

## Unit Y108 The Early Srts And The Origins Of The Civil War

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Even before it was a city, Vancouver was a property speculator's wet dream."There are more speculators about New Westminster and Victoria than there were in Winnipeg during the boom," CPR Chief WC Van Horne warned a friend in 1884, "and they are a much sharper lot. Nearly every person is more or less interested and you will have to be on your guard against all of them."Ever since Europeans first laid claim to the Squamish Nation territory in the 1870s, the real estate industry has held the region in its grip. Its influence has been grotesquely pervasive at every level of civic life, determining landmarks like Stanley Park and City Hall, as well as street names, neighbourhoods, even the name "Vancouver" itself. Land of Destiny aims to explore that influence, starting in 1862, with the first sale of land in the West End, and continuing up until the housing crisis of today. It will explore the backroom dealings, the skulduggery and nepotism, the racism and the obscene profits, while at the same time revealing that the same forces which made Vancouver what it is, speculation and global capital, are the same ones that shape it today, showing that more than anything else, the history of real estate and the history of Vancouver are one and the same.And it's been dirty as hell.About the Series: Land of Destiny is the first title in Anvil's new series "49.2: Tales from the Off Beat," an ongoing series dedicated to celebrating the eccentric and unusual parts of city history. From Jesse Donaldson, author of the Bill Duthie Booksellers' Choice Award finalist book This Day In Vancouver, and a host of other local historians, the series will be an in-depth examination of the weird, the wonderful, and the terrible, injecting fresh details into well-worn local lore, or digging deep into the obscure people, places, and happenings of the last 130 years. From psychedelic hospitals to town fools, from communist organizers to real estate scumbags, 49.2 will take pains to break down the myths surrounding the City of Glass.

A classic problem in mathematics is solving systems of polynomial equations in several unknowns. Today, polynomial models are ubiquitous and widely used across the sciences. They arise in robotics, coding theory, optimization, mathematical biology, computer vision, game theory, statistics, and numerous other areas. This book furnishes a bridge across mathematical disciplines and exposes many facets of systems of polynomial equations. It covers a wide spectrum of mathematical techniques and algorithms, both symbolic and

numerical. The set of solutions to a system of polynomial equations is an algebraic variety - the basic object of algebraic geometry. The algorithmic study of algebraic varieties is the central theme of computational algebraic geometry. Exciting recent developments in computer software for geometric calculations have revolutionized the field. Formerly inaccessible problems are now tractable, providing fertile ground for experimentation and conjecture. The first half of the book gives a snapshot of the state of the art of the topic. Familiar themes are covered in the first five chapters, including polynomials in one variable, Grobner bases of zero-dimensional ideals, Newton polytopes and Bernstein's Theorem, multidimensional resultants, and primary decomposition. The second half of the book explores polynomial equations from a variety of novel and unexpected angles. It introduces interdisciplinary connections, discusses highlights of current research, and outlines possible future algorithms. Topics include computation of Nash equilibria in game theory, semidefinite programming and the real Nullstellensatz, the algebraic geometry of statistical models, the piecewise-linear geometry of valuations and amoebas, and the Ehrenpreis-Palamodov theorem on linear partial differential equations with constant coefficients. Throughout the text, there are many hands-on examples and exercises, including short but complete sessions in MapleR, MATLABR, Macaulay 2, Singular, PHCpack, CoCoA, and SOSTools software. These examples will be particularly useful for readers with no background in algebraic geometry or commutative algebra. Within minutes, readers can learn how to type in polynomial equations and actually see some meaningful results on their computer screens. Prerequisites include basic abstract and computational algebra. The book is designed as a text for a graduate course in computational algebra.

In its 114th year, Billboard remains the world's premier weekly music publication and a diverse digital, events, brand, content and data licensing platform. Billboard publishes the most trusted charts and offers unrivaled reporting about the latest music, video, gaming, media, digital and mobile entertainment issues and trends.

A comprehensive guide to full-time degree courses, institutions and towns in Britain.

With the most comprehensive and up-to-date overview of structure-based drug discovery covering both experimental and computational approaches, Structural Biology in Drug Discovery: Methods, Techniques, and Practices describes principles, methods, applications, and emerging paradigms of structural biology as a tool for more efficient drug development. Coverage includes successful examples, academic and industry insights, novel concepts, and advances in a rapidly evolving field. The combined chapters, by authors writing from the frontlines of structural biology and drug discovery, give readers a valuable reference and resource that: Presents the benefits, limitations, and potentiality of major techniques in the field such as X-ray crystallography, NMR, neutron crystallography, cryo-EM, mass spectrometry and other biophysical techniques, and computational structural biology. Includes detailed chapters on druggability, allostery, complementary use of thermodynamic and kinetic information, and powerful approaches such as structural chemogenomics and fragment-based drug design. Emphasizes the need for the in-depth biophysical characterization of protein targets as well as of therapeutic proteins, and for a thorough quality assessment of experimental structures. Illustrates advances in the field of established therapeutic targets like kinases, serine proteinases, GPCRs, and epigenetic proteins, and of more challenging ones like protein-protein interactions and intrinsically disordered proteins.

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