Record Nr.	UNINA9910137167803321
Titolo	Fragment-based drug discovery : lessons and outlook / / edited by Daniel A. Erlanson and Wolfgang Jahnke
Pubbl/distr/stampa	Wiesbaden, Germany : , : Wiley-VCH Verlag GmbH & Co. KGaA, , 2016 ©2016
ISBN	3-527-68362-3 3-527-68360-7 3-527-68361-5
Descrizione fisica	1 online resource (527 p.)
Collana	Methods and Principles in Medicinal Chemistry
Disciplina	615.19
Soggetti	Drug development Drugs - Design Ligands (Biochemistry) Drug Discovery Llgands
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 at the end of each chapters and index.
Nota di contenuto	Fragment-based Drug Discovery: Lessons and Outlook; Contents; Contributors; Preface; A Personal Foreword; Part I: The Concept of Fragment-based Drug Discovery; 1. The Role of Fragment-based Discovery in Lead Finding; 1.1 Introduction; 1.2 What is FBLD?; 1.3 FBLD: Current Practice; 1.3.1 Using Fragments: Conventional Targets; 1.3.2 Using Fragments: Unconventional Targets; 1.4 What do Fragments Bring to Lead Discovery?; 1.5 How did We Get Here?; 1.5.1 Evolution of the Early Ideas and History; 1.5.2 What has Changed Since the First Book was Published in 2006? 1.6 Evolution of the Methods and Their Application Since 20051.6.1 Developments in Fragment Libraries; 1.6.2 Fragment Hit Rate and Druggability; 1.6.3 Developments in Fragment Screening; 1.6.4 Ways of Evolving Fragments; 1.6.5 Integrating Fragments Alongside Other Lead-Finding Strategies; 1.6.6 Fragments Can be Selective; 1.6.7 Fragment Binding Modes; 1.6.8 Fragments, Chemical Space, and

1.

Novelty; 1.7 Current Application and Impact; 1.8 Future Opportunities; References: 2. Selecting the Right Targets for Fragment-Based Drug **Discovery**; 2.1 Introduction 2.2 Properties of Targets and Binding Sites2.3 Assessing Druggability; 2.4 Properties of Ligands and Drugs; 2.5 Case Studies; 2.5.1 Case Study 1: Inhibitors of Apoptosis Proteins (IAPs); 2.5.2 Case Study 2: HCV-NS3; 2.5.3 Case Study 3: PKM2; 2.5.4 Case Study 4: Soluble Adenylate Cyclase; 2.6 Conclusions; References; 3. Enumeration of Chemical Fragment Space: 3.1 Introduction: 3.2 The Enumeration of Chemical Space; 3.2.1 Counting and Sampling Approaches; 3.2.2 Enumeration of the Chemical Universe Database GDB; 3.2.3 GDB Contents; 3.3 Using and Understanding GDB; 3.3.1 Drug Discovery 3.3.2 The MQN System3.3.3 Other Fingerprints; 3.4 Fragments from GDB; 3.4.1 Fragment Replacement; 3.4.2 Shape Diversity of GDB Fragments: 3.4.3 Aromatic Fragments from GDB: 3.5 Conclusions and Outlook; Acknowledgment; References; 4. Ligand Efficiency Metrics and their Use in Fragment Optimizations; 4.1 Introduction; 4.2 Ligand Efficiency; 4.3 Binding Thermodynamics and Efficiency Indices; 4.4 Enthalpic Efficiency Indices; 4.5 Lipophilic Efficiency Indices; 4.6 Application of Efficiency Indices in Fragment-Based Drug Discovery Programs: 4.7 Conclusions: References Part II: Methods and Approaches for Fragment-based Drug Discovery5. Strategies for Fragment Library Design; 5.1 Introduction; 5.2 Aims; 5.3 Progress; 5.3.1 BDDP Fragment Library Design: Maximizing Diversity; 5.3.2 Assessing Three-Dimensionality; 5.3.3 3DFrag Consortium; 5.3.4 Commercial Fragment Space Analysis; 5.3.5 BDDP Fragment Library Design: 5.3.6 Fragment Complexity: 5.3.6.1 Diversity-Oriented Synthesis-Derived Fragment-Like Molecules; 5.4 Future Plans; 5.5 Summary; 5.6 Key Achievements; References 6. The Synthesis of Biophysical Methods In Support of Robust Fragment-Based Lead Discovery