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It was probably the French chemist Portes, who first reported in 1880 that the mucin in the vitreous body, which he named hyalomucine, behaved differently from other mucoids in cornea and cartilage. Fifty four years later Karl Meyer isolated a new polysaccharide from the vitreous, which he named hyaluronic acid. Today its official name is hyaluronan, and modern-day research on this polysaccharide continues to grow. Expertly written by leading scientists in the field, this book provides readers with a broad, yet detailed review of the chemistry of hyaluronan, and the role it plays in human biology and pathology. Twenty-seven chapters present a sequence leading from the chemistry and biochemistry of hyaluronan, followed by its role in various pathological conditions, to modified hylauronans as potential therapeutic agents and finally to the functional, structural and biological properties of hyaluronidases. Chemistry and Biology of Hyaluronan covers the many interesting facets of this fascinating molecule, and all chapters are intended to reach the wider research community. Comprehensive look at the chemistry and biology of hyaluronans Essential to Chemists, Biochemists and Medical researchers Broad yet detailed review of this rapidly growing research area

Physical Biology of the Cell is a textbook for a first course in physical biology or biophysics for undergraduate or graduate students. It maps the huge and complex landscape of cell and molecular biology from the distinct perspective of physical biology. As a key organizing principle, the proximity of topics is based on the physical concepts that

Phenotypic Switching: Implications in Biology and Medicine provides a comprehensive examination of phenotypic switching across biological systems, including underlying mechanisms, evolutionary significance, and its role in biomedical science. Contributions from international leaders discuss conceptual and theoretical aspects of phenotypic plasticity, its influence over biological development, differentiation, biodiversity, and potential applications in cancer therapy, regenerative medicine and stem cell therapy, among other treatments. Chapters discuss fundamental mechanisms of phenotypic switching, including transition states, cell fate decisions, epigenetic factors, stochasticity, protein-based inheritance, specific areas of human development and disease relevance, phenotypic plasticity in melanoma, prostate cancer, breast cancer, non-genetic heterogeneity in cancer, hepatitis C, and more. This book is essential for active researchers, basic and translational scientists, clinicians, postgraduates and students in genetics, human genomics, pathology, bioinformatics, developmental biology, evolutionary biology and adaptive opportunities in yeast. Thoroughly addresses the conceptual, experimental and translational aspects that underlie phenotypic plasticity Emphasizes quantitative approaches, nonlinear dynamics, mechanistic insights and key methodologies to advance phenotypic plasticity studies Features a diverse range of chapter contributions from international leaders in the field

This book, first published in 2005, is a discussion for advanced physics students of

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how to use physics to model biological systems.

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Readers experience for themselves how the coloring of a carefully designed picture almost magically creates understanding. Indispensable for every biology student.

Advances in Protein Molecular and Structural Biology Methods offers a complete overview of the latest tools and methods applicable to the study of proteins at the molecular and structural level. The book begins with sections exploring tools to optimize recombinant protein expression and biophysical techniques such as fluorescence spectroscopy, NMR, mass spectrometry, cryo-electron microscopy, and X-ray crystallography. It then moves towards computational approaches, considering structural bioinformatics, molecular dynamics simulations, and deep machine learning technologies. The book also covers methods applied to intrinsically disordered

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proteins (IDPs) followed by chapters on protein interaction networks, protein function, and protein design and engineering. It provides researchers with an extensive toolkit of methods and techniques to draw from when conducting their own experimental work, taking them from foundational concepts to practical application. Presents a thorough overview of the latest and emerging methods and technologies for protein study Explores biophysical techniques, including nuclear magnetic resonance, X-ray crystallography, and cryo-electron microscopy Includes computational and machine learning methods Features a section dedicated to tools and techniques specific to studying intrinsically disordered proteins

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