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and 12 • III Acknowledgments This document has been updated from the 1996 IRP to include suggested achievement indicators, a more clear and succinct set of prescribed learning outcomes, a snapshot of the course's key elements, and

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IRP senior investigator Julie Segre, Ph.D., was one of four IRP researchers elected to the NAM in 2019. Dr. Dr. Segre leads the Translational and Functional Genomics Branch at the National Human Genome Research Institute (NHGRI), where she studies the way in which the skin forms a barrier between the body and the environment.

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This book provides a collection of applicable learning theories and their applications to science teaching. It presents a synthesis of historical theories while also providing practical implications for improvement of pedagogical practices aimed at advancing the field into the future. The theoretical viewpoints included in this volume span cognitive and social human development, address theories of learning, and describe approaches to teaching and curriculum development. The book presents and discusses

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humanistic, behaviourist, cognitivist, and constructivist theories. In addition, it looks at other theories, such as multiple intelligences theory, systems thinking, gender/sexuality theory and indigenous knowledge systems. Each chapter follows a reader-motivated approach anchored on a narrative genre. The book serves as a guide for those aiming to create optional learning experiences to prepare the next generation STEM workforce. Chapter "The Bildung Theory—From von Humboldt to Klafki and Beyond" is available open access under a Creative Commons Attribution 4.0

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book is addressed not only to researchers in mathematical programming, but to all scientists in various disciplines who use optimization methods in solving problems in computational chemistry and biology.

This book offers a meso-level description of demographics, science education, and science teacher education. Representing all 13 Canadian jurisdictions, the book provides local insights that serve as the basis for exploring the Canadian system as a whole and function as a common starting point from which to identify causal relationships that

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may be associated with Canada's successes. The book highlights commonalities, consistencies, and distinctions across the provinces and territories in a thematic analysis of the 13 jurisdiction-specific chapters. Although the analysis indicates a network of policy and practice issues warranting further consideration, the diverse nature of Canadian science education makes simple identification of causal relationships elusive. Canada has a reputation for strong science achievement. However, there is currently limited literature on science education in Canada at the general level or

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in specific areas such as Canadian science curriculum or science teacher education. This book fills that gap by presenting a thorough description of science education at the provincial/territorial level, as well as a more holistic description of pressing issues for Canadian science education.

The blood system is multi-scale, from the organism to the organs to cells to intracellular signaling pathways to macromolecule interactions. Blood consists of

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circulating cells, cellular fragments (platelets and microparticles), and plasma macromolecules. Blood cells and their fragments result from a highly-ordered process, hematopoiesis. Definitive hematopoiesis occurs in the bone marrow, where pluripotential stem cells give rise to multiple lineages of highly specialized cells. Highly-productive and continuously regenerative, hematopoiesis requires a microenvironment of mesenchymal cells and blood vessels. A Systems Biology Approach to Blood is divided into three main sections: basic components, physiological processes,

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and clinical applications. Using blood as a window, one can study health and disease through this unique tool box with reactive biological fluids that mirrors the prevailing hemodynamics of the vessel walls and the various blood cell types. Many blood diseases, rare and common can and have been exploited using systems biology approaches with successful results and therefore ideal models for systems medicine. More importantly, hematopoiesis offers one of the best studied systems with insight into stem cell biology, cellular interaction, development; lineage programming and

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reprogramming that are every day influenced by the most mature and understood regulatory networks.

Aluminum, bound almost exclusively to oxygen in various combinations, is the most abundant metal in the earth's crust and, therefore, of great commercial potential. Once methods were developed (in the 1880's) to free useable quantities of the element from oxygen, applications for the element began developing rapidly. This growth has resulted in the ubiquity of the metal in today's world. Therefore it can be found intentionally

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introduced in many products in direct contact with human beings. It is commonly known that soluble forms of aluminum are toxic to living organisms. However, aluminum is not known to be bioavailable under everyday conditions. In fact, the solubility product of common aluminum compounds, such as  $Al(OH)_3$  is so low as to make it essentially unavailable. This volume of Structure and Bonding seeks to provide in one source, a resource where the basic science related to aluminum toxicity may be obtained. It should be stressed that this volume is not intended to be a warning to avoid contact with aluminum. Living

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organisms have adequate defenses to prevent aluminum toxicity under normal conditions. Rather the volume was created to simply provide an understanding of the biological effects of aluminum. As such, the present volume should be considered in the context of the companion volumes in this three part series of Structure and Bonding. The first volume was devoted to fundamental developments in group 13 chemistry.

Metal ions in biology is an ever expanding area in science and medicine involving metal ions in proteins and enzymes, their



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biosynthesis, catalysis, electron transfer, metal ion trafficking, gene regulation and disease. While X-ray crystallography has provided snapshots of the geometric structures of the active site redox cofactors in these proteins, the application of high resolution EPR spectroscopy in conjunction with quantum chemistry calculations has enabled, in many cases, a detailed understanding of a metalloenzymes mechanism through investigations of the geometric and electronic structure of the resting, enzyme-substrate intermediates and product complexes. This volume, Part II of a two-

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volume set demonstrates the application of high resolution EPR spectroscopy in determining the geometric and electronic structure of active site metal ion centers in iron sulfur cluster containing metalloproteins, mononuclear molybdenum metalloenzymes, manganese-containing enzymes and novel metalloproteins.

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