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Experimental phasing opportunities for macromolecular crystallography at very long wavelengths

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Despite recent advances in cryo-electron microscopy and artificial intelligence-based model predictions, a significant fraction of structure determinations by macromolecular crystal-lography still requires experimental phasing, usually by means of single-wavelength anomalous diffraction (SAD) techniques. Most synchrotron beamlines provide highly brilliant beams of X-rays of between 0.7 and 2 Å wavelength. Use of longer wavelengths to access the absorption edges of biologically important lighter atoms such as calcium, potassium, chlorine, sulfur and phosphorus for native-SAD phasing is attractive but technically highly challenging. The long-wavelength beamline I23 at Diamond Light Source overcomes these limitations and extends the accessible wavelength range to $\lambda = 5$. Here we report 22 macromolecular structures solved in this extended wavelength range, using anomalous scattering from a range of elements which demonstrate the routine feasibility of lighter atom phasing. We suggest that, in light of its advantages, long-wavelength crystallography is a compelling option for experimental phasing.

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Macromolecular Crystallography

Alexander McPherson

Macromolecular Crystallography:

Introduction to Macromolecular Crystallography Alexander McPherson, 2011-09-20 A comprehensive and approachable introduction to crystallography now updated in a valuable new edition The Second Edition of this well received book continues to offer the most concise authoritative and easy to follow introduction to the field of crystallography Dedicated to providing a complete basic presentation of the subject that does not assume a background in physics or math the book's content flows logically from basic principles to methods such as those for solving phase problems interpretation of Patterson maps and the difference Fourier method the fundamental theory of diffraction and the properties of crystals and applications in determining macromolecular structure This new edition includes a vast amount of carefully updated materials as well as two completely new chapters on recording and compiling X ray data and growing crystals of proteins and other macromolecules Richly illustrated throughout to clarify difficult concepts this book takes a non technical approach to crystallography that is ideal for professionals and graduate students in structural biology biophysics biochemistry and molecular biology who are studying the subject for the first time Macromolecular Crystallography Mark R. Sanderson, Jane V. Skelly, 2007-08-23 Macromolecular crystallography is the study of macromolecules using X ray crystallographic techniques to determine their molecular structure This title synthesises contributions from a team of internationally recognized leaders offering chapters on conventional and high throughput methods Crystallography, Part D Charles W. Carter Jr., Charles W. Carter, Robert M. Sweet, 2003-12-10 Accurate molecular structures is vital for rational drug design and for structure based functional studies directed toward the development of effective therapeutic agents and drugs Crystallography can reliably predict structure both in terms of folding and atomic details of bonding Phases Map interpretation and refinement Analysis and software Macromolecular Crystallography Protocols, Volume 2 Sylvie Doublie, 2008-02-05 In the decade since publication of the first edition of Crystallographic Methods and Protocols the field has seen several major developments that have both accelerated the pace of structure determination and made crystallography accessible to a broader range of investigators Volume I Preparation and Crystallization of Macromolecules is dedicated to the crystallization and ways to increase the odds of obtaining crystals in macromolecules while Volume 2 Structure Determination covers both computational methods for characterizing crystals and solving structures Macromolecular Crystallography Protocols, Volume 1 Sylvie Doublie, 2008-02-04 Macromolecular Crystallography Protocols now in two volumes examines major developments that have occurred since publication of the acclaimed first edition nearly a decade ago Volume 1 Preparation and Crystallization of Macromolecules and Volume 2 Structure Determination explore recent advances that have accelerated the pace of structural determination and made crystallography accessible to a broader range of investigators Volume 1 is composed of detailed protocols for the preparation and optimization of crystals including tips from the experts on the best methods for inducing proteins to adopt their

crystalline form Volume 2 complements the first volume by addressing laboratory techniques for crystal handling and structural characterization as well as computational techniques for data collection phasing and refinement The volume concludes with a detailed and insightful survey of available crystallographic software. These volumes will be an indispensable reference for obtaining macromolecular crystals and determining their three dimensional structure Crystallography with Synchrotron Radiation John R. Helliwell, 1992 This highly illustrated monograph provides a comprehensive treatment of the study of the structure and function of the molecules of life proteins nucleic acids and viruses using synchrotron radiation and crystallography Beginning with chapters on the fundamentals of macromolecular crystallography and macromolecular structure the book goes on to review the sources and properties of synchrotron radiation instrumentation and monochromatic data collection There are also chapters on the Laue method on diffuse X ray scattering and on variable wavelength anomalous dispersion methods. The book concludes with a description and survey of applications including studies at high resolution the use of small crystals the study of large unit cells and time resolved crystallography particularly of enzymes Appendices are provided that present essential information for the synchrotron user as well as information about synchrotron facilities currently available **Macromolecular Crystallography** Charles W. Carter, Robert M. Sweet, 2003 Evolving Methods for Macromolecular Crystallography Randy J. Read, Joel Sussman, 2007-06-25 This volume draws on the expertise of leaders in the field of macromolecular crystallography to illuminate the dramatic developments that are accelerating progress in structural biology. Their contributions span the range of techniques from crystallization through data collection structure solution and analysis The book shows how modern high throughput methods are contributing to a deeper understanding of medical problems Macromolecular Crystallography Maria Armenia Carrondo, Paola Spadon, 2011-12-01 This volume is a collection of the contributions presented at the 42nd Erice Crystallographic Course whose main objective was to train the younger generation on advanced methods and techniques for examining structural and dynamic aspects of biological macromolecules. The papers review the techniques used to study protein assemblies and their dynamics including X ray diffraction and scattering electron cryo electron microscopy electro nanospray mass spectrometry NMR protein docking and molecular dynamics A key theme throughout the book is the dependence of modern structural science on multiple experimental and computational techniques and it is the development of these techniques and their integration that will take us forward in the future Introduction to MacRomolecular Crystallography McPherson, 2002-12-01 Methods in Macromolecular Crystallography North Atlantic Treaty Organization. Scientific Affairs Division, 2001 This volume contains 20 papers from two courses Methods for Macromolecular Crystallography and Chemical Prospective in Crystallography of Molecular Biology held during the late spring of 2000 in Reice Italy The papers discuss crystals synchrotrons detector development data processing ab initio phasing and high and low resolution molecular placement experimental phase measurement density modification methods

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Crystallography, Part C Charles W. Carter Jr.,2003-12-15 Accurate molecular structures are vital for rational drug design and for structure based functional studies directed toward the development of effective therapeutic agents and drugs Crystallography can reliably predict structure both in terms of folding and atomic details of bonding Methodological methods in crystals Methodological methods data analysis Structural Biology in Drug Discovery Jean-Paul Renaud,2020-01-27 With the most comprehensive and up to date overview of structure based drug discovery covering both experimental and computational approaches Structural Biology in Drug Discovery Methods Techniques and Practices describes principles

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Table of Contents Macromolecular Crystallography

- 1. Understanding the eBook Macromolecular Crystallography
 - The Rise of Digital Reading Macromolecular Crystallography
 - Advantages of eBooks Over Traditional Books
- 2. Identifying Macromolecular Crystallography
 - Exploring Different Genres
 - Considering Fiction vs. Non-Fiction
 - Determining Your Reading Goals
- 3. Choosing the Right eBook Platform
 - Popular eBook Platforms
 - Features to Look for in an Macromolecular Crystallography
 - User-Friendly Interface
- 4. Exploring eBook Recommendations from Macromolecular Crystallography
 - Personalized Recommendations
 - Macromolecular Crystallography User Reviews and Ratings
 - Macromolecular Crystallography and Bestseller Lists
- 5. Accessing Macromolecular Crystallography Free and Paid eBooks
 - Macromolecular Crystallography Public Domain eBooks
 - Macromolecular Crystallography eBook Subscription Services
 - Macromolecular Crystallography Budget-Friendly Options
- 6. Navigating Macromolecular Crystallography eBook Formats

- o ePub, PDF, MOBI, and More
- Macromolecular Crystallography Compatibility with Devices
- Macromolecular Crystallography Enhanced eBook Features
- 7. Enhancing Your Reading Experience
 - Adjustable Fonts and Text Sizes of Macromolecular Crystallography
 - Highlighting and Note-Taking Macromolecular Crystallography
 - Interactive Elements Macromolecular Crystallography
- 8. Staying Engaged with Macromolecular Crystallography
 - Joining Online Reading Communities
 - Participating in Virtual Book Clubs
 - Following Authors and Publishers Macromolecular Crystallography
- 9. Balancing eBooks and Physical Books Macromolecular Crystallography
 - Benefits of a Digital Library
 - Creating a Diverse Reading Collection Macromolecular Crystallography
- 10. Overcoming Reading Challenges
 - Dealing with Digital Eye Strain
 - Minimizing Distractions
 - Managing Screen Time
- 11. Cultivating a Reading Routine Macromolecular Crystallography
 - Setting Reading Goals Macromolecular Crystallography
 - Carving Out Dedicated Reading Time
- 12. Sourcing Reliable Information of Macromolecular Crystallography
 - Fact-Checking eBook Content of Macromolecular Crystallography
 - Distinguishing Credible Sources
- 13. Promoting Lifelong Learning
 - Utilizing eBooks for Skill Development
 - Exploring Educational eBooks
- 14. Embracing eBook Trends
 - Integration of Multimedia Elements
 - Interactive and Gamified eBooks

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