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Life Science Pfeb March 2015

**Ming Lei, Christopher L.-H. Huang, R.
John Solaro, Yunbo Ke**



Life Science Pfeb March 2015:

New Weapons to Control Bacterial Growth Tomas G. Villa, Miguel Vinas, 2016-03-24 Written by specialists in the different fields this book presents new perspectives and insights into strategies and weapons to fight microbial infections It also reviews the state of the art of alternative treatment approaches and new therapeutic agents to deal with infections caused by multidrug resistant microorganisms In an era of accumulated resistance to current antibiotics it is vital that this is undertaken without further delay Aspects discussed include the control of RNA synthesis the use of bacteriocins or enzybiotics bacteriophages or purified lysins the specific control of pathogenic clostridia the design of new drugs affecting DNA synthesis in bacteria the use of fecal matter transplant strategies the specific control of quorum sensing responses in bacteria the use of new peptides as antibiotics and new ways to control bacteria that cause cancer such as Helicobacter pylori cancers

The Cambridge Handbook of Information Technology, Life Sciences and Human Rights Marcello Ienca, Oreste Pollicino, Laura Liguori, Elisa Stefanini, Roberto Andorno, 2022-05-26 Debates on the human rights implications of new and emerging technologies have been hampered by the lack of a comprehensive theoretical framework for the complex issues involved This volume provides that framework bringing a multidisciplinary and international perspective to the evolution of human rights in the digital and biotechnological era It delves into the latest frontiers of technological innovation in the life sciences and information technology sectors such as neurotechnology robotics genetic engineering and artificial intelligence Leading experts from the technological medical and social sciences as well as law philosophy and business share their extensive knowledge about the transformation of the rights framework in response to technological innovation In addition to providing a comprehensive interdisciplinary and international state of the art descriptive analysis the volume also offers policy recommendations to protect and promote human rights in the context of emerging socio technological trends

Exhausted Nick Polizzi, Pedram Shojai, OMD, 2021-08-03 New in paperback from the New York Times best selling author of *The Urban Monk* and author of *The Sacred Science* draw a clear and thorough road map for healing from exhaustion inside and out We re facing an unprecedented healthcare crisis and much of it stems from the collapse of the energy production systems of our cells Exhausted is a timely book that sheds light on the root of this epidemic Mark Hyman M D New York Times best selling author of *Food Eat Fat Get Thin* and *The Blood Sugar Solution* The economics are simple you can t run on a deficit of energy and expect to feel well Yet that is what we ve come to expect from ourselves as we have pushed farther and farther down the hole of fatigue and away from any semblance of being healthy But what if you had a simple road map to reclaiming your health and vibrancy Nick Polizzi author of *The Sacred Science* and Pedram Shojai O M D and New York Times best selling author of *The Urban Monk* and *The Art of Stopping Time* provide Wisdom from the top scientists physicians and experts in traditional healing practices A concise nuts and bolts understanding of what energy is and how we use it An exploration of the key areas in which we have been depriving our body of energy From our immune

system to hormones to sleep patterns Polizzi and Shojai offer methods for evaluating your individual needs as well as safe easy remedies for whole body healing Maximize the potential energy from the essential parts of your body and life so you can finally feel fully alive and find the fulfillment you deserve both personally and professionally **Curiosity And Passion For Science And Art** Uwe B Sleytr, 2016-07-04 This book describes the accomplishments of a curious and imaginative scientist and his endeavours to translate or even to extrapolate scientific insights into the world of art The science section in this volume concerns studies on S layers a very important class of proteins found on the surface of numerous Bacteria and nearly all Archaea S layer proteins are one of the most abundant biopolymers on our planet and assemble into the simplest type of biological membrane Moreover they are unique building blocks and patterning elements for the production of complex supramolecular structures and nanoscale devices in nanobiotechnology molecular nanotechnology synthetic biology biomimetics and nanomedicine In the second part of this book the author goes on to passionately describe how his scientific activities stimulated his art work which in particular concerns the visualization of results and the potential of synthetic biology and evolutionary events induced by genetic manipulations Most importantly the engagement in art allowed him to leave the rather curtailed canon of science and reach a mental state of unlimited freedom of thoughts Mask like sculptures are used as examples to visualize the intersection between science and art and in particular the unpredictability and mystery of scientific visions *LuxR Solos Are Becoming Major Players in Cell-Cell Communication in Bacteria* Vittorio Venturi, Brian M.M. Ahmer, 2016-08-11 The most common quorum sensing QS system in Gram negative bacteria occurs via N acyl homoserine lactone AHLs signals An archetypical system consists of a LuxI family protein synthesizing the AHL signal which binds at quorum concentrations to the cognate LuxR family transcription factors which then control gene expression by binding to specific sequences in target gene promoters QS LuxR family proteins are approximately 250 amino acids long and made up of two domains at the N terminus there is an autoinducer binding domain whereas the C terminus contains a DNA binding helix turn helix HTH domain QS LuxRs display surprisingly low similarities 18-25% even if they respond to structurally similar AHLs 95% of LuxRs share 9 highly conserved amino acid residues six of these are hydrophobic or aromatic and form the cavity of the AHL binding domain and the remaining three are in the HTH domain With only very few exceptions the luxI R cognate genes of AHL QS systems are located adjacent to each other The sequencing of many bacterial genomes has revealed that many proteobacteria also possess LuxRs that do not have a cognate LuxI protein associated with them These LuxRs have been called orphans and more recently solos LuxR solos are widespread in proteobacterial species that possess a canonical complete AHL QS system as well as in species that do not In many cases more than one LuxR solo is present in a bacterial genome Scientists are beginning to investigate these solos Are solos responding to AHL signals If present in a bacterium which possesses a canonical AHL QS system are solos an integral part of the regulatory circuit Are LuxR solos eavesdropping on AHLs produced by neighboring bacteria Have they evolved to respond to different signals

instead of AHLs and are these signals endogenously produced or exogenously provided Are they involved in interkingdom signaling by responding to eukaryotic signals Recent studies have revealed that LuxR solos are involved in several mechanisms of cell cell communication in bacteria implicating them in bacterial intraspecies and interspecies communication as well as in interkingdom signaling by responding to molecules produced by eukaryotes LuxR solos are likely to become major players in signaling since they are widespread among proteobacterial genomes and because initial studies highlight their different roles in bacterial communication This Research Topic allows scientists studying or interested in LuxR solos to report their data and or express their hypotheses and thoughts on this important and currently understudied family of signaling proteins

Landscapes of Collectivity in the Life Sciences Snait B. Gissis,Ehud Lamm,Ayelet Shavit,2018-01-12 Broad perspective on collectivity in the life sciences from microorganisms to human consensus and the theoretical and empirical opportunities and challenges Many researchers and scholars in the life sciences have become increasingly critical of the traditional methodological focus on the individual This volume counters such methodological individualism by exploring recent and influential work in the life sciences that utilizes notions of collectivity sociality rich interactions and emergent phenomena as essential explanatory tools to handle numerous persistent scientific questions in the life sciences The contributors consider case studies of collectivity that range from microorganisms to human consensus discussing theoretical and empirical challenges and the innovative methods and solutions scientists have devised The contributors offer historical philosophical and biological perspectives on collectivity and describe collective phenomena seen in insects the immune system communication and human collectivity with examples ranging from cooperative transport in the longhorn crazy ant to the evolution of autobiographical memory They examine ways of explaining collectivity including case studies and modeling approaches and explore collectivity s explanatory power They present a comprehensive look at a specific case of collectivity the Holobiont notion the idea of a multi species collective a host and diverse microorganisms and the hologenome theory which posits that the holobiont and its hologenome are a unit of adaption The volume concludes with reflections on the work of the late physicist Eshel Ben Jacob pioneer in the study of collective phenomena in living systems Contributors Oren Bader John Beatty Dinah R Davison Daniel Dor Ofer Feinerman Raghavendra Gadagkar Scott F Gilbert Snait B Gissis Deborah M Gordon James Griesemer Zachariah I Grochau Wright Erik R Hanschen Eva Jablonka Mohit Kumar Jolly Anat Kolumbus Ehud Lamm Herbert Levine Arnon Levy Xue Fei Li Elisabeth A Lloyd Yael Lubin Eva Maria Luef Ehud Meron Richard E Michod Samir Okasha Simone Pika Joan Roughgarden Eugene Rosenberg Ayelet Shavit Yael Silver Alfred I Tauber Ilana Zilber Rosenberg

Research Handbook on Intellectual Property and the Life Sciences Duncan Matthews,Herbert Zech,2017-06-30 Intellectual property IP is a key component of the life sciences one of the most dynamic and innovative fields of technology today At the same time the relationship between IP and the life sciences raises new public policy dilemmas The Research Handbook on Intellectual Property and the Life Sciences comprises contributions by leading

experts from academia and industry to provide in depth analyses of key topics including pharmaceuticals diagnostics and genes plant innovations stem cells the role of competition law and access to medicines The Research Handbook focuses on the relationship between IP and the life sciences in Europe and the United States complemented by country specific case studies on Australia Brazil China India Japan Kenya South Africa and Thailand to provide a truly international perspective

Handbook of Software Solutions for ICME Georg J. Schmitz,Ulrich Prah,2016-09-20 As one of the results of an ambitious project this handbook provides a well structured directory of globally available software tools in the area of Integrated Computational Materials Engineering ICME The compilation covers models software tools and numerical methods allowing describing electronic atomistic and mesoscopic phenomena which in their combination determine the microstructure and the properties of materials It reaches out to simulations of component manufacture comprising primary shaping forming joining coating heat treatment and machining processes Models and tools addressing the in service behavior like fatigue corrosion and eventually recycling complete the compilation An introductory overview is provided for each of these different modelling areas highlighting the relevant phenomena and also discussing the current state for the different simulation approaches A must have for researchers application engineers and simulation software providers seeking a holistic overview about the current state of the art in a huge variety of modelling topics This handbook equally serves as a reference manual for academic and commercial software developers and providers for industrial users of simulation software and for decision makers seeking to optimize their production by simulations In view of its sound introductions into the different fields of materials physics materials chemistry materials engineering and materials processing it also serves as a tutorial for students in the emerging discipline of ICME which requires a broad view on things and at least a basic education in adjacent fields

Ca²⁺ Signaling and Heart Rhythm Ming Lei,Christopher L.-H. Huang,R. John Solaro,Yunbo Ke,2016-07-25 Ca²⁺ is a key second messenger in the intricate workings of the heart In cardiac myocytes Ca²⁺ signaling controls or modulates electrophysiological function excitation contraction coupling contractile function energy balance cell death and gene transcription Thus diverse Ca²⁺ dependent regulatory processes occur simultaneously within a cell Yet distinct signals can be resolved by local Ca²⁺ sensitive protein complexes and differential Ca²⁺ signal integration In addition to its importance to normal cardiac function such regulation is also crucial in disease conditions Ca²⁺ is likely involved in ectopic cardiac rhythms in both atrial and ventricular tissues through generating triggered activity often appearing as delayed afterdepolarisations particularly following cellular Ca overloading Recent studies also implicate Ca²⁺ in Na channel expression and properties with consequences for conduction velocity and therefore arrhythmic substrate At the cellular level such regulation involves control of the activity of membrane ion channels and Ca²⁺ handling proteins These in turn involve multiple extra and intracellular signaling pathways This e book assembles review and original articles from experts in this field It summarises major recent progress bearing on roles of Ca²⁺ in cardiac electrophysiological function encompassing both normal and

abnormal cardiac function These extend from physiological roles of Ca²⁺ signaling in pacemaker function in particular generation of sino atrial pacemaker potentials to pathological roles of abnormal Ca²⁺ signaling in both atrial and ventricular arrhythmogenesis It also seeks to bridge the gap between advances in basic science and development of new therapies

Amino Acids of the Glutamate Family: Functions beyond Primary Metabolism Sakiko Okumoto, Dietmar Funck, Maurizio Trovato, Giuseppe Forlani, 2016-10-07 The life of proteins starts and ends as amino acids In addition to the primary function as protein building blocks amino acids serve multiple other purposes to make a plant's life worth living This is true especially for the amino acids of the glutamate family namely glutamate Glu glutamine Gln proline Pro and arginine Arg as well as the product of Glu decarboxylation aminobutyric acid GABA Synthesis accumulation interconversion and degradation of these five compounds contribute in many ways to the regulation of plant development and to responses to environmental challenges Glu and Gln hold key positions as entry points and master regulators of nitrogen metabolism in plants and have a pivotal role in the regulatory interplay between carbon and nitrogen metabolism Pro and GABA are among the best studied compatible osmolytes that accumulate in response to water deficit yet the full range of protective functions is still to be revealed Arg with its exceptionally high nitrogen to carbon ratio has long been recognized as a major storage form of organic nitrogen Most of the enzymes involved in metabolism of the amino acids of the glutamate family in plants have been identified or can be predicted according to similarity with animal or microbial homologues However for some of these enzymes the detailed biochemical properties still remain to be determined in order to understand activities in vivo Additionally uncertainties regarding the subcellular localization of proteins and especially the lack of knowledge about intracellular transport proteins leave significant gaps in our understanding of the metabolic network connecting Glu Gln Pro GABA and Arg While anabolic reactions are distributed between the cytosol and chloroplasts catabolism of the amino acids of the glutamate family takes place in mitochondria and has been implicated in fueling energy demanding physiological processes such as root elongation recovery from stress bolting and pollen tube elongation Exceeding the metabolic functions the amino acids of the glutamate family were recently identified as important signaling molecules in plants Extracellular Glu GABA and a range of other metabolites trigger responses in plant cells that resemble the actions of Glu and GABA as neurotransmitters in animals Plant homologues of the Glu gated ion channels from mammals and protein kinase signaling cascades have been implicated in these responses Pollen tube growth and guidance depend on GABA signaling and the root architecture is specifically regulated by Glu GABA and Pro signaling or metabolism were shown to contribute to the orchestration of defense and programmed cell death in response to pathogen attacks Pro signaling was additionally proposed to regulate developmental processes and especially sexual reproduction Arg is tightly linked to nitric oxide NO production and signaling in plants although Arg dependent NO synthases could still not be identified Potentially Arg derived polyamines constitute the missing link between Arg and NO signaling in response to stress Taken together the amino acids of the

glutamate family emerge as important signaling molecules that orchestrate plant growth and development by integrating the metabolic status of the plant with environmental signals especially in stressful conditions This research topic collects contributions from different facets of glutamate family amino acid signaling or metabolism to bring together and integrate in a comprehensive view the latest advances in our understanding of the multiple functions of Glu derived amino acids in plants

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