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Titolo	Failed imagination? : the Anglo-American New World Order from Wilson to Bush // Andrew Williams
Pubbl/distr/stampa	Manchester, [England] ; ; New York, New York : , : Manchester University Press, , 2007 [Place of distribution not identified] : , : Palgrave, , [date of distribution not identified] ©2007
ISBN	0-7190-9513-1
Edizione	[Second edition.]
Descrizione fisica	1 online resource (358 pages)
Disciplina	909.82
Soggetti	World politics - 20th century International relations - History - 20th century International relations
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	Front matter -- Dedication -- Contents -- Preface to the second edition -- Acknowledgements -- List of abbreviations -- Introduction -- 1 The imagining of the Versailles NWO, 1914-19 -- 2 The Paris Peace Conference and the Treaty of Versailles, 1919 -- 3 The United States and the planning of an American NWO, 1939-44 -- 4 The 'first follower' and Roosevelt's NOW -- 5 Joint Allied proposals for a NOW -- 6 International organisation, global security and the NOW -- 7 The economic element of the NWO project -- 8 Self-determination and the NOW -- Conclusion -- Select bibliography -- Index
Sommario/riassunto	The main purpose of this book is to explain how (mainly) American, but also British and other Western, policy makers have planned and largely managed to create an international order in their own image, the so-called 'New World Order'. It shows how this seismic shift in international relations has developed through the major global wars of the twentieth and twenty-first centuries. It uses a wide variety of historical archival material to give the background to the current and historical American obsession with creating the world order, one that both reflects the American national interest but also can be said to

have established the major security, economic, organisational and normative pillars of our epoch. In addition it provides excellent background reading for the current debate about American foreign policy and the origins of 'neo-conservatism' in international relations. This edition updates a very successful first edition of the title, with additional material to take into account changes in the global order since 2001 and the beginning of the 'War on Terror'.

2. Record Nr.	UNINA9911019352003321
Titolo	Analysing gene expression : a handbook of methods: possibilities and pitfalls // edited by Stefan Lorkowski and Paul Cullen
Pubbl/distr/stampa	Weinheim ; ; New York, : Wiley-VCH, c2003
ISBN	9786610520428 9781280520426 1280520426 9783527605330 3527605339 9783527601493 352760149X
Descrizione fisica	1 online resource (984 p.)
Altri autori (Persone)	LorkowskiStefan CullenPaul, Dr.
Disciplina	572.8/65
Soggetti	Gene expression
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Livello bibliografico	Monografia
Note generali	Description based upon print version of record.
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	Analysing Gene Expression A Handbook of Methods Possibilities and Pitfalls; Foreword; Preface; Contents; List of Contributors; Volume 1; 1 Basic concepts of gene expression; 1.1 Introduction; 1.2 Basics of transcription and translation in the cell; 1.2.1 Introduction; 1.2.2 Transcription; 1.2.3 Translation; 1.2.4 Summary; 1.3 Regulation of transcription; 1.3.1 Introduction; 1.3.2 mRNA expression profiles - the transcriptome; 1.3.3 Protein expression profiles - the proteome; 1.3.4

Interaction between genes and proteins - the interactome; 1.3.5 The transcription machinery and core promoters
1.3.6 Regulatory promoters 1.3.7 Enhancers; 1.3.8 Locus control regions; 1.3.9 Matrix attachment regions; 1.3.10 Insulators; 1.3.11 RIDGEs - Regions of increased gene expression; 1.3.12 Enhanceosomes; 1.3.13 Chromatin; 1.3.14 Silencer elements; 1.3.15 Transcription factors, repressors and co-repressors; 1.3.16 Epigenetics; 1.3.17 Summary and conclusions; 1.4 Post-transcriptional regulation; 1.4.1 Introduction; 1.4.2 Regulation of RNA stability and degradation; 1.4.3 Regulation of transcription elongation; 1.4.3.1 Introduction; 1.4.3.2 Regulation of transcription elongation in prokaryotes
1.4.3.3 Regulation of transcription elongation in eukaryotes 1.4.3.4 Conclusions; 1.4.4 Differential/alternative pre-mRNA splicing; 1.4.5 Trans-RNA splicing; 1.4.6 Regulation of mRNA transport; 1.4.7 Directed intracellular mRNA localisation; 1.4.8 Regulation of mRNA polyadenylation; 1.4.9 Antisense RNA; 1.4.10 RNA editing; 1.4.11 Summary and conclusions; 1.5 Post-translational modification of proteins; 1.5.1 Introduction; 1.5.2 Proteolytic cleavage of proteins; 1.5.3 Acylation; 1.5.4 Prenylation; 1.5.5 Methylation; 1.5.6 Sulphation; 1.5.7 Phosphorylation; 1.5.8 Ubiquitination
1.5.9 Glycosylation 1.5.10 Conclusions; 1.6 Correlation of mRNA and protein expression; 1.6.1 Introduction; 1.6.2 Levels of mRNA and protein expression: correlations and discrepancies; 1.6.3 Conclusions; 1.7 Housekeeping genes, internal and external standards; 1.7.1 What are housekeeping genes?; 1.7.2 Survey of the most important housekeeping genes; 1.7.2.1 Glyceraldehyde-3-phosphate dehydrogenase; 1.7.2.2 -Actin; 1.7.3 Other commonly used housekeeping genes; 1.7.3.1 Ribosomal RNA (rRNA); 1.7.4 New identified 'maintenance genes'; 1.7.5 Methods of quantification 1.7.5.1 Normalisation (via internal standards) 1.7.5.2 Standard curves (via external standards); 1.7.5.3 Alternative methods (via internal standards); 1.7.6 Summary; 1.8 Classification of differential gene expression technologies; 1.8.1 Introduction; 1.8.2 From single genes to transcriptomes; 1.8.3 Methods of classification; 1.8.4 Conclusions; 1.9 Summary; 1.10 References; 2 Sample preparation and supplementary tools; 2.1 Introduction; 2.2 Preparation of cells and tissues; 2.2.1 Immunopurification of cells; 2.2.2 Differential-velocity centrifugation/countercurrent elutriation
2.2.3 Surface affinity chromatography

Sommario/riassunto

This book combines the experience of 225 experts on 900 pages. Scientists worldwide are currently overwhelmed by the ever-increasing number and diversity of genome projects. This handbook is your guide through the jungle of new methods and techniques available to analyse gene expression - the first to provide such a broad view of the measurement of mRNA and protein expression in vitro, in situ and even in vivo. Despite this broad approach, detail is sufficient for you to grasp the principles behind each method. In each case, the authors weigh up the advantages and disadvantages, paying parti
