

Mviewer

Cang Hui, Nick Isaac, Quentin Groom, Vernon Visser, Sandra MacFadyen

Computer Networks Andrzej Kwiecien, Piotr Gaj, Piotr Stera, 2013-05-27 This book constitutes the refereed proceedings of the 20th International Conference on Computer Networks, CN 2013, held in Lwówek Śląski, Poland, in June 2013. The 58 revised full papers presented were carefully reviewed and selected for inclusion in the book. The papers in these proceedings cover the following topics: computer networks, network architectural issues, Internet and wireless solutions, teleinformatics and communications, new technologies, queueing theory and queueing networks, innovative applications, networking in e-business, security aspects of hardware and software, industrial systems, quantum and bio-informatics, cloud networking and services.

Emerging Technologies in Computing Mahdi H. Miraz, Garfield Southall, Maaruf Ali, Andrew Ware, 2023-01-25 This book constitutes the refereed conference proceedings of the 5th EAI International Conference on Emerging Technologies in Computing, iCETiC 2022, held in Chester, UK, August 15-16, 2022. The 16 full papers included in this book were carefully reviewed and selected from 54 submissions. They were organized in topical sections as follows: Cloud, IoT and Distributed Computing; and AI, Expert Systems and Big Data Analytics.

Multi-Agent Oriented Programming Olivier Boissier, Rafael H. Bordini, Jomi Hubner, Alessandro Ricci, 2020-09-15 The main concepts and techniques of multi-agent oriented programming, which supports the multi-agent systems paradigm at the programming level. A multi-agent system is an organized ensemble of autonomous, intelligent, goal-oriented entities called agents, communicating with each other and interacting within an environment. This book introduces the main concepts and techniques of multi-agent oriented programming, (MAOP) which supports the multi-agent systems paradigm at the programming level. MAOP provides a structured approach based on three integrated dimensions, which the book examines in detail: the agent dimension, used to design the individual (interacting) entities; the environment dimension, which allows the development of shared resources and connections to the real world; and the organization dimension, which structures the interactions among the autonomous agents and the shared environment.

Graphics programming with GDI+ and DirectX Alex Polyakov, Vitaly Brusentsev, 2005 Providing a look ahead at some of the most promising innovations in graphics technologies, this guide covers the GDI+ graphical library, the .NET Framework platform, C#, and the DirectX library. Theoretical aspects of graphics, algorithms for solving common problems, and examples of their practical implementation are covered. The support of popular graphic file formats, such as BMP, GIF, TIFF, JPEG, EXI, PNG, ICO, WMF, and EMF, as well as bitmap redrawing with the application of the external alpha channel are discussed. Resizing, stretching, distorting, and rotating bitmaps are addressed in detail, as is gradient filling.

Журнал «Хакер» No08/2014, 2017-05-20 «Хакер» – это культовый журнал для тех, кто страстно увлечен современными технологиями. Для тех, кто хочет развиваться в IT или просто быть на острие. В каждом номере: подробные HOWTO, практические материалы по разработке и администрированию, интервью с выдающимися людьми, создавших технологические продукты и известные IT-компании, и, конечно, экспертные статьи о хакерстве и информационной безопасности. Мы предельно открыто пишем о существующих проблемах, рассказывая, как их могут использовать злоумышленники. При этом легкость изложения, даже невероятно сложных тем, – наш конек. У издания нет аналогов ни в России, ни в мире. В номере: Натягиваем Blizzard Warden Лишаем зрения античит-систему многопользовательских игр Легалайз в компьютере Избавляемся от пиратского ПО: лицензией, freeware, огнем и мечом История большого поиска С чего начинались продукты Google Живой в мире мертвых Как остаться анонимным при использовании смартфона Шелл-коды под ARM Учимся детектировать вредоносные нагрузки для популярной платформы Математика для программиста Изучать или забивать? Что

нового в NewSQL? Погружение в новейшие базы данных и многое другое

Computational Epigenomics and Epitranscriptomics Pedro H. Oliveira, 2023-02-01 This volume details state-of-the-art computational methods designed to manage, analyze, and generally leverage epigenomic and epitranscriptomic data. Chapters guide readers through fine-mapping and quantification of modifications, visual analytics, imputation methods, supervised analysis, and integrative approaches for single-cell data. Written in the highly successful *Methods in Molecular Biology* series format, chapters include introductions to their respective topics, lists of the necessary materials and reagents, step-by-step, readily reproducible laboratory protocols, and tips on troubleshooting and avoiding known pitfalls. Cutting-edge and thorough, *Computational Epigenomics and Epitranscriptomics* aims to provide an overview of epigenomic protocols, making it easier for researchers to extract impactful biological insight from their data.

Introduction to Data Mining and Analytics Kris Jamsa, 2020-02-03 Data Mining and Analytics provides a broad and interactive overview of a rapidly growing field. The exponentially increasing rate at which data is generated creates a corresponding need for professionals who can effectively handle its storage, analysis, and translation.

Parallel Processing and Applied Mathematics Roman Wyrzykowski, Jack Dongarra, Ewa Deelman, Konrad Karczewski, 2023-04-27 This two-volume set, LNCS 13826 and LNCS 13827, constitutes the proceedings of the 14th International Conference on Parallel Processing and Applied Mathematics, PPAM 2022, held in Gdansk, Poland, in September 2022. The 77 regular papers presented in these volumes were selected from 132 submissions. For regular tracks of the conference, 33 papers were selected from 62 submissions. The papers were organized in topical sections named as follows: Part I: numerical algorithms and parallel scientific computing; parallel non-numerical algorithms; GPU computing; performance analysis and prediction in HPC systems; scheduling for parallel computing; environments and frameworks for parallel/cloud computing; applications of parallel and distributed computing; soft computing with applications and special session on parallel EVD/SVD and its application in matrix computations. Part II: 9th Workshop on Language-Based Parallel Programming (WLPP 2022); 6th Workshop on Models, Algorithms and Methodologies for Hybrid Parallelism in New HPC Systems (MAMHYP 2022); first workshop on quantum computing and communication; First Workshop on Applications of Machine Learning and Artificial Intelligence in High Performance Computing (WAML 2022); 4th workshop on applied high performance numerical algorithms for PDEs; 5th minisymposium on HPC applications in physical sciences; 8th minisymposium on high performance computing interval methods; 7th workshop on complex collective systems.

Handbook of Fire and the Environment Brian J. Meacham, Margaret McNamee, 2022-07-28 The fundamental purpose of this handbook is to raise awareness about environmental impacts of fire and fire suppression, primarily within the fire engineering and firefighting communities, but also within the environmental engineering and planning disciplines. The Handbook provides readers with a fundamental understanding of the problem and its magnitude and includes a set of tools and methods for assessing environmental, social and financial impacts, and a set of tools for identifying and selecting appropriate mitigation options.

Applications Interface Programming Using Multiple Languages Ying Bai, 2003 Annotation This book provides a detailed description about the practical considerations in multiple languages programming as well as the interfaces among different languages in the Window environment. Authentic examples and detailed explanations are combined together in this book to provide the readers a clear picture as how to handle the multiple languages programming in Windows.

Structural Health Monitoring 2015 Fu-Kuo Chang ,Fotis Kopsaftopoulos,2015-10-01 Proceedings of the Tenth International Workshop on Structural Health Monitoring, September 1-3, 2015. Selected research on the entire spectrum of structural health techniques and areas of application Available in print, complete online text download or individual articles. Series book comprising two volumes provides selected international research on the entire spectrum of structural health monitoring techniques used to diagnose and safeguard aircraft, vehicles, buildings, civil infrastructure, ships and railroads, as well as their components such as joints, bondlines, coatings and more. Includes special sections on system design, signal processing, multifunctional materials, sensor distribution, embedded sensors for monitoring composites, reliability and applicability in extreme environments. The extensive contents can be viewed below.

High-Performance Computational Solutions in Protein Bioinformatics Dariusz Mrozek,2014-06-04 Recent developments in computer science enable algorithms previously perceived as too time-consuming to now be efficiently used for applications in bioinformatics and life sciences. This work focuses on proteins and their structures, protein structure similarity searching at main representation levels and various techniques that can be used to accelerate similarity searches. Divided into four parts, the first part provides a formal model of 3D protein structures for functional genomics, comparative bioinformatics and molecular modeling. The second part focuses on the use of multithreading for efficient approximate searching on protein secondary structures. The third and fourth parts concentrate on finding 3D protein structure similarities with the support of GPUs and cloud computing. Parts three and four both describe the acceleration of different methods. The text will be of interest to researchers and software developers working in the field of structural bioinformatics and biomedical databases.

Beyond Databases, Architectures and Structures. Facing the Challenges of Data Proliferation and Growing Variety Stanisław Kozielski,Dariusz Mrozek,Paweł Kasprowski,Bożena Małysiak-Mrozek,Daniel Kostrzewa,2018-09-07 This book constitutes the refereed proceedings of the 14th International Conference entitled Beyond Databases, Architectures and Structures, BDAS 2018, held in Poznań, Poland, in September 2018, during the IFIP World Computer Congress. It consists of 38 carefully reviewed papers selected from 102 submissions. The papers are organized in topical sections, namely big data and cloud computing; architectures, structures and algorithms for efficient data processing; artificial intelligence, data mining and knowledge discovery; text mining, natural language processing, ontologies and semantic web; image analysis and multimedia mining.

Beyond Databases, Architectures and Structures. Towards Efficient Solutions for Data Analysis and Knowledge Representation Stanisław Kozielski,Dariusz Mrozek,Paweł Kasprowski,Bożena Małysiak-Mrozek,Daniel Kostrzewa,2017-05-16 This book constitutes the refereed proceedings of the 13th International Conference entitled Beyond Databases, Architectures and Structures, BDAS 2017, held in Ustroń, Poland, in May/June 2017. It consists of 44 carefully reviewed papers selected from 118 submissions. The papers are organized in topical sections, namely big data and cloud computing; artificial intelligence, data mining and knowledge discovery; architectures, structures and algorithms for efficient data processing; text mining, natural language processing, ontologies and semantic web; bioinformatics and biological data analysis; industrial applications; data mining tools, optimization and compression.

Scalable Big Data Analytics for Protein Bioinformatics Dariusz Mrozek,2018-09-25 This book presents a focus on proteins and their structures. The text describes various scalable solutions for protein structure similarity searching, carried out at main representation levels and for prediction of 3D structures of proteins. Emphasis is placed on techniques that can be used to accelerate similarity searches and protein structure modeling processes.

The content of the book is divided into four parts. The first part provides background information on proteins and their representation levels, including a formal model of a 3D protein structure used in computational processes, and a brief overview of the technologies used in the solutions presented in the book. The second part of the book discusses Cloud services that are utilized in the development of scalable and reliable cloud applications for 3D protein structure similarity searching and protein structure prediction. The third part of the book shows the utilization of scalable Big Data computational frameworks, like Hadoop and Spark, in massive 3D protein structure alignments and identification of intrinsically disordered regions in protein structures. The fourth part of the book focuses on finding 3D protein structure similarities, accelerated with the use of GPUs and the use of multithreading and relational databases for efficient approximate searching on protein secondary structures. The book introduces advanced techniques and computational architectures that benefit from recent achievements in the field of computing and parallelism. Recent developments in computer science have allowed algorithms previously considered too time-consuming to now be efficiently used for applications in bioinformatics and the life sciences. Given its depth of coverage, the book will be of interest to researchers and software developers working in the fields of structural bioinformatics and biomedical databases.

Administrasi MongoDB Agus Kurniawan, Buku ini membahas secara singkat dan ringkas serta praktis mengenai manajemen administrasi MongoDB. MongoDB adalah salah satu platform NoSQL. Topik yang dibahas meliputi: * Mengenal MongoDB * Sintak MongoDB Shell * Manajemen MongoDB * Export & Import Data * Backup & Restore

Biodiversity Informatics: Building a Lifeboat for High Functionality Data to Decision Pipeline Cang Hui, Nick Isaac, Quentin Groom, Vernon Visser, Sandra MacFadyen, 2024-03-06

ARTnews ,1991 Vols. for 19--1949/50 include: Art news annual (title varies slightly). issued as a separate section of a regular number; 195--1959 issued as a separate volume.

Aperture, Getting Started Apple Computer, Inc, 2006

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Understanding mass balance for food compliance Nov 6,

2022 – Mass balance, in relationship to food production, can be defined as being the ability to account for all quantities of raw materials, waste, ... Tolerance on Mass Balance for Recall/withdrawal for BRC Aug 3, 2016 – Tolerance on Mass Balance for Recall/withdrawal for BRC – posted in BRCGS ... For example, if you have used 100 Kg of raw materials and 1000 donut ... BRC Auditing – What To Expect Under Food Issue 8 Oct 17, 2019 – The mass balance is the quantity of incoming raw material against the quantity used in the resulting finished products, taking process waste and ... The Mass Balance Approach in Feedstock Substitution An established method to foster sustainability in existing infrastructure · Benefits of the Mass Balance Approach · Biomass balance and ChemCycling · ChemCycling ... 8. Mass Balance Mass-balance analysis may also be referred to as. “Material Flow Analysis” or “Substance Flow Analysis.” Table 8.1 provides several examples of possible inputs,. Mass Balance Approach in the Chemical Industry The mass balance Approach (MBA) is a process for determining the use of chemically recycled or bio-based feedstock in a final product when both recycled and ... BRC 3.9.2 Trace Exercise Sample Procedure to conduct a mass balance check · 1. Select a raw material lot number used in a finished product made within the last 6 months. · 2. Review storage ... UNDERSTANDING VULNERABILITY ASSESSMENT Table 6 provides examples of PRNs for different raw materials. Table 6 Priority ... Mass balance exercises at critical points in the supply chain – the mass ... ISSUE 8 FOOD SAFETY – Frequently Asked Questions – a worked example from the raw material supplier, which ... to conduct a mass balance test every 6 months for each claim or a single mass balance test every. 4000 Years of Christmas: A Gift from the Ages it is an excellent publiucation showing the origins of many Christmas traditions. This includes originally pagan customs that were later Christianized, with the ... 4000 Years of Christmas: A Gift from the Ages A detailed look at the origins of Christmas celebrations ranges from before Jesus's birth and includes Rome's pagan

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