Record Nr. UNINA9910969611203321 Autore Cristini Vittorio <1970-> Titolo Multiscale modeling of cancer: an integrated experimental and mathematical modeling approach / / Vittorio Cristini, John Lowengrub Cambridge:,: Cambridge University Press,, 2010 Pubbl/distr/stampa 1-107-21140-9 **ISBN** 1-282-77073-X 9786612770739 0-511-90148-8 0-511-79912-8 0-511-90227-1 0-511-79772-9 0-511-78145-8 0-511-90069-4 Edizione [1st ed.] Descrizione fisica 1 online resource (xix, 278 pages): digital, PDF file(s) Disciplina 362.196/994 Soggetti Cancer - Research - Mathematical models Multiscale modeling Lingua di pubblicazione Inglese **Formato** Materiale a stampa Livello bibliografico Monografia Title from publisher's bibliographic system (viewed on 05 Oct 2015). Note generali Nota di bibliografia Includes bibliographical references and index. Part I. Theory: 1. Introduction; 2. Biological background; 3. Continuum Nota di contenuto tumor modeling: single phase; 4. Analysis and calibration of singlephase continuum tumor models; 5. Continuum tumor modeling: multiphase; 6. Discrete cell modeling; 7. Hybrid continuum -- discrete models; 8. Numerical schemes -- Part II. Applications: 9. Continuum tumor modeling: a multidisciplinary approach; 10. Agent-based cell modeling: application to breast cancer. Sommario/riassunto Mathematical modeling, analysis and simulation are set to play crucial roles in explaining tumor behavior, and the uncontrolled growth of cancer cells over multiple time and spatial scales. This book, the first to

integrate state-of-the-art numerical techniques with experimental data, provides an in-depth assessment of tumor cell modeling at multiple scales. The first part of the text presents a detailed biological

background with an examination of single-phase and multi-phase continuum tumor modeling, discrete cell modeling, and hybrid continuum-discrete modeling. In the final two chapters, the authors guide the reader through problem-based illustrations and case studies of brain and breast cancer, to demonstrate the future potential of modeling in cancer research. This book has wide interdisciplinary appeal and is a valuable resource for mathematical biologists, biomedical engineers and clinical cancer research communities wishing to understand this emerging field.