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Physics 702: Torque and Rotational Equilibrium | Georgia ... Note Taking Guide – Episode 702 PHYSICS Fundamentals © 2004, GPB 7-05 When an object is in rotational equilibrium: Ex. Problem #1: Rotational Equilibrium A uniform seesaw is 3.2 m long, with its pivot at the center. A girl weighing 580 N sits at one end. Where would a boy weighing 670 N need to sit to balance the seesaw?

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Problem : Find the % by mass of oxygen in water. $(16/18) * 100 = 88.9\%$ Percentage by mass of element in a compound = $(\text{mass of element in 1 mol of compound} \div \text{molar mass of compound}) * 100\%$ (after completing lab #1)
Find the % of carbon in sodium bicarbonate (NaHCO_3).

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CHEMISTRY: A Study of Matter © 2004, GPB 7.15 Example Problem: Find the % by mass of oxygen in

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Example Problem: Find the % by mass of oxygen in water.

Percentage by mass of element in a compound = (mass of element in 1 mol of compound ÷ molar mass of compound) x 100 %.

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As the United Nations Decade on Biodiversity 2011 – 2020 comes to a close and countries prepare to adopt a post-2020 global biodiversity framework, this edition of The State of the World ' s Forests (SOFO) examines the contributions of forests, and of the people who use and manage them, to the conservation and sustainable use of biodiversity. Forests cover just over 30 percent of the global

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land area, yet they provide habitat for the vast majority of the terrestrial plant and animal species known to science. Unfortunately, forests and the biodiversity they contain continue to be under threat from actions to convert the land to agriculture or unsustainable levels of exploitation, much of it illegal. The State of the World ' s Forests 2020 assesses progress to date in meeting global targets and goals related to forest biodiversity and examines the effectiveness of policies, actions and approaches, in terms of both conservation and sustainable development outcomes. A series of case studies provide examples of innovative practices that combine conservation and sustainable use of forest biodiversity to create

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balanced solutions for both people and the planet.

A how-to guide for assessing the impact of fiscal policy on inequality and poverty Inequality has emerged in recent years as a major topic of economic and political discussion, but it is often unclear whether governments can or should do something about it, and if so, what that something might be. This unique volume, edited by Nora Lustig, an equity expert at Tulane University, helps fill that void. Developed by the Commitment to Equity Institute at Tulane, the book examines both the theory and the practical methods for determining the impact of taxation and public spending on inequality and

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poverty. It provides a step-by-step guide for policymakers, economists, and social planners when analyzing whether fiscal policy has narrowed or widened inequality. The book also has user-written software for conducting a Commitment to Equity Assessment, along with several country studies of these assessments. In addition to serving as a manual, the book can be used as a stand-alone reference for those interested in the methods for assessing the impact on equity of fiscal policy. It also serves as a textbook for advanced undergraduate and graduate courses on public finance and income distribution.

The Paralysis Resource Guide,

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produced by the Christopher & Dana Reeve Foundation, is a reference and lifestyle tool for people affected by paralysis. The book includes details on medical and clinical subjects related to all causes of paralysis, as well as health maintenance information. The fully-illustrated book provides a detailed overview of biomedical research, assistive technology, sports and recreation activities, legal and civil rights, social security and benefits, and numerous lifestyle options.

Since the first attempts at structure-based drug design about four decades ago, molecular modelling techniques for drug design have developed enormously, along with the

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increasing computational power and structural and biological information of active compounds and potential target molecules. Nowadays, molecular modeling can be considered to be an integral component of the modern drug discovery and development toolbox. Nevertheless, there are still many methodological challenges to be overcome in the application of molecular modeling approaches to drug discovery. The eight original research and five review articles collected in this book provide a snapshot of the state-of-the-art of molecular modeling in drug design, illustrating recent advances and critically discussing important challenges. The topics covered include virtual screening and

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pharmacophore modelling, chemoinformatic applications of artificial intelligence and machine learning, molecular dynamics simulation and enhanced sampling to investigate contributions of molecular flexibility to drug – receptor interactions, the modeling of drug – receptor solvation, hydrogen bonding and polarization, and drug design against protein – protein interfaces and membrane protein receptors.

The first notable feature of this book is its innovation: Computational intelligence (CI), a fast evolving area, is currently attracting lots of researchers' attention in dealing with many complex problems. At present, there are quite a lot competing

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books existing in the market. Nevertheless, the present book is markedly different from the existing books in that it presents new paradigms of CI that have rarely mentioned before, as opposed to the traditional CI techniques or methodologies employed in other books. During the past decade, a number of new CI algorithms are proposed. Unfortunately, they spread in a number of unrelated publishing directions which may hamper the use of such published resources. These provide us with motivation to analyze the existing research for categorizing and synthesizing it in a meaningful manner. The mission of this book is really important since those algorithms are going to be a new revolution in

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computer science. We hope it will stimulate the readers to make novel contributions or even start a new paradigm based on nature phenomena. Although structured as a textbook, the book's straightforward, self-contained style will also appeal to a wide audience of professionals, researchers and independent learners. We believe that the book will be instrumental in initiating an integrated approach to complex problems by allowing cross-fertilization of design principles from different design philosophies. The second feature of this book is its comprehensiveness: Through an extensive literature research, there are 134 innovative CI algorithms covered in this book.

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Recent advances in drug discovery have been rapid. The second edition of *Bioinformatics and Drug Discovery* has been completely updated to include topics that range from new technologies in target identification, genomic analysis, cheminformatics, protein analysis, and network or pathway analysis. Each chapter provides an extended introduction that describes the theory and application of the technology. In the second part of each chapter, detailed procedures related to the use of these technologies and software have been incorporated. Written in the highly successful *Methods in Molecular Biology*™ series format, the chapters include the kind of detailed description and implementation advice that is

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crucial for getting optimal results in the laboratory. Thorough and intuitive, *Bioinformatics and Drug Discovery, Second Edition* seeks to aid scientists in the further study of the rapidly expanding field of drug discovery.

This is the first book, after J. E. Sandys, to cover the multifaceted field of “ancient scholarship” from the beginnings to the fall of Byzantium. It is worth underlining the benefits of a work with multiple expert voices in a field so complex. The book is based on the four historiographical chapters of *Brill's Companion to Ancient Greek Scholarship* (2015), which have been updated and rethought.

Various antiseptic agents, such as

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chlorhexidine, are used for different applications, e.g. in healthcare, veterinary medicine, animal production and household products, including cosmetics. However, not all antiseptic agents provide significant health benefits, especially in some products used in human medicine (alcohol-based hand rubs, antimicrobial soaps). While some products (antimicrobial soaps, surface disinfectants, instrument disinfectants, wound antiseptics) may contain one or more biocidal agents with a comparable antimicrobial efficacy but large differences in their potential for microbial adaptation and tolerance. An increased bacterial resistance has been described for various antimicrobial agents, sometimes

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including a cross-resistance to antibiotics. The book is the first comprehensive reference resource on antiseptic agents, including their efficacy, natural and acquired resistance, adaptation, and cross-resistance. It also discusses their and appropriate use in terms of a balance between their efficacy and the risk of acquired bacterial resistance / tolerance. Focusing on human and veterinary medicine and household products, it helps readers make informed decisions concerning against antiseptic products based on their composition. The book contributes to reduce any unnecessary selection pressure towards emerging pathogens and to keep the powerful antiseptic agents for all those applications that have a

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clear benefit (e.g. reduction of healthcare-associated infection).

This book reviews the advances and challenges of structure-based drug design in the preclinical drug discovery process, addressing various diseases, including malaria, tuberculosis and cancer. Written by internationally recognized researchers, this edited book discusses how the application of the various in-silico techniques, such as molecular docking, virtual screening, pharmacophore modeling, molecular dynamics simulations, and residue interaction networks offers insights into pharmacologically active novel molecular entities. It presents a clear concept of the molecular mechanism of different

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drug targets and explores methods to help understand drug resistance. In addition, it includes chapters dedicated to natural-product-derived medicines, combinatorial drug discovery, the CryoEM technique for structure-based drug design and big data in drug discovery. The book offers an invaluable resource for graduate and postgraduate students, as well as for researchers in academic and industrial laboratories working in the areas of chemoinformatics, medicinal and pharmaceutical chemistry and pharmacoinformatics.

Radio Design in Nanometer Technologies is the first volume that looks at the integrated radio

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design problem as a "piece of a big puzzle", namely the entire chipset or single chip that builds an entire wireless system. This is the only way to successfully design radios to meet the stringent demands of today ' s increasingly complex wireless systems.

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