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Nota di contenuto	 Preface; Glossary; Contents; The Oocyte Determinants of Early Reprogramming; 1 Controversies Surrounding Oocyte Reprogramming; 2 A Brief History of SCNT; 3 The Uniqueness of the Oocyte: More than Totipotency; 4 Qualitative Aspects of Oocyte-Mediated Nuclear Reprogramming; 4.1 Characteristic Traits of the Oocyte Are Species Specific; 4.2 Transcriptional Activity in Germinal Vesicle-Stage Oocytes; 4.3 Different Maturation States and Accompanying Chromatin Configuration of Recipient Oocytes for SCNT; 5 Quantitative Aspects of Oocyte-Mediated Nuclear Reprogramming 5.1 Amount of Reprogramming Factors5.2 Kinetics of Reprogramming; 6 Gene Expression in Oocytes; 7 The Elusive Reprogramming Factors; 8 Using Transcriptomics and Proteomics to Search for the Molecular Fingerprint Indicative of an Oocyte's Reprogramming Potenti; 8.1 Candidate Gene Approach; 8.2 Transcriptome Analysis; 8.3 Proteome

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	Analysis; Conclusion; References; Stella and Zygotic Reprogramming; 1 Stella; 1.1 Identification of Stella; 1.2 Expression Pattern of Stella; 1.3 Gene Disruption Analysis of Stella; 2 Epigenetic Reprogramming in Zygotes 2.1 Active Loss of 5mC in Zygotic Paternal Genome2.2 Protection of Imprinted Genes and Repetitive Sequences in Zygotes; 2.3 Involvement of 5hmC and Tet Proteins in Active Loss of 5mC; 2.4 Histone Modification in Zygotes; 3 Mechanism of Stella-Mediated Protection of 5mC in Zygotes; 3.1 Stella Protects Active Loss of 5mC in Zygotes; 3.2 Stella Protects Imprinted Genes and Repetitive Sequences; 3.3 Protective Function of Stella Depends on H3K9me2; 3.4 Mechanism of Stella-Mediated Protection of 5mC in Zygotes; 4 Perspectives; References; Histone Variants and Reprogramming in Early Development 1 Mammalian Development: Context and Early Epigenetic Reprogramming2 Mechanisms of Epigenetic Reprogramming and Chromatin Remodelling in the Early Embryo; 3 The Components of the Chromatin Change as Development Proceeds; 4 Histone Variants as Regulators of Epigenetic Information During Reprogramming; 5 H3.3 and De Novo Establishment of Heterochromatin; 6 H3 Variants: A Conserved Function in the Germline; 7 Variants of H2A: The Case of MacroH2A; 8 H2A.Z Shows a Dynamic Localisation During Early Reprogramming in Embryos 9 High Endogenous Levels of Phosphorylated H2A.X Are Characteristic of Early Embryos10 Barr Body-Deficient H2A: H2A.B; References; DNA Methylation Reprogramming in Preimplantation Development; 1 Epigenetic Profiles in Mature Gametes; 2 DNA Methylation Reprogramming in the Zygote; 3 The Mechanisms of DNA Demethylation; 4 DNA Methylation Reprogramming in Further Preimplantation Development; 5 Roles of Domts in Reprutation Davelopment; 6 Prose
	Preimplantation Development; 5 Roles of Dnmts in Regulating DNA Methylation in Early Embryos; 5.1 Dnmt1; 5.2 Dnmt3a, Dnmt3L and Dnmt3b; 6 DNA Methylation Reprogramming in SCNT-Derived Embryos 7 Histone Modifications and DNA Methylation Crosstalk
Sommario/riassunto	The ability of a single genome to give rise to hundreds of functionally distinct cell type programs is in itself remarkable. Pioneering studies over the past few decades have demonstrated that this plasticity is retained throughout development, a phenomenon of epigenetic programming and reprogramming that remains one of the most fascinating areas of modern biology, with major relevance to human health and disease. This book presents the basic biology involved, including key mechanistic insights into this rapidly growing field.