

1. Record Nr.	UNINA9910455242303321
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Titolo	Christian mysticism [[electronic resource]] : an introduction to contemporary theoretical approaches // Louise Nelstrop, Kevin Magill, and Bradley B. Onishi
Pubbl/distr/stampa	Farnham, Surrey, England ; ; Burlington, VT, : Ashgate, 2009
ISBN	1-317-16669-8 1-315-57188-9 1-282-34470-6 9786612344701 0-7546-8695-7
Descrizione fisica	1 online resource (288 p.)
Altri autori (Persone)	MagillKevin J. <1972-> OnishiBradley B
Disciplina	248.2/2
Soggetti	Mysticism Philosophy and religion Electronic books.
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Description based upon print version of record.
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	Cover; Contents; Preface and Acknowledgements; Introduction: Contemporary Theoretical Approaches; Part I Key Themes and Motifs; 1 Platonism and Christian Mysticism; 2 Negative or Apophatic Language; 3 Selfhood and Interiority; 4 Erotic Imagery; 5 Hierarchy; 6 Symbolism and Scripture's 'Mystical Sense'; Part II Later Developments in Christian Mysticism; 7 Female Mysticism; 8 Visionary Mysticism; 9 Imaginative Mysticism, Meditation and Experience; 10 Mysticism and Heresy; III Postmodern Re-Readings of Pseudo-Denys and Augustine 11 Exploring Negative Language through Pseudo-Denys: Derrida and Marion12 Exploring Selfhood through Augustine: Marion, Derrida and Lyotard; Conclusion: What is Christian Mysticism?; Bibliography; Index
Sommario/riassunto	This book introduces students to Christian mysticism and modern critical responses to it. Christianity has a rich tradition of mystical theology that first emerged in the writings of the early church fathers, and flourished during the Middle Ages. Today Christian mysticism is

increasingly recognised as an important Christian heritage relevant to today's spiritual seekers. The book sets out to provide students and other interested readers with access to the main theoretical approaches to Christian mysticism - including those propounded by William James, Steven Katz, Bernard McGinn, Michael Sells

2. Record Nr.	UNINA9910831192003321
Titolo	The epigenome [[electronic resource]] : molecular hide and seek // edited by S. Beck and A. Olek
Pubbl/distr/stampa	Weinheim ; ; [Cambridge], : Wiley-VCH, c2003
ISBN	1-280-52085-X 9786610520855 3-527-60597-5 3-527-60151-1
Descrizione fisica	1 online resource (190 p.)
Altri autori (Persone)	OlekA (Alexander) BeckStephan (Stephan G.)
Disciplina	599.935 611.01816
Soggetti	Human genome Medical genetics Nature and nurture Genome, Human Genomics - methods Cytosine - physiology
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Description based upon print version of record.
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	The Epigenome Molecular Hide and Seek; Preface; Contents; 1 Five Not Four: History and Significance of the Fifth Base; Summary; 1.1 Historical Introduction; 1.2 Sequencing 5-methylcytosine (5-mC) Residues in Genomic DNA; The Bisulfite Method; 1.3 Gene Silencing; 1.4 Development; 1.5 Abnormal DNA Methylation in Cancer Cells; 1.6

Nuclear Transfer; 1.7 Aging; 1.8 The Future; References; 2 (Epi)genetic Signals: Towards a Human Genome Sequence of All Five Nucleotides; Summary; 2.1 A Linguistic Prologue; 2.2 Towards the Complete Sequence of the Human Genome with Five Nucleotides
2.3 Patterns of DNA Methylation - the Scaffold for Building a Functional Genome
2.4 DNA Methylation Patterns in Segments of the Human Genome and in Viral Genomes; 2.4.1 On Viral Genomes and Foreign DNA Integrates (Table 2.1); 2.4.2 DNA Methylation Patterns in the Human Genome (Table 2.1); 2.5 Insertions of Foreign DNA into Established Mammalian Genomes; 2.6 De Novo Methylation of Integrated Foreign DNA; 2.6.1 Ad12 Genomes in Hamster Tumor Cells; 2.6.2 De Novo Methylation of Foreign DNA Integrated into the Mouse Genome by Homologous or Heterologous Recombination [23]
2.7 Genome-wide Perturbations in the Mammalian Genome upon Foreign DNA Insertion
2.8 Outlook and Recommendations; References;
3 Epi Meets Genomics: Technologies for Finding and Reading the 5(th) Base; Summary; 3.1 The Development of 5(th)-Base Technologies; 3.1.1 Unusual DNA-cutting Enzymes; 3.1.2 A Unique Chemical Reaction that Modifies Methylated DNA; 3.1.3 The Advance of 5(th) Base Technologies in Epigenomic Research; 3.2 Restriction Landmark Genomic Scanning (RLGS): Finding the 5(th)-base Signposts in the Genomic Atlas; 3.2.1 Principle; 3.2.2 How Does RLGS Work?; 3.2.3 Applications
3.3 Methylation-sensitive Arbitrarily Primed (AP) PCR: Fishing for the 5 (th) Bases in Genomic Ponds
3.3.1 Principle; 3.3.2 How Does MS AP-PCR Work?; 3.3.3 Applications; 3.4 Differential Methylation Hybridization (DMH): Identifying the 5(th) Bases in the Genomic Crossword Puzzle; 3.4.1 Principle; 3.4.2 How Does DMH Work?; 3.4.3 Applications; 3.5 MethyLight: Finding 5(th)-base Patterns in Genomic Shadows; 3.5.1 Principle; 3.5.2 How Does MethyLight Work?; 3.5.3 Applications; 3.6 Exploring the Epigenome; References;
4 Mammalian Epigenomics: Reprogramming the Genome for Development and Therapy
Summary
4.1 Introduction; 4.2 DNA Methylation; 4.3 Histone Modifications; 4.4 Imprinting; 4.5 Reprogramming and Cloning; 4.6 Epimutations and Epigenetic Inheritance; 4.7 Epigenomics: The Future; 4.7 Conclusions; References;
5 At the Controls: Genomic Imprinting and the Epigenetic Regulation of Gene Expression; Summary; 5.1 Introduction; 5.2 Genomic Imprinting; 5.2.1 The Role of DNA Methylation in Imprinted Gene Expression; 5.2.3 Organization of Imprinted Genes
5.2.4 The Mechanism of Imprinting at the Mouse Igf2r Imprinted Domain Requires a Cis-acting Noncoding Antisense Transcript Regulated by DNA Methylation (Fig. 5.2 a)

Sommario/riassunto

This is the first book that describes the role of the Epigenome (cytosine methylation) in the interplay between nature and nurture. It focuses and stimulates interest in what will be one of the most exciting areas of post-sequencing genome science: the relationship between genetics and the environment. Written by the most reputable authors in the field, this book is essential reading for researchers interested in the science arising from the human genome sequence and its implications on health care, industry and society.
