling a wave of research that sought to unravel the mysteries of apoptosis [1]. Amidst these explorations, the mid-20th century heralded significant breakthroughs. Pioneering electron microscopy techniques provided unprecedented resolution, enabling scientists to delve deeper into cellular structures and observe the orchestrated disassembly of dying cells. It was during this period that the morphological hallmarks of apoptosis—cell shrinkage, membrane blebbing, chromatin condensation, and formation of apoptotic bodies—were meticulously documented, offering a profound understanding of this fundamental process (Table 1) [5].

In the latter half of the 20th century, the scientific community witnessed the emergence of groundbreaking discoveries. A key milestone came in the form of the identification of caspases, the enzymatic executioners orchestrating apoptosis. The elucidation of these proteases' role in cleaving vital cellular substrates offered a molecular perspective, transforming apoptosis from a mere morphological phenomenon into a finely regulated biochemical cascade. As the 21st century dawned, the exploration of cell death expanded beyond apoptosis. Necrosis, once deemed a chaotic and uncontrolled form of cell death, was redefined. It became evident that necrosis, too, had regulatory elements, especially in the context of ischemic injury and inflammation. Concurrently, autophagy, a cellular recycling mechanism, garnered increasing attention. Its role in maintaining cellular homeostasis, particularly during periods of stress and starvation, unfolded as researchers decoded the intricate machinery of autophagosome formation and cargo recognition. The modern era of cell death research is marked by a profound transformation catalysed by advances in molecular biology, genetics, and high-

CHAPTER 14

Stem Cells: Breakthroughs in Medicine and Therapeutics

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Abstract: Stem cells are usually referred to as undifferentiated cell masses, which can differentiate into many cell types. Stem cells are present in all organisms in various evolved forms. These cells can self-renew. Additionally, stem cells can be broadly classified based on their origin and differentiation potential. Thus, the therapeutic potential of these cells can be harnessed and utilized to treat various disorders, including blood disorders, tissue and organ regeneration, and others, which can be classified under the division of stem cell therapy. Although stem cells possess the capacity for use in disease treatment, their implementation also has its own risks. This chapter covers the basics of stem cells, including the associated cell types, applications, epigenetics, and recent trends.

Keywords: Blood disorders, Differentiation potential, Epigenetics, Recent trends, Stem cell therapy, Stem cells, Tissue regeneration.

INTRODUCTION

Stem cells are defined as undifferentiated masses of cells present in all living organisms. Furthermore, stem cells represent the building blocks of all tissues and organs, with major properties including:

- i. Extensive proliferation
- ii. An ability to arise from a single cell (clonal).
- iii. An ability to differentiate to form different cell types (potency).

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However, these properties vary among the different cell types. For instance, Embryonic Stem Cells (ESCs) undergo extensive proliferation, whereas Adult Stem Cells (ASCs) primarily differentiate into tissue-specific cells.

Thus, stem cells are currently employed in cell therapy and drug development processes. Additionally, stem cells have improved understanding of pathogenesis. Nonetheless, despite these significant advances in stem cell biology, certain ethical issues remain that limit the utilization of these cells [1].

STEM CELL FUNCTIONS

Stem cells are predominantly used due to their ability to self-renew and differentiate into various types of cells. In adult organisms, stem cells can either actively replenish tissues to repair damage or remain dormant, as is the case with neural stem cells in the mammalian brain. There are two types of stem cell division: Symmetric and asymmetric. Symmetric division occurs when the cells reproduce to produce more identical cells, while asymmetric division occurs when the cell differentiates to form a specific cell type. In the asymmetric model, the stem cell forms an exact copy and one differentiated cell, which helps maintain the homeostasis of the stem cell pool at the individual level. A major disadvantage of asymmetric division is that the stem cell pool cannot be replenished at the time of injury. However, this limitation can be resolved by the symmetric division, which maintains the homeostasis of the stem cell pool at the population level. Meanwhile, two types of symmetric division exist: the first is a proliferation division, which leads to the formation of two stem cells; the second is a differentiation division, which leads to the formation of two differentiated cells. Proliferation and differentiation of these cells occur due to the signals from the surrounding tissues as well as the stem cell [2, 3].

TYPES OF STEM CELLS

Stem cells are classified based on the following factors:

- i. Differentiation potential
- ii. Origin

Stem Cells Based on Differentiation Potential

The ability of stem cells to differentiate is based on their origin and associated derivation; thus, stem cells can be divided into five major categories.

Totipotent/Omnipotent Stem Cells

These are the cells that have the potential to differentiate into the various cell types found in the entire organism. Totipotent cells have the highest potential to differentiate. For example, the zygote is a totipotent cell that can differentiate into all cell types, including the three germ layers and extraembryonic tissues, such as the placenta. Meanwhile, totipotent cells found in an embryo differentiate into the inner cell mass (ICM) and the extra-embryonic cell lineage, such as the trophoblast. These totipotent embryos exhibit a unique transcriptome, characterized by dramatic changes in epigenetic and chromatin features, including the *de novo* assembly of nucleosomes, demethylation of DNA, chromatin remodeling, and histone modifications.

Pluripotent Stem Cells

These stem cells play a crucial role in forming the three primary germ layers, which, in turn, help develop all tissues and organs: ectoderm, mesoderm, and endoderm. These stem cells are only short-lived during the early stages of embryo development; however, pluripotent stem cells can develop into either fetal cells or adult cells, but not into a complete organism. However, these cell lines can be immortalized and grown *in vitro*. These cells can then differentiate into all cell types through various mechanisms, similar to those employed by ESCs.

Multipotent Stem Cells

Meanwhile, Multipotent Stem Cells (MSCs) have a more restricted differentiation potential, transitioning to specific cell types within a particular lineage. For example, Hematopoietic Stem Cells (HSCs) can differentiate into different types of blood cells. These cells are considered ASCs due to their limited capacity for differentiation and multipotency. MSCs are found in various organs, including the periosteum, adipose tissue, placenta, dermis, and umbilical cord blood. MSCs can be identified by their adherence and the ability to differentiate into cartilage, fat, and bone, as well as the expression of markers, such as CD90, CD105, and CD73. Currently, these cells are being used in tissue healing, development, and defense.

Oligopotent Stem Cells

These cells possess the ability for self-renewal and can differentiate into various types of cells. Moreover, these stem cells can be viewed as progenitor cells for ASCs and can differentiate into specific cell types; for example, myeloid stem cells can differentiate into white blood cells but not red blood cells. Furthermore, these progenitor-oligo cells have also been observed to differentiate into more mature cells, such as alveolar epithelial cells and bronchiolar epithelium.

CHAPTER 15

Cancer Stem Cells: Catalysts of Cancer Progression

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Abstract: Within tumors, a small number of cells exist alongside the majority of more rapidly dividing cells that constitute most of the tumor. This slowly growing subset of cells is referred to as cancer stem cells (CSCs), and they can regenerate themselves, allowing for the continuous growth of the tumor. CSCs were first reported in acute myeloid leukemia by Bonnet and Dick in 1997 and have been identified in other forms of cancer, including the brain, breast, lung, and liver tumors. These cells also show a striking resemblance to normal stem cells, for instance, in their ability to give rise to both the progenitor and malignant cells via asymmetric mitosis. These cells act on their own and play a significant role in the progression of tumors, inducing metastatic foci and bearing chemoresistance and radiotherapy resistance to standard treatments. This resistance can be attributed to several mechanisms, including active DNA repair, non-cycling state, and efflux of the drugs. They are components of the immune system that allow them to interact with other immune cells, thus escaping any immune responses while at the same time aiding in the recurrence of the tumors. Many crucial signaling pathways in the body, such as Wnt/ β -catenin, Notch, and Hedgehog, control neural stem cells' actions. New therapies have been integrated against cancer aimed at cancer stem cells, such as specific marker blocking, miRNA/LncRNA therapy, and immunotherapy. There is also great potential in new strategies, such as nanotechnology-based targeting of cancer stem cells to reduce the chances of tumor recurrence. Management techniques seek to eradicate the cancer stem cell population to mitigate the chances of recurrence and enhance treatment success.

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Keywords: ALDH, Asymmetric division, Cancer heterogeneity, Cancer stem cells, Chemoresistance, Differentiation, DNA repair, Epigenetics, Epithelial-mesenchymal transition, Hedgehog pathway, Immunotherapy, Markers, Metastasis, Microenvironment, Notch pathway, Proliferation, Quiescence, Self-renewal, Tumor growth, Tumor microenvironment.

INTRODUCTION TO CANCER STEM CELLS

Cancer stem cells are small subunits of cells formed within a solid tumor that are capable of self-renewal, leading to continuous tumor growth. They are similar to normal stem cells and can divide to form undifferentiated masses of cells. The mitosis of a cancer stem cell is asymmetric and can give rise to both progenitor and malignant cells. The identification and characterization of these cells were first done by Bonnet and Dick (1997) in acute myeloid leukemia (AML) in CD34⁺ CD38⁻ sets; later, cancer stem cells were also spotted in other types of tumors in the brain, breasts, lung, and liver. These stem cells, which are originally undifferentiated cells, would differentiate for the renewal of cells or tumorigenesis.

History and Evolution of Cancer Stem Cells

1997: Scientists Bonnet and John Dick identified masses of cells that would divide uncontrollably in leukemia when those cells were put in mice. They would then provide evidence that supported the hypothesis of the formation of cancer stem cells.

2000: Cancer stem cells were studied in a broader spectrum, and their existence was attributed to solid tumors that would be present in the brain, breast, and colon cancer. The solid tumors were identified when the cell samples were tested with molecular/surface markers and functional assays.

2010: Later, around 2010, researchers began looking in depth for the functional understanding of cancer stem cells. They focused on differentiating cancer stem cells from normal stem cells and their interactions with other cells, the pathways associated with the cells, and how to treat cancers that are represented by cancer stem cells.

Current Concerns and Challenges

With ongoing research on cancer stem cells, researchers have developed target therapies that work under the mechanism of targeting the specific cancer stem cell and working to eliminate them. However, cancer stem cells are very complex

structures and are heterogeneous in nature. Different cancers have different types of cancer stem cells, and to use target technology to destroy them is extremely complicated; hence, newer and more effective approaches have been developed to destroy these cells in cancerous tumors.

CHARACTERISTICS OF CANCER STEM CELLS

Cancer stem cells can be caused by differentiation and self-renewal in multipotent, tissue-specific, mature, or progenitor stem cells. In self-renewal, stem cells can continue to exist throughout an organism's lifetime, whereas in differentiation, progenitors and mature cells are passed on for regeneration purposes or tissue genesis. Differentiation is typically a one-way process, as after division, the cell loses the self-renewal property, but in the case of self-renewal in stem cells, the progenitor cells end up dividing to form mature cells and do not lose the self-renewal characteristic (Figs. 1 and 2) [1, 2].

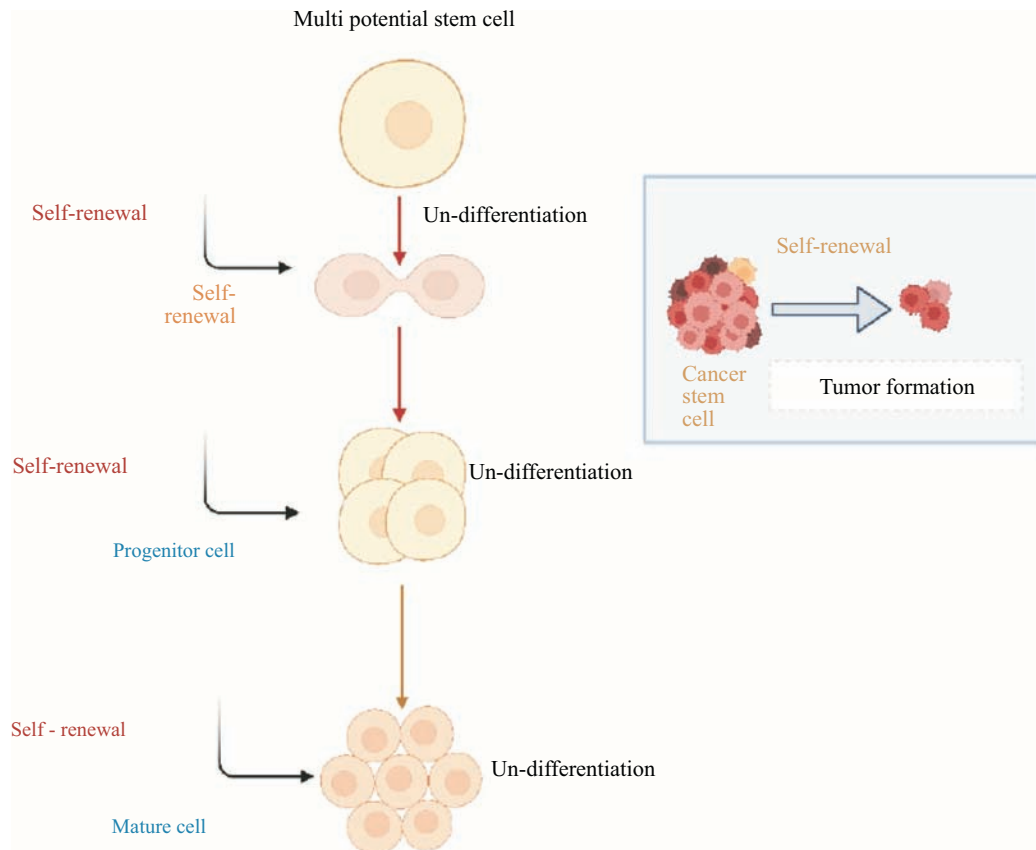


Fig. (1). Differentiation of multi-potent stem cells.

SUBJECT INDEX

A

Abiogenesis 1, 2
 Acidic keratins 183, 191
 Actin 184, 185, 186, 187, 188, 189, 190
 bundles 185, 186
 filaments 184, 185, 186, 187, 188, 189, 190
 networks 185
 Active transport 138, 140, 151
 Adult stem cells (ASCs) 290, 293, 298
 Alcoholic liver disease 297
 Aldehyde dehydrogenase 326, 335
 Alpha keratins 190, 191
 Alzheimer's disease 134, 155, 171, 193, 229, 263
 Amyotrophic lateral sclerosis 193, 229
 Angiogenesis 332, 346,
 Antigen-dependent cellular cytotoxicity (ADCC) 270
 Anoikis 256, 257, 258, 259, 260, 261
 Apoptotic protease-activating factor 1 (Apaf-1) 252
 Aquaporin 10, 126, 151
 Aspartylglucosaminuria 115
 Autolysosomes 103, 109, 110, 132
 Autophagy 15, 30, 47, 52, 106, 109, 110, 111, 112, 118, 119, 120, 124, 131

B

Basal cell carcinoma 341
 Biconcave cells 9
 Binary fission 3, 10, 29
 Biofuel production 20, 30
 Bioremediation 22, 26, 31
 Bitopic 141, 142
 Blastocyst 292, 301
 Blood disorders 289, 298, 314, 316
 Bone marrow transplantation 93, 305
 Brain organoids 300

Breast cancer 95, 98, 156, 325, 328, 329, 334, 388, 389

C

Cajal bodies 37, 38
 Calcium-activated potassium channel 149
 Callogenesis 307
 Callus 296, 306, 307
 Cancer 300, 303, 321, 322, 325, 326, 329
 heterogeneity 322, 329
 stem cells 300, 303, 321, 325, 326
 Cardiac cells 149
 Cardiomyocytes 57, 199, 222, 305
 CAR-T cell therapy 138
 Caseous necrosis 274, 275
 Caspases 247, 252, 278, 280
 Cell
 adhesion 24, 134, 256, 258, 325, 346
 communication 209, 210, 212, 251
 cycle 10, 125, 209, 305, 315, 339, 346
 division 10, 29, 35, 189, 207, 290, 308, 315, 346
 fusion 311
 junctions 11
 motility 10, 182, 216
 shrinkage 247, 249, 253, 254, 280
 signaling 8, 24, 165, 194, 206, 207, 09, 212, 213
 surface receptors 208, 215, 216, 218
 therapy 138, 289, 290, 296, 297, 304, 305
 Cerebral myopathy 171
 Cervical cancer 345, 346
 Chemotherapy 87, 156, 230, 303, 314, 324, 329, 333, 335, 340, 341, 346, 347
 Cirrhosis 297
 Clathrin 64, 65, 110, 119, 157
 Coagulative necrosis 272, 273, 274
 Colorectal cancer 328, 343, 344
 Combination therapy 342, 343

**K.N. Aruljothi, Prakash Gangadaran, Krishnan Anand, Satish Ramalingam, K. Kumaran & Kruthika
 Prakash
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Co-translational translocation 48, 79
CRISPR-Cas9 34, 42, 299, 312

D

Detoxification 123, 124, 125, 126, 135, 265, 297, 308, 309, 312, 313
DNA 37, 39, 40, 42, 96, 142, 195, 249, 253, 254, 267, 268, 270, 280, 338
 fragmentation 249, 253, 254, 267, 268, 270, 280
 methylation 308, 309, 312, 313
 repair 37, 39, 40, 42, 96, 267, 268, 338
Drug delivery systems 122, 138, 157

E

Electron transport chain 166, 167
Embryogenesis 207, 310, 328
Embryonic stem cells 113, 128, 167, 183, 194
Endocytosis 143, 157, 220
Endocrine signaling 208
Endolysosomes 103
Endosymbiosis 163, 173
Enzyme replacement therapy 114, 118
Epigenetics 289, 308, 309, 316, 322
Epithelial-mesenchymal transition 192, 261, 330

F

Ferroptosis 248, 249, 262, 263, 264, 265, 266, 267, 281, 282
Fetal stem cells 292
Fluid mosaic model 138, 140, 142, 158

G

Gangrenous necrosis 275, 276
Gap junctions 210
Gaucher disease 70, 117, 118
Genome editing 161, 170, 172, 178, 312
Glial fibrillary acidic proteins 192, 196
Glycosylation 49, 64, 65, 66, 69, 71, 74, 107, 213
Growth factors 207, 208, 213, 214, 216, 217, 225, 314, 327
Guanine nucleotide-binding proteins 216

H

Hematopoietic stem cells 291, 314
Heterochromatin 37, 310, 313
Heteroplasmy 164, 169
Hedgehog pathway 328
Histone modification 308, 310
Huntington's disease 68, 171, 311
Hydrolytic enzymes 102, 131

I

Immune checkpoint inhibitors 256, 324
Immunotherapy 256, 327, 340, 342, 343, 344, 349
Induced pluripotent stem cells (iPSCs) 292, 294
Insulin 207, 208, 211, 214, 217
Integral proteins 141, 147, 150
Intrinsic pathway 251, 252, 258, 259, 264
Ion 130, 138, 215, 217, 222, 232
 channels 138, 215, 217, 222, 232
 regulation 130
Isotonic solutions 146

J

Juxtacrine signaling 207, 208

K

Keratin 190, 195, 328
Kinesins 197, 198

L

Lamellipodia 189
Laminopathies 39, 42, 43, 194
Leber's hereditary optic neuropathy 171
Ligand-gated ion channels 215, 217, 232
Lipid 139, 140, 142, 143, 145, 147, 153, 154, 155, 215, 232, 248, 263, 264, 265, 266, 281
 bilayer 139, 140, 143, 145, 147, 153, 232
 metabolism 215, 264, 265, 266
 peroxidation 248, 263, 264, 265, 266, 281
 rafts 142, 154, 155
Liquefactive necrosis 273, 274, 275
Liver failure 41, 93, 297

Lysosomal 70, 102, 107, 108, 110, 113, 114, 115, 116
dysfunction 70, 102, 114
membrane proteins 107, 108, 110, 113, 114
storage disorders (LSDs) 115, 116

M

Macrophages 104, 143, 212, 255, 270, 271, 276, 324
Melanoma 337, 341
Membrane blebbing 249, 253, 254, 270
Mesenchymal stem cells 292, 293, 303, 310, 311, 313
Metabolic disorders 54, 55
Metastasis 10, 73, 98, 145, 228, 261, 302, 322, 330, 333, 339, 341, 348, 349
Microfilaments 182, 183, 184, 189, 201
Mitochondrial DNA 161, 168, 170, 171, 178, 269, 301
Molecular chaperones 49, 196
Monotopic 141, 142
Multipotent stem cells 291
Muscular dystrophy 118, 119, 194
Myelodysplastic syndrome 92
Myofibrillar myopathy 192
Myosin 186, 187, 188, 192, 196

N

Nanodelivery 72
Necroptosis 249, 269, 276, 277, 278, 279, 281
Necrosis 246, 247, 252, 262, 267, 272, 273, 277, 279, 281, 282
NETosis 248, 249, 269, 270, 281
Neural cells 296, 305, 309
Neurodegenerative diseases 68, 73, 74, 134, 182, 192, 200, 248, 255, 268, 269, 278, 305, 313
Neurofilament proteins 193
Neurotransmitters 207, 211, 215, 217, 222
Notch pathway 303, 322, 328
Nuclear 34, 37, 39, 142, 194
pore complexes 34, 37
lamins 39, 142, 194

O

Oligonucleotide fingerprinting 82

Oligopotent stem cells 291
Onco-ribosomes 95
Oncogenes 278, 304, 348
Oogenesis 163
Organ regeneration 289, 297
Osmosis 146
Osmoregulation 122, 124, 125, 128
Osteoblasts 309, 314
Oxidative phosphorylation 161, 165, 171, 172, 178, 279

P

Paracrine signaling 207
Parthanatos 267, 268, 269, 281
Pauci-molecular theory 140
Phagolysosomes 103, 110, 111
Phagophore 106, 112, 132, 133
PI3K/Akt pathway 98, 259, 261
Pluripotent stem cells 291, 292, 294, 296, 300, 301, 305, 315
Poly (ADP-ribose) polymerase 267, 338
Polytopic 141, 142
Progenitor cells 92, 95, 291, 305, 314, 315, 323
Protein 8, 24, 35, 48, 49, 50, 51, 64, 65, 66, 77, 78, 79, 80, 81, 83, 85, 96, 97, 183
glycosylation 64, 65
folding 48, 49, 50, 51, 66, 80
synthesis 8, 24, 35, 48, 77, 78, 79, 80, 81, 83, 85, 96, 97, 183
Protofilament 195, 196, 197
Proto-lysosomes 109, 110
Pyroptosis 248, 249, 282

Q

Quiescence 322, 338, 346

R

Reactive oxygen species 166, 169, 170, 178, 270, 277
Regenerative medicine 294, 300, 311, 315, 316
Retinitis pigmentosa 114, 222, 310
Ribosomal proteins 77, 78, 175
Ribosome 90, 91, 97, 98, 99
profiling 97, 98, 99
recycling 90, 91

Subject Index

Ribosomopathies 91, 95, 99

S

Sarcomere 186, 187, 188
Schistosomiasis 164
Signal transduction 196, 206, 207, 208, 209,
213, 214, 215, 216, 217, 218, 231, 232
Single-cell RNA sequencing 256, 261, 281
Skin cancer 339, 341, 342
Sliding filament model 186
Sphingolipidoses 153, 154
Sphingolipids 142, 153, 154,
Spinal muscular atrophy 39, 43
Squamous cell carcinoma 341
Synaptic 154, 217
 plasticity 154
 signaling 217

T

Tay-Sachs disease 134, 154
Therapeutic potential 47, 77, 232, 289, 298,
301, 303, 314, 316
Thylakoids 174, 176
Tissue homeostasis 119, 212, 249, 252, 254,
255, 313
Tonoplast 123, 126, 127, 128, 135
Trans-golgi network (TGN) 64, 71, 105
Tumor 43, 223, 232, 304, 325, 327, 326, 330,
331, 332, 334, 346, 347
 angiogenesis 330, 331, 332
 microenvironment 326, 331, 334, 346, 347
 progression 223, 232, 325, 327, 330, 332,
 suppressor genes 43, 304
Turgor pressure 122, 123, 125, 128, 130, 135

U

Ulcerative colitis 54
Umbilical cord 291, 292, 294, 305, 314
Unipotent stem cells 292
Uniparental inheritance 164

V

Vacuolar pH 134
Vimentin 191, 192, 195, 196, 331
Voltage-gated potassium channel 150

Cell Biology: Basics to Breakthroughs 357

W

Wnt/ β -catenin pathway 328
Wound healing 207, 211, 213, 315

Z

Zygote 30, 291



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