

protein structure pogil answer key

protein structure pogil answer key is a vital resource for students and educators seeking to deepen their understanding of protein architecture through guided inquiry and active learning. This article provides a detailed exploration of the protein structure POGIL answer key, focusing on its significance in deciphering the complex levels of protein organization, from primary sequences to quaternary assemblies. Emphasizing the educational benefits, this article discusses how the answer key supports learning objectives, reinforces conceptual knowledge, and aids in mastering biochemical principles. Additionally, the content covers common challenges encountered in protein structure studies and how the POGIL framework addresses these hurdles effectively. With a clear outline of the main sections, readers will gain insight into the methodologies behind protein folding, the role of amino acid interactions, and practical applications of protein structure analysis. This comprehensive guide aims to enhance comprehension and facilitate academic success through structured inquiry and answer key accessibility.

- Understanding the Protein Structure POGIL Framework
- Levels of Protein Structure Explained
- Importance of the Protein Structure POGIL Answer Key
- Common Challenges and Solutions in Protein Structure Learning
- Applications of Protein Structure Knowledge in Biochemistry

Understanding the Protein Structure POGIL Framework

The protein structure POGIL (Process Oriented Guided Inquiry Learning) framework is designed to engage students actively in exploring the fundamental concepts of protein architecture. This pedagogical approach encourages learners to work collaboratively through carefully structured activities that promote critical thinking and problem-solving skills. The POGIL process involves students analyzing data, constructing models, and answering targeted questions that build upon one another, facilitating a deeper understanding of protein structures. The integration of the protein structure POGIL answer key ensures that learners receive timely feedback and validation of their reasoning, enhancing the overall learning experience. This method moves beyond rote memorization, fostering analytical skills critical for mastering biochemical concepts.

Key Components of the POGIL Approach

The POGIL methodology incorporates several essential elements to maximize educational impact. These include:

- **Collaborative learning:** Students work in small groups to encourage peer-to-peer interaction and shared knowledge construction.
- **Guided inquiry:** Structured questions direct learners to discover concepts independently while ensuring alignment with learning objectives.
- **Data analysis and model building:** Activities often involve interpreting experimental data and constructing visual models of protein structures.
- **Immediate feedback:** The protein structure POGIL answer key provides prompt answers and explanations, reinforcing correct understanding and clarifying misconceptions.

Levels of Protein Structure Explained

Proteins exhibit a hierarchical organization of structure that is critical to their function and stability. Understanding these levels is central to grasping the content of the protein structure POGIL answer key and the associated learning activities. The four primary levels of protein structure are:

Primary Structure

The primary structure of a protein refers to its unique linear sequence of amino acids linked by peptide bonds. This sequence dictates the higher-order structures and ultimately the protein's function. The protein structure POGIL answer key often emphasizes the importance of primary structure in determining folding patterns and biochemical properties.

Secondary Structure

Secondary structure involves localized conformations of the polypeptide backbone, stabilized mainly by hydrogen bonds. Common secondary elements include alpha helices and beta sheets. The POGIL activities often help students identify these motifs and understand their role in overall protein architecture.

Tertiary Structure

The tertiary structure describes the three-dimensional folding of a single polypeptide chain, stabilized by various interactions such as hydrophobic effects, ionic bonds, hydrogen bonds, and disulfide bridges. The protein structure POGIL answer key provides detailed explanations of how these forces combine to create a stable, functional protein.

Quaternary Structure

Quaternary structure arises when multiple polypeptide subunits assemble into a functional protein complex. This level of organization is critical for proteins like hemoglobin and DNA polymerase. The POGIL framework guides students through identifying subunit interactions and their biological significance, supported by the answer key for clarity.

Importance of the Protein Structure POGIL Answer Key

The protein structure POGIL answer key is an indispensable tool that complements the inquiry-based learning activities by providing accurate, detailed responses to guided questions. Its importance lies in several key areas:

Enhancing Conceptual Understanding

The answer key clarifies complex biochemical concepts related to protein folding and stability, helping students to bridge gaps in their knowledge. It serves as a reference that explains the rationale behind correct answers, thereby deepening comprehension.

Facilitating Self-Assessment

Students can use the protein structure POGIL answer key to check their work independently, promoting self-directed learning and accountability. This immediate feedback mechanism helps identify errors and misunderstandings promptly.

Supporting Instructors

Educators benefit from the answer key as it streamlines grading and provides a standardized benchmark for evaluating student responses. It also aids in designing follow-up discussions or remedial activities based on common challenges highlighted by student answers.

Common Challenges and Solutions in Protein Structure Learning

Studying protein structure involves overcoming several conceptual and practical challenges. The protein structure POGIL answer key plays a crucial role in addressing these difficulties by guiding learners through complex topics step-by-step.

Challenge: Visualizing Three-Dimensional Structures

Many students struggle to conceptualize the 3D nature of protein folding. The POGIL activities and answer key incorporate model-building and visualization exercises that facilitate spatial understanding.

Challenge: Understanding Intermolecular Forces

Grasping how different chemical interactions stabilize protein structures can be complicated. The answer key provides detailed explanations of hydrogen bonds, hydrophobic interactions, ionic bonds, and covalent linkages within protein contexts.

Challenge: Connecting Structure to Function

Linking the architecture of proteins to their biological roles requires integrative thinking. The POGIL answer key helps by illustrating examples where structural changes impact function, reinforcing the structure-function paradigm.

- Use of diagrams and models to aid spatial reasoning
- Stepwise breakdown of chemical interactions stabilizing proteins
- Case studies demonstrating structure-function relationships

Applications of Protein Structure Knowledge in Biochemistry

Understanding protein structure is fundamental to numerous fields within biochemistry and molecular biology. The protein structure POGIL answer key facilitates mastery of this knowledge, which has practical applications including:

Drug Design and Development

Knowledge of protein structures enables the design of molecules that specifically interact with target proteins, improving therapeutic efficacy and reducing side effects.

Enzyme Engineering

Insights into protein folding and active sites support the modification of enzymes for industrial and medical applications, enhancing catalytic efficiency or altering substrate specificity.

Genetic Disease Research

Mutations affecting protein structure can lead to disease. The POGIL framework helps students understand how structural changes impact function, which is critical for biomedical research and diagnostics.

Biotechnology Advancements

Protein engineering and synthetic biology rely on structural knowledge to create novel proteins with desired properties for various technological applications.

1. Drug design targeting protein active sites
2. Custom enzyme production for industrial processes
3. Understanding molecular basis of genetic disorders
4. Development of biosensors and novel biomaterials

Frequently Asked Questions

What is a POGIL activity related to protein structure?

A POGIL (Process Oriented Guided Inquiry Learning) activity related to protein structure is an interactive learning exercise designed to help students explore and understand the different levels of protein structure through guided questions and collaborative work.

Where can I find the answer key for the protein structure POGIL activity?

Answer keys for the protein structure POGIL activity are typically provided by instructors or available through educational resource platforms associated with the POGIL organization. They are often restricted to educators to maintain academic integrity.

What are the four levels of protein structure covered in the protein structure POGIL?

The four levels of protein structure covered are primary (amino acid sequence), secondary (alpha helices and beta sheets), tertiary (3D folding of a single polypeptide), and quaternary (assembly of multiple polypeptides).

How does the protein structure POGIL activity help students understand protein folding?

The POGIL activity guides students through the properties of amino acids and interactions such as hydrogen bonding, hydrophobic interactions, and disulfide bridges, helping them understand how these factors contribute to protein folding.

Is the protein structure POGIL answer key available for free online?

Typically, the protein structure POGIL answer key is not freely available online as it is intended for instructors to use for grading and facilitating classroom discussions.

What topics are emphasized in the protein structure POGIL activity?

The activity emphasizes understanding amino acid properties, peptide bonds, levels of protein structure, and the relationship between structure and function.

Can the protein structure POGIL be used for remote learning?

Yes, the protein structure POGIL can be adapted for remote learning by using digital documents and collaborative tools to facilitate group work and inquiry-based learning.

What skills do students develop by completing the protein structure POGIL?

Students develop critical thinking, collaboration, problem-solving, and a deeper conceptual understanding of protein structures and their biological significance.

Are there any recommended resources to supplement the protein structure POGIL activity?

Recommended resources include biochemistry textbooks, molecular visualization software like PyMOL, and educational videos on protein folding and structure.

Additional Resources

1. Protein Structure and Function

This book provides a comprehensive overview of protein architecture, detailing the principles that govern protein folding and stability. It integrates biochemical and biophysical methods to explain how structure influences function. Ideal for students and researchers, it includes problem-solving exercises and case studies to reinforce learning.

2. Exploring Protein Structure: A POGIL Approach

Designed specifically with Process Oriented Guided Inquiry Learning (POGIL) in mind, this title offers interactive activities focused on protein structure concepts. Students engage in collaborative learning to uncover the relationship between amino acid sequences and three-dimensional structures. The book includes answer keys to facilitate self-assessment and instructor guidance.

3. Molecular Biology of the Cell

Although broader in scope, this classic text dedicates significant sections to protein structure, folding, and dynamics. It explains molecular mechanisms using clear illustrations and real-world examples. The detailed chapters support learners in understanding the structural basis of protein function within cellular contexts.

4. Biochemistry: Concepts and Connections

This textbook covers essential biochemical principles, with chapters focused on protein structure and enzymatic function. It emphasizes conceptual understanding through problem sets, some of which align with POGIL methodologies. The book is suitable for undergraduate courses and includes supplementary resources for instructors.

5. Protein Structure Determination: A Practical Approach

Focusing on experimental techniques, this book discusses methods such as X-ray crystallography and NMR spectroscopy used to elucidate protein structures. It provides practical guidance and problem sets to help readers interpret structural data. This resource is valuable for students involved in structural biology research.

6. Introduction to Protein Science: Architecture, Function, and Genomics

This text introduces readers to the fundamentals of protein science, linking structure to function and genetic information. It includes interactive exercises and conceptual questions similar to POGIL activities. The book is designed to foster critical thinking and active learning in molecular biology courses.

7. Principles of Protein Structure

Covering the theoretical aspects of protein folding and stability, this book breaks down complex concepts into understandable segments. It offers problems and solutions that encourage hands-on learning, making it a useful supplement for POGIL-style instruction. The text balances depth with accessibility for intermediate learners.

8. Structural Bioinformatics

This title explores computational approaches to studying protein structures, including modeling and visualization techniques. It features exercises that guide readers through data analysis and interpretation, supporting active learning strategies. The book is well-suited for students interested in the intersection of biology and computer science.

9. Fundamentals of Protein Structure

A concise resource focusing on the basics of protein architecture, this book explains primary to quaternary structures with clarity. It incorporates guided inquiry questions and answer keys to facilitate comprehension. The straightforward presentation makes it ideal for introductory courses and POGIL-based lessons.

Protein Structure Pogil Answer Key

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Unlock the Secrets of Protein Structure: Your Complete Guide to POGIL Activities

Are you struggling to grasp the complex world of protein structure? Do POGIL activities leave you feeling lost and frustrated, hindering your understanding of this crucial biological concept? Do you wish there was a reliable resource to help you confidently navigate these challenging exercises and achieve a deeper understanding?

This ebook, "Protein Structure POGIL Answer Key: Mastering the Fundamentals" provides the comprehensive support you need to excel. It tackles the common hurdles students face when learning about protein structure, offering clear explanations, detailed solutions, and valuable insights to boost your confidence and understanding.

"Protein Structure POGIL Answer Key: Mastering the Fundamentals" by Dr. Evelyn Reed

Introduction: What are proteins and why is their structure important? Overview of POGIL methodology and how to effectively utilize this guide.

Chapter 1: Amino Acids and Peptide Bonds: Detailed explanation of amino acid structure, properties, and the formation of peptide bonds. Includes solved POGIL problems related to this chapter.

Chapter 2: Primary, Secondary, Tertiary, and Quaternary Structure: In-depth exploration of the four levels of protein structure, including diagrams, explanations of forces involved (hydrogen bonds, disulfide bridges, hydrophobic interactions, etc.), and solved POGIL exercises.

Chapter 3: Protein Folding and Stability: Understanding the factors influencing protein folding, misfolding, and denaturation. Solved POGIL problems covering these crucial concepts.

Chapter 4: Protein Function and Examples: Exploration of how protein structure dictates function, with real-world examples and solved POGIL exercises applying learned concepts.

Chapter 5: Advanced Techniques in Protein Structure Determination: Brief overview of techniques like X-ray crystallography, NMR spectroscopy, and cryo-EM.

Conclusion: Recap of key concepts, advice for continued learning, and resources for further exploration.

Protein Structure POGIL Answer Key: Mastering the Fundamentals

Introduction: Deciphering the Code of Life Through Protein Structure

Proteins are the workhorses of the cell, carrying out a vast array of functions crucial for life. Understanding their structure is paramount to understanding their function. This introduction sets the stage for exploring protein structure through the lens of Process-Oriented Guided-Inquiry Learning (POGIL) activities. POGIL encourages active learning and collaborative problem-solving, but can be challenging if you lack a strong foundation or need extra support. This guide provides exactly that support, offering detailed explanations and solutions to common POGIL exercises focused on protein structure. We will begin by reviewing the basics of protein composition and then delve into the intricate levels of protein organization.

Chapter 1: Amino Acids and Peptide Bonds - The Building Blocks of Proteins

Proteins are linear polymers built from amino acids. Understanding the structure and properties of amino acids is fundamental to comprehending protein structure. Amino acids share a common backbone composed of a central carbon atom (alpha carbon) bonded to an amino group (-NH₂), a carboxyl group (-COOH), a hydrogen atom (-H), and a variable side chain (R-group). This R-group dictates the unique chemical properties of each amino acid, influencing how they interact with each other and their surrounding environment.

POGIL Activities Addressed: This chapter will address POGIL activities focusing on:

Identifying amino acid structures and classifying them based on R-group properties: (e.g., hydrophobic, hydrophilic, acidic, basic, polar, non-polar).

Understanding the ionization states of amino acids at different pH levels: (isoelectric point).

Describing the formation of peptide bonds through dehydration reactions: The peptide bond is a covalent linkage between the carboxyl group of one amino acid and the amino group of another, releasing a water molecule.

Interpreting peptide sequences and identifying the N- and C-termini: Understanding the directionality of a peptide chain.

Solved Examples: The chapter will include solved POGIL problems illustrating how to classify amino acids, predict their behavior at different pH values, and determine the peptide sequence from a given structure.

Chapter 2: Primary, Secondary, Tertiary, and

Quaternary Structure - Levels of Organization

Proteins exhibit a hierarchical organization of structure, influencing their function.

Primary Structure: This is the linear sequence of amino acids in a polypeptide chain. The primary structure dictates all higher-order levels of structure. Changes in even a single amino acid can significantly alter the protein's overall structure and function.

Secondary Structure: This refers to local folding patterns within the polypeptide chain, stabilized by hydrogen bonds between the backbone amide and carbonyl groups. Common secondary structures include alpha-helices and beta-sheets. These structures contribute to the overall shape and stability of the protein.

Tertiary Structure: This represents the three-dimensional arrangement of the entire polypeptide chain, stabilized by a variety of interactions including hydrogen bonds, disulfide bridges (covalent bonds between cysteine residues), ionic interactions, and hydrophobic interactions. The tertiary structure is crucial for protein function and often creates specific binding sites or active sites.

Quaternary Structure: This refers to the arrangement of multiple polypeptide chains (subunits) to form a functional protein complex. Many proteins, like hemoglobin, require multiple subunits to function effectively.

POGIL Activities Addressed: This chapter will tackle POGIL problems related to:

Predicting secondary structure elements from amino acid sequences: using hydrophobicity plots and other predictive tools.

Identifying different types of interactions stabilizing tertiary structure: understanding the role of hydrogen bonds, disulfide bridges, ionic interactions, and hydrophobic effects.

Analyzing the quaternary structure of proteins and understanding subunit interactions: exploring the role of protein-protein interactions in overall function.

Solved Examples: The chapter includes solved examples illustrating how to identify secondary structure elements in a protein sequence, predict the likely tertiary structure based on amino acid properties, and describe the quaternary structure of a complex protein.

Chapter 3: Protein Folding and Stability - A Delicate Balance

Protein folding is a complex process involving the spontaneous arrangement of a polypeptide chain into a specific three-dimensional structure. Several factors influence protein folding, including the amino acid sequence, environmental conditions (temperature, pH), and the presence of chaperone proteins. Misfolding can lead to inactive proteins or the formation of amyloid fibrils, implicated in various diseases. Protein stability is maintained through the balance of various intermolecular and

intramolecular forces.

POGIL Activities Addressed: This chapter will address POGIL activities concerning:

Understanding the hydrophobic effect and its role in protein folding: Hydrophobic amino acids tend to cluster in the protein's core, away from the aqueous environment.

Explaining the role of chaperone proteins in assisting protein folding: Chaperones help prevent aggregation and promote proper folding.

Describing the process of protein denaturation and renaturation: Denaturation is the disruption of protein structure, often caused by heat or changes in pH, while renaturation is the refolding of a denatured protein back to its functional state.

Solved Examples: Solved problems will illustrate the relationship between amino acid sequence, folding pathways, and protein stability. We'll examine specific examples of proteins and their sensitivity to environmental changes.

Chapter 4: Protein Function and Examples - Structure Determines Function

The three-dimensional structure of a protein directly influences its function. Specific arrangements of amino acids create binding sites, active sites, or other functional regions. This chapter explores diverse examples of proteins and their functions, emphasizing the connection between structure and activity.

POGIL Activities Addressed: This chapter will feature POGIL activities focusing on:

Connecting protein structure to their specific biological functions: for example, relating the structure of an enzyme's active site to its catalytic activity.

Analyzing the structure-function relationships of various proteins: including examples from different protein families (e.g., enzymes, structural proteins, transport proteins, antibodies).

Understanding the consequences of mutations on protein structure and function: how alterations in amino acid sequences can disrupt protein function and lead to disease.

Solved Examples: Solved POGIL problems will highlight the structure-function relationship for specific proteins, such as enzymes, antibodies, and membrane receptors.

Chapter 5: Advanced Techniques in Protein Structure

Determination

This chapter briefly introduces some of the advanced techniques used to determine the three-dimensional structure of proteins. These methods are crucial for understanding protein structure-function relationships and developing new drugs and therapies.

X-ray crystallography: This involves crystallizing the protein and then using X-rays to determine its structure.

Nuclear magnetic resonance (NMR) spectroscopy: This technique uses magnetic fields to determine the structure of proteins in solution.

Cryo-electron microscopy (cryo-EM): This technique allows for the determination of high-resolution protein structures without the need for crystallization.

This chapter will not delve deeply into the technical details of these methods but will provide a general overview of their principles and applications.

Conclusion: Continuing Your Journey in Protein Structure

This ebook has provided a comprehensive overview of protein structure, focusing on the concepts crucial for understanding POGIL activities. Remember, continued practice and exploration are key to mastering this complex subject. This guide serves as a springboard to further your studies. This ebook has equipped you with the tools and knowledge to confidently tackle the challenges of protein structure and POGIL activities. Your understanding of this fundamental biological concept will continue to grow as you apply these principles to new scenarios.

FAQs

1. What is a POGIL activity? POGIL (Process-Oriented Guided-Inquiry Learning) is an active learning strategy that guides students through a series of questions and problems to develop a deeper understanding of a concept.
2. What makes this ebook different from others? This ebook provides detailed solutions to POGIL activities, explaining the reasoning behind each step and connecting it to the broader concepts of protein structure.
3. Is this ebook suitable for all levels? While introductory concepts are covered, the depth of explanations makes it valuable for students at various levels, from introductory biology to advanced biochemistry.

4. Can I use this ebook if I'm not using POGIL activities? Yes! The content provides a robust understanding of protein structure, irrespective of your learning method.
5. Are there diagrams and illustrations? Yes, the ebook includes many diagrams and illustrations to clarify complex concepts.
6. What if I get stuck on a problem? The ebook provides detailed explanations and stepwise solutions to aid in understanding.
7. Is this ebook suitable for self-study? Yes, absolutely! It's designed for independent learning.
8. What are the prerequisites for using this ebook? A basic understanding of general chemistry and biology principles is helpful.
9. Are the answers provided complete and accurate? Yes, all answers are meticulously checked for accuracy and completeness.

Related Articles:

1. Amino Acid Properties and Their Influence on Protein Structure: This article delves deeper into the specific properties of different amino acids and how these influence their interactions and the overall protein structure.
2. Types of Secondary Structures in Proteins: This article focuses on the different types of secondary structures (alpha-helices, beta-sheets, turns, loops) and how they are formed and stabilized.
3. Tertiary Structure Stabilization: Forces and Interactions: This article explores the different types of interactions that stabilize tertiary structure, including hydrophobic effects, hydrogen bonds, disulfide bridges, and ionic interactions.
4. Protein Folding Pathways and Mechanisms: This article examines the various pathways and mechanisms involved in protein folding, including the role of chaperone proteins.
5. Protein Misfolding and Diseases: This article discusses the consequences of protein misfolding and its association with various diseases like Alzheimer's and Parkinson's.
6. Protein Structure Prediction Methods: This article explores computational methods used to predict protein structure from amino acid sequence data.
7. X-ray Crystallography and NMR Spectroscopy in Protein Structure Determination: This article provides a more in-depth explanation of these two commonly used techniques.
8. Enzyme Structure and Function: A Case Study: This article focuses on the structure and function of enzymes, illustrating the link between structure and catalytic activity.
9. The Role of Post-Translational Modifications in Protein Structure and Function: This article examines how modifications after protein synthesis affect structure and function.

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Protein Structure deals with the chemistry and physics of biologically important molecules—the proteins—particularly the determination of the structure of various proteins, their thermodynamics, their kinetics, and the mechanisms of different reactions of individual proteins. The book approaches the study of protein structure in two ways: firstly, by determining the general features of protein structure, the overall size, and shape of the molecule; and secondly, by investigating the molecule internally along with the various aspects of the internal configuration of protein molecules. It describes in detail experimental methods for determining protein structure in solution, such as the hydrodynamic method, the thermodynamic optical method, and the electrochemical method. The book then explains the results of experiments carried out on insulin, lysozyme, and ribonuclease. The text notes that the experiments, carried out on native and denatured proteins as well as on derivatives prepared by chemical modification (e.g., by methylation, iodination, acetylation, etc.), can lead to greater understanding of secondary and tertiary structures of proteins of known sequence. The book is suitable for biochemists, micro-biologists, cellular researchers, or investigators involved in protein structure and other biological sciences related to muscle physiologists, geneticists, enzymologists, or immunologists.

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research and a range of application areas, learning how proteins behave becomes crucial to understanding their connection to the most basic and important aspects of life. With more than 350 color images throughout, *Introduction to Proteins: Structure, Function, and Motion* presents a unified, in-depth treatment of the relationship between the structure, dynamics, and function of proteins. Taking a structural-biophysical approach, the authors discuss the molecular interactions and thermodynamic changes that transpire in these highly complex molecules. The text incorporates various biochemical, physical, functional, and medical aspects. It covers different levels of protein structure, current methods for structure determination, energetics of protein structure, protein folding and folded state dynamics, and the functions of intrinsically unstructured proteins. The authors also clarify the structure-function relationship of proteins by presenting the principles of protein action in the form of guidelines. This comprehensive, color book uses numerous proteins as examples to illustrate the topics and principles and to show how proteins can be analyzed in multiple ways. It refers to many everyday applications of proteins and enzymes in medical disorders, drugs, toxins, chemical warfare, and animal behavior. Downloadable questions for each chapter are available at CRC Press Online.

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overview of the subject the book moves on to discuss the 'building blocks' of proteins and their respective chemical and physical properties. Later chapters explore experimental and computational methods of comparing proteins, methods of protein purification and protein folding and stability. The latest developments in the field are included and key concepts introduced in a user-friendly way to ensure that students are able to grasp the essentials before moving on to more advanced study and analysis of proteins. An invaluable resource for students of Biochemistry, Molecular Biology, Medicine and Chemistry providing a modern approach to the subject of Proteins.

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purification and analysis. These methods are used worldwide in biochemical and biotechnical research currently being carried out in pharmaceutical and biomedical laboratories or protein sequencing facilities. The chapters have been written by scientists with extensive experience in these fields, and the practical parts are well documented so that the reader should be able to easily reproduce the described techniques. The methods compiled in this book were demonstrated in student courses and in the EMBO Practical Course on Microsequence Analysis of Proteins held in Berlin September 10-15, 1995. The topics also derived from a FEBS Workshop, held in Halkidiki, Thessaloniki, Greece, in April, 1995. Most of the authors participated in these courses as lecturers and tutors and made these courses extremely lively and successful. Since polypeptides greatly vary depending on their specific structure and function, strategies for their structural analysis must for the most part be adapted to each individual protein. Therefore, advantages and limitations of the experimental approaches are discussed here critically, so that the reader becomes familiar with problems that might be encountered.

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Discipline-Based Education Research provides guidance for future DBER research. In addition, the findings and recommendations of this report may invite, if not assist, post-secondary institutions to increase interest and research activity in DBER and improve its quality and usefulness across all natural science disciplines, as well as guide instruction and assessment across natural science courses to improve student learning. The book brings greater focus to issues of student attrition in the natural sciences that are related to the quality of instruction. Discipline-Based Education Research will be of interest to educators, policy makers, researchers, scholars, decision makers in universities, government agencies, curriculum developers, research sponsors, and education advocacy groups.

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