1. Record Nr. UNINA990003834430403321

Autore	Deaglio, Mario <1943- >
Titolo	La fine dell'euforia / Mario Deaglio
Pubbl/distr/stampa	Milano : Guerrini e associati, c2001
ISBN	88-8335-213-0
Descrizione fisica	XII, 201 p. ; 23 cm
Disciplina	337.09049
Locazione	SE SDECBC DECSE
Collocazione	F/1.411 DEA/ D/8.15 SEB F/1.411 DEA/6 DEA337.09049B DEA337.09049C DEA337.09049D DEA337.09049A DEA337.09049E DEA337.09049H DEA337.09049F DEA337.09049G SE116.04.04-
Lingua di pubblicazione	Italiano
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Nell'occh.: Sesto rapporto sull'economia globale e l'Italia

2. Record Nr. UNINA9910135859403321

Titolo	IEEE Std 488.2-1992 : IEEE Standard Codes, Formats, Protocols, and Common Commands for Use With IEEE Std 488.1-1987, IEEE Standard Digital Interface for Programmable Instrumentation / / Institute of Electrical and Electronics
Pubbl/distr/stampa	Piscataway, NJ, USA : , : IEEE, , 1991
ISBN	0-7381-0665-8
Descrizione fisica	1 online resource (254 pages)
Disciplina	629.8
Soggetti	Digital control systems
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Sommario/riassunto	A set of codes and formats to be used by devices connected via the IEEE 488.1 bus is specified. This standard also defines communication protocols that are necessary to effect application-independent and device-dependent message exchanges, and furthere defines common commands and characteristics useful in instrument system applications. It is intended to apply to small-scale to medium-scale instrument systems comprised mainly of measurement, stimulus, and interconnect devices with an instrumentation controller. The standard may also apply to certain devices outside the scope of the instrument system environment. IEEE 488.1 subsets, standard message-handling protocols including error handling, unambiguous program and response-message syntactic structures, common commands useful in a wide range of instrument system applications, standard status reporting structures, and system configuration and synchronization protocols are covered. IEEE Std 488.2-1992 , 488.2.

3. Record Nr. UNINA9910416107703321

Titolo	Next Generation Kinase Inhibitors : Moving Beyond the ATP Binding/Catalytic Sites / / edited by Paul Shapiro
Pubbl/distr/stampa	Cham : , : Springer International Publishing : , : Imprint : Springer, , 2020
ISBN	3-030-48283-9
Edizione	[1st ed. 2020.]
Descrizione fisica	1 online resource (XII, 217 p. 59 illus., 56 illus. in color.)
Disciplina	572.792
Soggetti	Cancer - Research Cancer Research Proteïnes quinases Llibres electrònics
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di contenuto	Chapter 1: Introduction to Kinases, Cellular Signaling, and Kinase Inhibitors -- Chapter 2: Overview of Current Type I/II Kinase Inhibitors -- Chapter 3: Avoiding or Co-opting ATP Inhibition: Type III, IV, V, and VI Kinase Inhibitors -- Chapter 4: Structural Features Regulating Kinase Interactions with Regulatory and Substrate Proteins -- Chapter 5: Developing Kinase Inhibitors using Computer-Aided Drug Design Approaches -- Chapter 6: A Toolbox of Structural Biology and Enzyme Kinetics Reveals the Case for ERK Docking Site Inhibition -- Chapter 7: Novel Stabilized Peptide Inhibitors of Protein Kinases -- Chapter 8: Novel peptide-based inhibitors of protein kinases -- Index.
Sommario/riassunto	Protein kinases are fascinating enzymes that maintain the proper function of nearly every task performed by the cells of the human body. By extracting a phosphate from the energy molecule ATP and linking it to another protein, protein kinases alter the structure and ultimate function of other proteins. In this way, protein kinases help monitor the extracellular environment and integrate signaling cues that, for the most part, are beneficial for human health and survival. However, protein kinases are often dysregulated and responsible for the initiation and progression of many types of cancers, inflammatory disorders, and other diseases. Thus, decades of research have revealed much about

how protein kinases are regulated and approaches to inhibit these enzymes to treat disease. However, nearly 30 years since the identification of the first clinically beneficial small molecule protein kinase inhibitor, there are only a few examples where these drugs provide sustained and durable patient responses. The goal of this book is to provide biomedical scientists, graduate, and professional degree students insight into different approaches using small molecules to block specific protein kinase functions that promote disease. .