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Foundation Tier Paper 2F

F

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As May June Biology 2014 Paper

Bernhard Ganss, Megan Pugach



As May June Biology 2014 Paper:

The Coevolution of IDO1 and AhR in the Emergence of Regulatory T Cells in Mammals Ursula Grohmann, Paolo Puccetti, 2016-01-28 Indoleamine 2,3-dioxygenase IDO1 is an ancestral enzyme that initially confined to the regulation of tryptophan availability in local tissue microenvironments is now considered to play a wider role that extends to homeostasis and plasticity of the immune system. Thus IDO biology has implications for many aspects of immunopathology including viral infections, neoplasia, autoimmunity and chronic inflammation. Its immunoregulatory effects are mainly mediated by dendritic cells (DCs) and involve not only tryptophan deprivation but also production of kynurenines that act on IDO DCs thus rendering an otherwise stimulatory DC capable of regulatory effects as well as on T cells. The aryl hydrocarbon receptor AhR is a ligand-operated transcription factor originally recognized as the effector mediating the pathologic effects of dioxins and other pollutants. However, it is now well established that AhR activation by endogenous ligands can produce immunoregulatory effects. The IDO1 mechanism appears to have been selected through phylogenesis primarily to prevent overreacting responses to TLR-recognized pathogen-associated molecular patterns and only later did it become involved in the response to T cell receptor-recognized antigens. As a result, in mammals IDO1 has become pivotal in fetomaternal tolerance at a time when regulatory T cells emerged to meet the same need, namely protecting the fetus. IDO1 and regulatory T (Treg) cells may have then coevolved to broaden their function well beyond their initial task of protecting the fetus such that in acquired immunity IDO1 with its dual enzymic and signaling function has turned into an important component of the peripheral generation and effector function of regulatory T cells. AhR in turn, which has a role in regulatory T cell generation, is presumed to have evolved from invertebrates where it served a ligand-independent role in normal development processes. Evolution of the receptor in vertebrates resulted in the ability to bind structurally different ligands including xenobiotics and microbiota-derived catabolites. Considering the inability of invertebrate AhR homologs to bind dioxins, the adaptive role of the AhR to act as a regulator of xenobiotic-metabolizing enzymes may have been a vertebrate innovation to later acquire an additional immune regulatory role by coevolutionary pressure in mammals by IDO1 and regulatory T cells. Thus an entirely new paradigm in immunology and more specifically in immune tolerance is the coevolution of three systems, namely the IDO1 mechanism, AhR-driven gene transcription, and T cell regulatory activity that originating from the initial need of protecting the fetus in mammals have later turned into a pivotal mechanism of peripheral tolerance in autoimmunity, transplantation and neoplasia.

Enamel Research: Mechanisms and Characterization Bernhard Ganss, Megan Pugach, 2016-11-23 The rodent incisor is a good model system to study the molecular and cellular events that are involved in enamel biomineralization. Incisors in rodents continuously erupt during their lifespan, thus allowing the study of all stages of enamel synthesis, deposition, mineralization and maturation in the same tissue section. This model system has provided invaluable insight into the specifics of enamel formation as a basis to understand human pathologies such as amelogenesis imperfecta. Furthermore

the rodent incisor allows exploration and understanding of some of the most fundamental mechanisms that govern biomineralization Enamel is the most mineralized hardest tissue in the body It is formed within a unique organic matrix that unlike other hard tissues such as bone and dentin does not contain collagen The formation of enamel can be divided into two main stages the secretory and maturation stage During the secretory stage a highly ordered arrangement of hydroxyapatite crystals is formed under the influence of structural matrix proteins such as amelogenin ameloblastin and enamelin During the maturation stage the organic matrix is removed and hydroxyapatite crystals expand to ultimately yield a functional hard structure consisting of over 96% mineral Research efforts over the past decades have mainly focused on the secretory stage providing novel insights into the concept of biomineralization However the events that occur during the maturation stage have not been yet explored in detail likely because the physiological roles of the enamel forming ameloblasts are more diverse and complex at this stage Mature ameloblasts are involved in the regulation of calcium transport in large amounts phosphate and protein fragments in and out of the maturing enamel and provide regulatory mechanisms for the control of the pH In recent years increased efforts have been dedicated towards defining the molecular events during enamel maturation The development of an ever increasing number of transgenic animal models has clearly demonstrated the essential roles of matrix and non matrix proteins during enamel formation Multiple traditional and modern analytical techniques are applied for the characterization of enamel in these animals The need for this Research Topic therefore stems from new information that has been generated on molecular events during the enamel maturation stage and the development and application of highly advanced analytical techniques to characterize dental enamel The benefits and limitations of these techniques need to be reviewed and their application standardized for valid comparative studies

Epigenetics as a Deep Intimate Dialogue between Host and Symbionts Ilaria Negri, Eva Jablonka, 2016-06-30 Symbiosis is an intimate relationship between different living entities and is widespread in virtually all organisms It was critical for the origin and diversification of Eukaryotes and represents a major driving force in evolution Indeed symbiosis may support a wide range of biological processes including those underlying the physiology development reproduction health behavior ecology and evolution of the organisms involved in the relationship Although often confused with mutualism when both organisms benefit from the association symbiosis actually encompasses several and variable relationships Among them is parasitism when one organism benefits but the other is harmed and commensalism when one organism benefits and the other remains unaffected Even if many symbiotic lifestyles do exist in nature in many cases the intimacy between the partners is so deep that the symbiont sensu strictu resides into the tissues and or cells of the other partner Since the partners frequently belong to different kingdoms e g bacteria fungi protists and viruses living in association with animal and plant hosts their shared language should be a basic and ancient form of communication able to effectively blur the boundaries between extremely different living entities In recent years studies on the role of epigenetics in shaping host symbiont interactions have been flourishing Epigenetic changes include but are not

limited to DNA methylation remodelling of chromatin structure through histone chemical modifications and RNA interference. In this E book we present a series of papers exploring the fascinating developmental and evolutionary relationship between symbionts and hosts by focusing on the mediating epigenetic processes that enable the communication to be effective and robust at both the individual, the ecological and the evolutionary time scales. In particular the papers consider the role of epigenetic factors and mechanisms in the interactions among different species comprising the holobiont and host-parasite relationships. On the whole, since epigenetics is fast acting and reversible, enabling dynamic developmental communication between hosts and symbionts at several different time scales, we argue that it could account for the enormous plasticity that characterizes the interactions between all the organisms living symbiotically on our planet. **Cyanobacteria: The Green**

E. coli Anne M Ruffing, Toivo Kallas, 2016-03-16 As the world struggles to reduce its dependence on fossil fuels and curb greenhouse gas emissions, industrial biotechnology is also going green. *Escherichia coli* has long been used as a model Gram-negative bacterium not only for fundamental research but also for industrial applications. Recently, however, cyanobacteria have emerged as candidate chassis for the production of commodity fuels and chemicals utilizing CO₂ and sunlight as the main nutrient requirements. In addition to their potential for reducing greenhouse gas emissions and lowering production costs, cyanobacteria have naturally efficient pathways for the production of metabolites such as carotenoids, which are of importance in the nutraceutical industry. The unique metabolic and regulatory pathways present in cyanobacteria present new challenges for metabolic engineers and synthetic biologists. Moreover, their requirement for light and the dynamic regulatory mechanisms of the diurnal cycle further complicate the development and application of cyanobacteria for industrial applications. Consequently, significant advancements in cyanobacterial engineering and strain development are necessary for the development of a green *E. coli*. This Research Topic will focus on cyanobacteria as organisms of emerging industrial relevance, including research focused on the development of genetic tools for cyanobacteria, the investigation of new cyanobacterial strains, the construction of novel cyanobacterial strains via genetic engineering, the application of omics tools to advance the understanding of engineered cyanobacteria, and the development of computational models for cyanobacterial strain development. **Dual-use life science research and biosecurity in the 21st Century: Social,**

Technical, Policy, and Ethical Challenges Jonathan E. Suk, Kathleen M. Vogel, Amanda Jane Ozin, 2015-06-04 In September 2011, scientists announced new experimental findings that would not only threaten the conduct and publication of influenza research but would have significant policy and intelligence implications. The findings presented a modified variant of the H5N1 avian influenza virus, hereafter referred to as the H5N1 virus, that was transmissible via aerosol between ferrets. These results suggested a worrisome possibility: the existence of a new airborne and highly lethal H5N1 virus that could cause a deadly global pandemic. In response, a series of international discussions on the nature of dual-use life science arose. These discussions addressed the complex social, technical, political, security, and ethical issues related to dual-use research.

This Research Topic will be devoted to contributions that explore this matrix of issues from a variety of case study and international perspectives Immune system modeling and analysis Ramit Mehr,Miles Davenport,2015-04-22 The rapid development of new methods for immunological data collection from multicolor flow cytometry through single cell imaging to deep sequencing presents us now for the first time with the ability to analyze and compare large amounts of immunological data in health aging and disease The exponential growth of these datasets however challenges the theoretical immunology community to develop methods for data organization and analysis Furthermore the need to test hypotheses regarding immune function and generate predictions regarding the outcomes of medical interventions necessitates the development of mathematical and computational models covering processes on multiple scales from the genetic and molecular to the cellular and system scales The last few decades have seen the development of methods for presentation and analysis of clonal repertoires those of T and B lymphocytes and phenotypic surface marker based repertoires of all lymphocyte types and for modeling the intricate network of molecular and cellular interactions within the immune systems This e Book which has first appeared as a Frontiers in Immunology research topic provides a comprehensive online open access snapshot of the current state of the art on immune system modeling and analysis **Recent Advances in Symbiosis Research: Integrative**

Approaches M. Pilar Francino,Mónica Medina,2017-02-02 Traditionally symbiosis research has been undertaken by researchers working independently of one another and often focused on a few cases of bipartite host symbiont interactions New model systems are emerging that will enable us to fill fundamental gaps in symbiosis research and theory focusing on a broad range of symbiotic interactions and including a variety of multicellular hosts and their complex microbial communities In this Research Topic we invited researchers to contribute their work on diverse symbiotic networks since there are a large variety of symbioses with major roles in the proper functioning of terrestrial or aquatic ecosystems and we wished the Topic to provide a venue for communicating findings across diverse taxonomic groups A synthesis of recent investigations in symbiosis can impact areas such as agriculture where a basic understanding of plant microbe symbiosis will provide foundational information on the increasingly important issue of nitrogen fixation climate change where anthropogenic factors are threatening the survival of marine symbiotic ecosystems such as coral reefs animal and human health where unbalances in host microbiomes are being increasingly associated with a wide range of diseases and biotechnology where process optimization can be achieved through optimization of symbiotic partnerships Overall our vision was to produce a volume of works that will help define general principles of symbiosis within a new conceptual framework in the road to finally establish symbiology as an overdue central discipline of biological science *Protein Phosphorylation in Health and Disease* Allegra Via,Andreas Zanzoni,2016-08-11 Protein phosphorylation is one of the most abundant reversible post translational modifications in eukaryotes It is involved in virtually all cellular processes by regulating protein function localization and stability and by mediating protein protein interactions Furthermore aberrant protein phosphorylation is implicated in the

onset and progression of human diseases such as cancer and neurodegenerative disorders In the last years tens of thousands of in vivo phosphorylation events have been identified by large scale quantitative phospho proteomics experiment suggesting that a large fraction of the proteome might be regulated by phosphorylation This data explosion is increasingly enabling the development of computational approaches often combined with experimental validation aiming at prioritizing phosphosites and assessing their functional relevance Some computational approaches also address the inference of specificity determinants of protein kinases phosphatases and the identification of phosphoresidue recognition domains In this context several challenging issues are still open regarding phosphorylation including a better understanding of the interplay between phosphorylation and allosteric regulation agents and mechanisms disrupting or promoting abnormal phosphorylation in diseases the identification and modulation of novel phosphorylation inhibitors and so forth Furthermore the determinants of kinase and phosphatase recognition and binding specificity are still unknown in several cases as well as the impact of disease mutations on phosphorylation mediated signaling The articles included in this Research Topic illustrate the very diverse aspects of phosphorylation ranging from structural changes induced by phosphorylation to the peculiarities of phosphosite evolution Some also provide a glimpse into the huge complexity of phosphorylation networks and pathways in health and disease and underscore that a deeper knowledge of such processes is essential to identify disease biomarkers on one hand and design more effective therapeutic strategies on the other

[Sperm Differentiation and Spermatozoa Function:](#)

[Mechanisms, Diagnostics, and Treatment](#) Tomer Avidor-Reiss,Zhibing Zhang,Xin Zhiguo Li,2020-05-22

Push Button

Agriculture K. R. Krishna,2017-03-16 This book covers three main types of agricultural systems the use of robotics drones unmanned aerial vehicles and satellite guided precision farming methods Some of these are well refined and are currently in use while others are in need of refinement and are yet to become popular The book provides a valuable source of information on this developing field for those involved with agriculture and farming and agricultural engineering The book is also applicable as a textbook for students and a reference for faculty

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