

Bind Tuning

SJ Ball

Evolution, Emerging Functions and Structure of Actin-Binding Proteins Lei-Miao Yin, Michael Schnoor, Chang-Duk Jun, 2022-02-02

Performance Optimization and Tuning Techniques for IBM Power Systems Processors Including IBM POWER8 Brian Hall, Peter Bergner, Alon Shalev Housfater, Madhusudanan Kandasamy, Tulio Magno, Alex Mericas, Steve Munroe, Mauricio Oliveira, Bill Schmidt, Will Schmidt, Bernard King Smith, Julian Wang, Suresh Warriar, David Wendt, IBM Redbooks, 2017-03-31 This IBM® Redbooks® publication focuses on gathering the correct technical information, and laying out simple guidance for optimizing code performance on IBM POWER8® processor-based systems that run the IBM AIX®, IBM i, or Linux operating systems. There is straightforward performance optimization that can be performed with a minimum of effort and without extensive previous experience or in-depth knowledge. The POWER8 processor contains many new and important performance features, such as support for eight hardware threads in each core and support for transactional memory. The POWER8 processor is a strict superset of the IBM POWER7+™ processor, and so all of the performance features of the POWER7+ processor, such as multiple page sizes, also appear in the POWER8 processor. Much of the technical information and guidance for optimizing performance on POWER8 processors that is presented in this guide also applies to POWER7+ and earlier processors, except where the guide explicitly indicates that a feature is new in the POWER8 processor. This guide strives to focus on optimizations that tend to be positive across a broad set of IBM POWER® processor chips and systems. Specific guidance is given for the POWER8 processor; however, the general guidance is applicable to the IBM POWER7+, IBM POWER7®, IBM POWER6®, IBM POWER5, and even to earlier processors. This guide is directed at personnel who are responsible for performing migration and implementation activities on POWER8 processor-based systems. This includes system administrators, system architects, network administrators, information architects, and database administrators (DBAs).

Expert Oracle RAC Performance Diagnostics and Tuning Murali Vallath, 2014-10-13 Expert Oracle RAC Performance Diagnostics and Tuning provides comprehensive coverage of the features, technology and principles for testing and tuning RAC databases. The book takes a deep look at optimizing RAC databases by following a methodical approach based on scientific analysis rather than using a speculative approach, twisting and turning knobs and gambling on the system. The book starts with the basic concepts of tuning methodology, capacity planning, and architecture. Author Murali Vallath then dissects the various tiers of the testing implementation, including the operating system, the network, the application, the storage, the instance, the database, and the grid infrastructure. He also introduces tools for performance optimization and thoroughly covers each aspect of the tuning process, using many real-world examples, analyses, and solutions from the field that provide you with a solid, practical, and replicable approach to tuning a RAC environment. The book concludes with troubleshooting guidance and quick reference of all the scripts used in the book. Expert Oracle RAC Performance Diagnostics and Tuning covers scenarios and details never discussed before in any other performance tuning books. If you have a RAC database, this book is a requirement. Get your copy today. Takes you through optimizing the various tiers of the RAC environment. Provides real life case studies, analysis and solutions from the field. Maps a methodical approach to testing, tuning and diagnosing the cluster

Current Perspectives on Insulin-like Growth Factor Binding Protein (IGFBP) Research Andreas Hoefflich, John Pintar, Briony Forbes, 2019-02-06 The family of IGFBPs has been developed by the duplication of genes and genomes and contributes to genetic and functional diversity. Due to the different protein domains present in the molecule, IGFBPs can be seen as mediators of tissue-specific IGF-functions. However, IGFBPs also have IGF-independent functions both inside and outside the cell. These diverse genetic, molecular and functional aspects of IGFBPs are discussed within this Research Topic. Accumulating data provide evidence for the regulation of IGFBP-functions by proteases, which may acutely regulate bioactivity of the IGFs. However, during proteolytic degradation IGFBP-fragments with novel functions can also be formed and are located both intra- and extracellularly. Distinct IGFBP-fragments can even be found in the perinuclear compartment or within the nucleus, where they can impact on gene expression. Several contributions presented in the current Research Topic particularly stress the relevance of structural aspects in IGFBP research. The current lack of comprehensive structural information is dramatically limiting the biomarker potential of particular IGFBPs. Finally, the Research Topic also provides novel functions of the IGFBP family from model organisms, farm animals and humans. Thereby, the biomarker potential not only relates to normal and malignant growth but also to metabolism and animal welfare. One important aim of the Research Topic is to encourage next generation IGFBP research reflecting subject-individual, conditional, and hormonal parameters but also structural aspects of the IGFBPs.

Mechanisms, thermodynamics and kinetics of ligand binding revealed from molecular simulations and machine learning Yinglong Miao, Weiliang Zhu, Chia-en A. Chang, J. Andrew McCammon, 2023-02-24

The Neuronal Functions of EF-hand Ca(2+)-binding Proteins 2nd Edition Michael R. Kreutz, José R. Naranjo, Karl-Wilhelm Koch, Beat Schwaller, 2016-11-01 Ca²⁺ signaling in neurons is characterized by highly restricted and dynamic gradients called Ca²⁺ waves, spikes, transients and puffs depending upon their corresponding spatial and temporal features. Based on this strict segmentation the Ca²⁺ ion provides a versatile basis for complex signaling in neuronal subcompartments with a spatial resolution of micro- and nanodomains. The multitude of Ca²⁺-regulated processes requires specialized downstream processing machinery, translating the Ca²⁺ signal into alterations of cellular processes. The broad range of different

Ca²⁺-triggered phenomena in neurons, ranging from neurotransmission to gene expression, is reflected by the existence of a multitude of different Ca²⁺-binding proteins (CaBPs) from which numerous belong to the EF-hand super-family. EF-hand proteins can be subdivided into Ca²⁺ buffer and Ca²⁺ sensor proteins. Whereas the first group has a very high affinity for Ca²⁺, exhibits little conformational change in the Ca²⁺-bound state and is thought to mainly chelate Ca²⁺, the second group has a lower affinity for Ca²⁺ and shows considerable conformational changes upon Ca²⁺-binding, which usually triggers a target interaction. Neuronal calcium sensor (NCS) proteins and the related Caldendrin/CaBP/Calneuron (nCaBPs) proteins are members of this latter group. They resemble the structure of their common ancestor Calmodulin (CaM) with four EF-hand Ca²⁺-binding motifs, of which not all are functional. However, despite their structural homology with CaM, NCS as well as nCaBPs are quite diverse in amino acid sequence. It is therefore surprising that relatively few binding partners have been identified that are not CaM targets and this raises the question of the specificity and function of these interactions. In terms of function, binding of NCS and nCaBP has frequently different consequences than binding of CaM, which substantially increases the versatility of the Ca²⁺ tool kit. The general idea of this special issue is to provide an overview on the function of neuronal EF-hand calcium-binding proteins in health and disease. But we will not just provide a mere collection of articles to stress the function of each protein. The issue will mainly deal with emerging concepts on Ca²⁺-signaling/buffering mediated by EF-hand Ca²⁺-binding proteins. This includes questions like features that define the functional role of a EF-hand calcium sensor in neurons, the conditions that make physiological relevance of a given interaction of a CaBP with its target plausible, the emerging synaptic role of these proteins, and mounting evidence for their role in the regulation of protein trafficking. Structural aspects and biophysical studies will be covered. Another aspect will be the role of CaBPs in brain disease states. This aspect includes studies showing that CaBPs are targets of drugs in clinical use, studies showing that expression levels of calcium-binding proteins are frequently altered in brain disease states as well as reports on mutations in EF-hand calcium sensors linked to human disease.

Cellular Fatty Acid-Binding Proteins II Jan F.C. Glatz, Ger J. van der Vusse, 2012-12-06 Twenty years have elapsed since cytoplasmic proteins exhibiting high-affinity binding of long-chain fatty acids were first identified (Ockner et al., Science 177:56-58, 1972). These cellular fatty acid-binding proteins (FABPs) are now well established to comprise a ligand-defined group of macromolecules belonging to a family of cytoplasmic lipid binding proteins. Unique features of the FABPs are the existence of distinct types of FABP and that these are found in a variety of tissues in remarkable abundance, with some cells expressing more than one type. The physiological significance of the FABPs has only partly been elucidated. By increasing the cytoplasmic solubilization of fatty acids, the cellular FABPs are considered to function primarily in intracellular fatty acid transport, but may also be assigned important regulatory roles in cellular lipid homeostasis as well as in the modulation of cell growth and differentiation. The broad interests in cellular FABPs has led to the organization of the 1st International Workshop on Fatty Acid-Binding Protein, held in Maastricht, the Netherlands, in 1989. Prompted by the success of the first meeting, the 2nd International Workshop on Fatty-Acid-Binding Proteins, which was held again in Maastricht, on August 31 and September 1, 1992, brought together scientific specialists in the field of FABP research for two days of intensive and fruitful discussion. This volume is a collection of selected papers from this conference, and thus provides the state-of-the-art knowledge of cellular FABPs. The contributors to this issue represent pioneering as well as new investigators, and also reflect the multidisciplinary nature of research in this exciting and rapidly progressing field.

POWER7 and POWER7+ Optimization and Tuning Guide Brian Hall, Mala Anand, Bill Buros, Miso Cilimdžić, Hong Hua, Judy Liu, John MacMillan, Sudhir Maddali, K Madhusudanan, Bruce Mealey, Steve Munroe, Francis P O'Connell, Sergio Reyes, Raul Silvera, Randy Swanberg, Brian Twichell, Brian F Veale, Julian Wang, Yaakov Yaari, IBM Redbooks, 2013-03-04 This IBM® Redbooks® publication provides advice and technical information about optimizing and tuning application code to run on systems that are based on the IBM POWER7® and POWER7+™ processors. This advice is drawn from application optimization efforts across many different types of code that runs under the IBM AIX® and Linux operating systems, focusing on the more pervasive performance opportunities that are identified, and how to capitalize on them. The technical information was developed by a set of domain experts at IBM. The focus of this book is to gather the right technical information, and lay out simple guidance for optimizing code performance on the IBM POWER7 and POWER7+ systems that run the AIX or Linux operating systems. This book contains a large amount of straightforward performance optimization that can be performed with minimal effort and without previous experience or in-depth knowledge. This optimization work can: Improve the performance of the application that is being optimized for the POWER7 system Carry over improvements to systems that are based on related processor chips Improve performance on other platforms The audience of this book is those personnel who are responsible for performing migration and implementation activities on IBM POWER7-based servers, which includes system administrators, system architects, network administrators, information architects, and database administrators (DBAs).

Tuning Autophagy-Inducing Activity and Toxicity for Lanthanide Nanocrystals Yunjiao Zhang, 2022-01-09 This thesis presents a simple, yet highly effective surface engineering solution that uses non-covalent binding peptides to control the autophagy-inducing activity of nanomaterials and nanodevices. The author presents RE-1, a short synthetic peptide that sequence-specifically binds to lanthanide (LN) oxide and upconversion

nanocrystals with high affinity, which was discovered using an innovative phage display approach. RE-1 effectively inhibits the autophagy-inducing activity and toxicity of these nanocrystals by forming a stable coating layer on the surface of the nanoparticles, and by reducing their sedimentation and cell interaction. RE- 1 and its variants provide a versatile tool for tuning cell interactions in order to achieve the desired level of autophagic response and are useful for the various diagnostic and therapeutic applications of LN-based nanomaterials and nanodevices.

Molecular Biology Nancy L. Craig,Rachel R. Green,Carol C. Greider,Cynthia Wolberger,Gisela G. Storz,2021-01-20 A fresh, distinctive approach to the teaching of molecular biology. With its focus on key principles, its emphasis on the commonalities that exist between the three kingdoms of life, and its integrated coverage of experimental methods and approaches, Molecular Biology is the perfect companion to any molecular biology course.

Tied and Bound: a Comparative View on Manuscript Binding Alessandro Bausi,Michael Friedrich,2023-08-07 The present volume contains twelve chapters authored by specialists of Asian, African and European manuscript cultures reflecting on the cohesion of written artefacts, particularly manuscripts. Assuming that 'codicological units' exist in every manuscript culture and that they are usually composed of discrete elements (such as clay tablets, papyrus sheets, bamboo slips, parchment bifolios, palm leaves), the issue of the cohesion of the constituents is a general one. The volume presents a series of case studies on devices and strategies adopted to achieve this cohesion by manuscript cultures distant in space (from China to West Africa) and time (from the third millennium bce to the present). This comparative view provides the frame for the understanding of a phenomenon that appears to be of essential importance for the study of the structure of written artefacts. Regardless of the way in which cohesion is realised, all strategies and devices that allow the constituents to be kept together are subsumed under the term 'binding'. Thus, it is possible to highlight similarities, convergences, and unique physical and technical methods adopted by various manuscript cultures to face a common challenge.

Oracle Performance Tuning for 10gR2 Gavin JT Powell,2011-04-08 Tuning of SQL code is generally cheaper than changing the data model. Physical and configuration tuning involves a search for bottlenecks that often points to SQL code or data model issues. Building an appropriate data model and writing properly performing SQL code can give 100%+ performance improvement. Physical and configuration tuning often gives at most a 25% performance increase. Gavin Powell shows that the central theme of Oracle10gR2 Performance Tuning is four-fold: denormalize data models to fit applications; tune SQL code according to both the data model and the application in relation to scalability; create a well-proportioned physical architecture at the time of initial Oracle installation; and most important, mix skill sets to obtain the best results. Fully updated for version 10gR2 and provides all necessary transition material from version 9i Includes all three aspects of Oracle database tuning: data model tuning, SQL & PL/SQL code tuning, physical plus configuration tuning Contains experienced guidance and real-world examples using large datasets Emphasizes development as opposed to operating system perspective

Workload Optimized Systems: Tuning POWER7 for Analytics Dino Quintero,Thomas Ancel,Graeme Cassie,Rodrigo Ceron,Amr Darwish,Guilherme Felix,Jian Jun He,Bharathraj Keshavamurthy,Sreenivas Makineedi,Girish Nikalje,Sudipto Pal,Zubair Salie,Ashu Tiwary,IBM Redbooks,2013-04-14 This IBM® Redbooks® publication addresses topics to help clients to take advantage of the virtualization strengths of the POWER® platform to solve system resource utilization challenges and maximize system throughput and capacity. This publication examines the tools, utilities, documentation, and other resources available to help technical teams provide business solutions and support for Cognos® Business Intelligence (BI) and Statistical Package for the Social Sciences (SPSS®) on Power Systems™ virtualized environments. This book addresses topics to help address complex high availability requirements, help maximize the availability of systems, and provide expert-level documentation to the worldwide support teams. This book strengthens the position of the Cognos and SPSS solutions with a well-defined and documented deployment model within a POWER system virtualized environment. This model provides clients with a planned foundation for security, scaling, capacity, resilience, and optimization. This book is targeted toward technical professionals (BI consultants, technical support staff, IT Architects, and IT Specialists) who are responsible for providing Smart Analytics solutions and support for Cognos and SPSS on Power Systems.

Molecular Biology Nancy Lynn Craig,Rachel Green,Orna Cohen-Fix,Carol Greider,Gisela Storz,2014 Molecular Biology: Principles of Genome Function offers a fresh, distinctive approach to the teaching of molecular biology. It is an approach that reflects the challenge of teaching a subject that is in many ways unrecognizable from the molecular biology of the 20th century - a discipline in which our understanding has advanced immeasurably, but about which many intriguing questions remain to be answered. It is written with severalguiding themes in mind: - A focus on key principles provides a robust conceptual framework on which students can build a solid understanding of the discipline; - An emphasis on thecommonalities that exist between the three kingdoms of life, and the discussion of differences between the three kingdoms where such differences offer instructive insights into molecular processes and components, gives students an accurate depiction of our current understanding of the conserved nature of molecular biology, and the differences that underpin biological diversity; - An integrated approach demonstrates how certain molecular phenomena have diverse impacts on genomefunction by presenting them as themes that recur throughout the book, rather than as artificially separated topics At heart, molecular biology is an experimental science, and a centralelement to the understanding of molecular biology is an appreciation of the approaches taken to yield the

information from which concepts and principles are deduced. Yet there is also the challenge of introducing the experimental evidence in a way that students can readily comprehend. Molecular Biology responds to this challenge with Experimental Approach panels, which branch off from the text in a clearly-signposted way. These panels describe pieces of research that have been undertaken, and which have been particularly valuable in elucidating difference aspects of molecular biology. Each panel is carefully cross-referenced to the discussion of key molecular biology tools and techniques, which are presented in a dedicated chapter at the end of the book. Beyond this, Molecular Biology further enriches the learning experience with full-colour, custom-drawn artwork; end-of-chapter questions and summaries; relevant suggested further readings grouped by topic; and an extensive glossary of key terms. Among the students being taught today are the molecular biologists of tomorrow; these individuals will be in a position to ask fascinating questions about fields whose complexity and sophistication become more apparent with each year that passes. Molecular Biology: Principles of Genome Function is the perfect introduction to this challenging, dynamic, but ultimately fascinating discipline.

Oracle PL/SQL Tuning Timothy S. Hall, 2006 Oracle experts know that PL/SQL tuning makes a huge difference in execution speed. As one of the world's most popular and respected experts, Dr. Tim Hall shares his secrets for tuning Oracle PL/SQL. This indispensable book shows how to hypercharge Oracle applications gaining as much as 30x improvement in execution speed using under-documented code tricks. Packed with working examples, learn how to re-write SQL into PL/SQL and how to use advanced Oracle bulk array processing techniques to achieve super high performance. You can save your company millions of dollars in hardware costs by making your applications run at peak efficiency. Targeted at the Senior Oracle DBA and developer, this advanced book illustrates powerful techniques that can make PL/SQL run faster than ever before. This book is not for beginners and should only be purchased by seasoned Oracle professionals who must turbocharge their applications. Your time savings from a single script is worth the price of this great book.

Oracle Database 11g Performance Tuning Recipes Sam Alapati, Darl Kuhn, Bill Padfield, 2011-10-18 Performance problems are rarely problems per se. They are more often crises during which you're pressured for results by a manager standing outside your cubicle while your phone rings with queries from the help desk. You won't have the time for a leisurely perusal of the manuals, nor to lean back and read a book on theory. What you need in that situation is a book of solutions, and solutions are precisely what Oracle Database 11g Performance Tuning Recipes delivers. Oracle Database 11g Performance Tuning Recipes is a ready reference for database administrators in need of immediate help with performance issues relating to Oracle Database. The book takes an example-based approach, wherein each chapter covers a specific problem domain. Within each chapter are recipes, showing by example how to perform common tasks in that chapter's domain. Solutions in the recipes are backed by clear explanations of background and theory from the author team. Whatever the task, if it's performance-related, you'll probably find a recipe and a solution in this book. Provides proven solutions to real-life Oracle performance problems Offers relevant background and theory to support each solution Written by a team of experienced database administrators successful in their careers

Actin-Monomer-Binding Proteins Pekka Lappalainen, 2008-05-14 This book provides a comprehensive view on actin monomer-binding proteins and the mechanisms by which they contribute to actin dynamics and various actin-dependent cellular processes. This new title contains chapters that describe the basic mechanisms of actin dynamics as well as the structural principles by which various actin-binding proteins interact with actin. The book is suitable further reading for students in advanced cell biology classes.

BIOS Instant Notes in Medicinal Chemistry Graham Patrick, 2019-10-10 Instant Notes in Medicinal Chemistry provides concise coverage for undergraduates studying medicinal chemistry as part of a science, pharmacy or medical course. It is a truly multidisciplinary subject involving such subject specialities as organic chemistry, pharmacology, biochemistry, physiology, microbiology, toxicology, genetics and computer mod

Drum Tuning, 2002 Miscellaneous Percussion Music - Mixed Levels

Regulation of Intracellular Ca²⁺-release Channels by Ca²⁺ and Ca²⁺-binding Proteins Nael Nadif Kasri, 2004 This is a Ph.D. dissertation. Almost everything we do is controlled by Ca²⁺, how we move, how our hearts beat and how our brains process information and store memories. Ca²⁺ is indeed a ubiquitous second messenger involved in a diversity of cellular activities such as muscular contraction, neurotransmitter release, synaptic plasticity, gene transcription, cell proliferation and apoptosis. The average cytosolic concentration of Ca²⁺ in resting cells ranges from 50nM to 100nM, reaching values close to 1uM when the cell is stimulated by a variety of physiological stimuli, while in the extracellular milieu this concentration is about 1mM. This large concentration gradient allows intracellular Ca²⁺ to work as a useful second messenger. Contents include: General Introduction, Objectives, Results, General Discussion, Summary.

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