

Methods in ENZYMOLOGY

Volume 483

Cryo-EM, Part C

Analyses, Interpretation, and Case Studies

Edited by

Grant J. Jensen



Cryo Em Part C Volume 483 Analyses Interpretation And Case Studies Methods In Enzymology

Carl Wu,C. David Allis



Cryo Em Part C Volume 483 Analyses Interpretation And Case Studies Methods In Enzymology:

Cryo-EM, Part C ,2010-09-30 This volume along with Part A and Part B is dedicated to a description of the instruments samples protocols and analyses that belong to cryo EM It emphasizes the relatedness of the ideas instrumentation and methods underlying all cryo EM approaches which allow practitioners to easily move between them Within each section the articles are ordered according to the most common symmetry of the sample to which their methods are applied Includes time tested core methods and new innovations applicable to any researcher Methods included are useful to both established researchers and newcomers to the field Relevant background and reference information given for procedures can be used as a guide

Research on Nitrification and Related Processes, Part B ,2011-06-15 The global nitrogen cycle is the one most impacted by mankind The past decade has changed our view on many aspects of the microbial biogeochemical cycles including the global nitrogen cycle which is mainly due to tremendous advances in methods techniques and approaches Many novel processes and the molecular inventory and organisms that facilitate them have been discovered only within the last 5 to 10 years and the process is in progress Research on Nitrification and Related Processes Part B provides state of the art updates on methods and protocols dealing with the detection isolation and characterization of macromolecules and their hosting organisms that facilitate nitrification and related processes in the nitrogen cycle as well as the challenges of doing so in very diverse environments Provides state of the art update on methods and protocols Deals with the detection isolation and characterization of macromolecules and their hosting organisms Deals with the challenges of very diverse environments

Nucleosomes, Histones and Chromatin Part A Carl Wu,C. David Allis,2012-08-31 This new volume of Methods in Enzymology continues the legacy of this premier serial by containing quality chapters authored by leaders in the field The first of 2 volumes covering nucleosomes histones and chromatin it has chapters on methods applied to the study of protein arginine methylation high resolution identification of intra and interchromosomal DNA interactions by 4C technology and peptide arrays to interrogate the binding specificity of chromatin binding proteins Continues the legacy of this premier serial by containing quality chapters authored by leaders in the field The first of 2 volumes covering nucleosomes histones and chromatin Chapters on methods applied to the study of protein arginine methylation high resolution identification of intra and interchromosomal DNA interactions by 4C technology and peptide arrays to interrogate the binding specificity of chromatin binding proteins

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other biophysical techniques and computational structural biology Includes detailed chapters on druggability allostery complementary use of thermodynamic and kinetic information and powerful approaches such as structural chemogenomics and fragment based drug design Emphasizes the need for the in depth biophysical characterization of protein targets as well as of therapeutic proteins and for a thorough quality assessment of experimental structures Illustrates advances in the field of established therapeutic targets like kinases serine proteinases GPCRs and epigenetic proteins and of more challenging ones like protein protein interactions and intrinsically disordered proteins

Nucleosomes, Histones & Chromatin Carl Wu, C. David Allis, 2012 Covers nucleosomes histones and chromatin with chapters on dynamic mapping of histone DNA interactions in nucleosomes by unzipping single molecules of DNA Digital DNase technology and Genome wide Analysis of Chromatin Transition

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Ghrelin Masayasu N. Kojima, Kenji Kangawa, 2012-09-25 This new volume of Methods in Enzymology continues the legacy of this premier serial by containing quality chapters authored by leaders in the field The volume covers ghrelin and has chapters on such topics as orphan gpcrs and methods for identifying their ligands ghrelin o acyltransferase assays and inhibition and thermogenic characterization of ghrelin receptor null mice Contains quality chapters authored by leaders in the field Has chapters on such topics as orphan gpcrs and methods for identifying their ligands ghrelin o acyltransferase assays and inhibition and thermogenic characterization of ghrelin receptor null mice

G Protein Coupled Receptors, 2013-01-24 This new volume of Methods in Enzymology continues the legacy of this premier serial by containing quality chapters authored by leaders in the field This volume covers G protein coupled receptors and includes chapters on such topics as G protein coupled receptor trafficking motifs structure based virtual screening and automation friendly high throughput assays for identification of pharmacoperone drugs Continues the legacy of this premier serial with quality chapters authored by leaders in the field Covers G protein coupled receptors Contains chapters on such topics as G protein coupled receptor trafficking motifs

structure based virtual screening and automation friendly high throughput assays for identifying pharmacoporene drugs

Biothermodynamics, Part D, 2011-02-16 The use of thermodynamics in biological research can be equated to an energy book keeping system While the structure and function of a molecule is important it is equally important to know what drives the energy force This volume presents sophisticated methods for estimating the thermodynamic parameters of specific protein protein protein DNA and small molecule interactions Elucidates the relationships between structure and energetics and their applications to molecular design aiding researchers in the design of medically important molecules Provides a must have methods volume that keeps MIE buyers and online subscribers up to date with the latest research Offers step by step lab instructions including necessary equipment from a global research community *Fragment Based Drug Design*

Lawrence C. Kuo, 2011-03-09 There are numerous excellent reviews on fragment based drug discovery FBDD but there are to date no hand holding guides or protocols with which one can embark on this orthogonal approach to complement traditional high throughput screening methodologies This Methods in Enzymology volume offers the tools practical approaches and hit to lead examples on how to conduct FBDD screens The chapters in this volume cover methods that have proven to be successful in generating leads from fragments including chapters on how to apply computational techniques nuclear magnetic resonance surface plasma resonance thermal shift and binding assays protein crystallography and medicinal chemistry in FBDD Also elaborated by experienced researchers in FBDD are sample preparations of fragments proteins and GPCR as well as examples of how to generate leads from hits Offers the tools practical approaches and hit to lead examples on how to conduct FBDD screens The chapters in this volume cover methods that have proven to be successful in generating leads from fragments including chapters on how to apply computational techniques nuclear magnetic resonance surface plasma resonance thermal shift and binding assays protein crystallography and medicinal chemistry in FBDD

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