Giovanni Bottegoni

List of Publications by Year in descending order

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62 papers 3,756 citations

147566 31 h-index 60 g-index

68 all docs 68
docs citations

68 times ranked 8134 citing authors

#	Article	IF	CITATIONS
1	Role of Molecular Dynamics and Related Methods in Drug Discovery. Journal of Medicinal Chemistry, 2016, 59, 4035-4061.	2.9	797
2	Four-Dimensional Docking: A Fast and Accurate Account of Discrete Receptor Flexibility in Ligand Docking. Journal of Medicinal Chemistry, 2009, 52, 397-406.	2.9	172
3	Recipes for the Selection of Experimental Protein Conformations for Virtual Screening. Journal of Chemical Information and Modeling, 2010, 50, 186-193.	2.5	160
4	A catalytically silent FAAH-1 variant drives anandamide transport in neurons. Nature Neuroscience, 2012, 15, 64-69.	7.1	150
5	Combining Galantamine and Memantine in Multitargeted, New Chemical Entities Potentially Useful in Alzheimer's Disease. Journal of Medicinal Chemistry, 2012, 55, 9708-9721.	2.9	129
6	The role of fragment-based and computational methods in polypharmacology. Drug Discovery Today, 2012, 17, 23-34.	3.2	110
7	Structure of the complement C5a receptor bound to the extra-helical antagonist NDT9513727. Nature, 2018, 553, 111-114.	13.7	110
8	Non-ATP Competitive Protein Kinase Inhibitors. Current Medicinal Chemistry, 2010, 17, 2804-2821.	1.2	108
9	Multitarget Drug Discovery for Alzheimer's Disease: Triazinones as BACEâ€1 and GSKâ€3β Inhibitors. Angewandte Chemie - International Edition, 2015, 54, 1578-1582.	7.2	107
10	Consistent Improvement of Cross-Docking Results Using Binding Site Ensembles Generated with Elastic Network Normal Modes. Journal of Chemical Information and Modeling, 2009, 49, 716-725.	2.5	106
11	Dual inhibition of REV-ERB \hat{l}^2 and autophagy as a novel pharmacological approach to induce cytotoxicity in cancer cells. Oncogene, 2015, 34, 2597-2608.	2.6	100
12	Versatility of the Curcumin Scaffold: Discovery of Potent and Balanced Dual BACE-1 and GSK-3 \hat{l}^2 Inhibitors. Journal of Medicinal Chemistry, 2016, 59, 531-544.	2.9	100
13	The ligand binding mechanism to purine nucleoside phosphorylase elucidated via molecular dynamics and machine learning. Nature Communications, 2015, 6, 6155.	5.8	98
14	BACE-1 Inhibitors: From Recent Single-Target Molecules to Multitarget Compounds for Alzheimer's Disease. Journal of Medicinal Chemistry, 2018, 61, 619-637.	2.9	90
15	Systematic Exploitation of Multiple Receptor Conformations for Virtual Ligand Screening. PLoS ONE, 2011, 6, e18845.	1.1	82
16	Molecular Dynamics Simulations and Kinetic Measurements to Estimate and Predict Protein–Ligand Residence Times. Journal of Medicinal Chemistry, 2016, 59, 7167-7176.	2.9	81
17	A new method for ligand docking to flexible receptors by dual alanine scanning and refinement (SCARE). Journal of Computer-Aided Molecular Design, 2008, 22, 311-325.	1.3	74
18	Irreversible Protein Kinase Inhibitors. Current Medicinal Chemistry, 2011, 18, 2981-2994.	1.2	67

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19	A Computational Study of the Binding of Propidium to the Peripheral Anionic Site of Human Acetylcholinesterase. Journal of Medicinal Chemistry, 2004, 47, 3991-3999.	2.9	61
20	Multi-Kinase Inhibitors. Current Medicinal Chemistry, 2015, 22, 695-712.	1,2	61
21	Kinetic and Structural Insights into the Mechanism of Binding of Sulfonamides to Human Carbonic Anhydrase by Computational and Experimental Studies. Journal of Medicinal Chemistry, 2016, 59, 4245-4256.	2.9	60
22	3,4-Dihydro-1,3,5-triazin-2(1 <i>H</i>)-ones as the First Dual BACE-1/GSK-3β Fragment Hits against Alzheimer's Disease. ACS Chemical Neuroscience, 2015, 6, 1665-1682.	1.7	54
23	A Comparative Study on the Application of Hierarchicalâ^'Agglomerative Clustering Approaches to Organize Outputs of Reiterated Docking Runs. Journal of Chemical Information and Modeling, 2006, 46, 852-862.	2.5	52
24	Low molecular weight, non-peptidic agonists of TrkA receptor with NGF-mimetic activity. Cell Death and Disease, 2012, 3, e339-e339.	2.7	48
25	BiKi Life Sciences: A New Suite for Molecular Dynamics and Related Methods in Drug Discovery. Journal of Chemical Information and Modeling, 2018, 58, 219-224.	2.5	48
26	Diaryl Urea: A Privileged Structure in Anticancer Agents. Current Medicinal Chemistry, 2016, 23, 1528-1548.	1,2	47
27	Benzophenone-based derivatives: A novel series of potent and selective dual inhibitors of acetylcholinesterase and acetylcholinesterase-induced beta-amyloid aggregation. European Journal of Medicinal Chemistry, 2011, 46, 1682-1693.	2.6	43
28	Small Molecule Aurora Kinases Inhibitors. Current Medicinal Chemistry, 2009, 16, 1949-1963.	1,2	42
29	ACIAP, Autonomous hierarchical agglomerative Cluster Analysis based protocol to partition conformational datasets. Bioinformatics, 2006, 22, e58-e65.	1.8	41
30	Discovery of a New Class of Highly Potent Inhibitors of Acid Ceramidase: Synthesis and Structure–Activity Relationship (SAR). Journal of Medicinal Chemistry, 2013, 56, 3518-3530.	2.9	41
31	Structure-Based Predictions of Activity Cliffs. Journal of Chemical Information and Modeling, 2015, 55, 1062-1076.	2.5	34
32	Mapping Cholesterol Interaction Sites on Serotonin Transporter through Coarse-Grained Molecular Dynamics. PLoS ONE, 2016, 11, e0166196.	1.1	29
33	Inflammation causes remodeling of mitochondrial cytochrome <i>c</i> oxidase mediated by the bifunctional gene <i>C15orf48</i> Science Advances, 2021, 7, eabl5182.	4.7	29
34	Design, Synthesis, Structure–Activity Relationship Studies, and Three-Dimensional Quantitative Structure–Activity Relationship (3D-QSAR) Modeling of a Series of <i>O</i> belia Fig. 10 Series as Dual Modulators of Dopamine D3 Receptor and Fatty Acid Amide Hydrolase. Journal of Medicinal Chemistry, 2017, 60, 2287-2304.	2.9	28
35	Synthesis of Monomeric Derivatives To Probe Memoquin's Bivalent Interactions. Journal of Medicinal Chemistry, 2011, 54, 8299-8304.	2.9	27
36	Polo-Like Kinases Inhibitors. Current Medicinal Chemistry, 2012, 19, 3937-3948.	1.2	26

3

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37	In silico antitarget screening. Drug Discovery Today: Technologies, 2004, 1, 209-215.	4.0	24
38	Applying a multitarget rational drug design strategy: the first set of modulators with potent and balanced activity toward dopamine D3 receptor and fatty acid amide hydrolase. Chemical Communications, 2014, 50, 4904-4907.	2.2	23
39	Combining Dyad Protonation and Active Site Plasticity in BACE-1 Structure-Based Drug Design. Journal of Chemical Information and Modeling, 2012, 52, 1079-1085.	2.5	22
40	A Triazolotriazineâ€Based Dual GSKâ€3β/CKâ€1Î′ Ligand as a Potential Neuroprotective Agent Presenting Two Different Mechanisms of Enzymatic Inhibition. ChemMedChem, 2019, 14, 310-314.	1.6	22
41	Fluorinated benzophenone derivatives: Balanced multipotent agents for Alzheimer's disease. European Journal of Medicinal Chemistry, 2014, 78, 157-166.	2.6	21
42	Benzimidazole Derivatives as Kinase Inhibitors. Current Medicinal Chemistry, 2014, 21, 2284-2298.	1.2	21
43	SERAPhiC: A Benchmark for in Silico Fragment-Based Drug Design. Journal of Chemical Information and Modeling, 2011, 51, 2882-2896.	2.5	20
44	Cyclin-dependent kinases: bridging their structure and function through computations. Future Medicinal Chemistry, 2011, 3, 1551-1559.	1.1	19
45	Multitarget drug design strategy in Alzheimer's disease: focus on cholinergic transmission and amyloid-l ² aggregation. Future Medicinal Chemistry, 2017, 9, 953-963.	1.1	19
46	Multi-target dopamine D3 receptor modulators: Actionable knowledge for drug design from molecular dynamics and machine learning. European Journal of Medicinal Chemistry, 2020, 188, 111975.	2.6	19
47	Development and Application of a Virtual Screening Protocol for the Identification of Multitarget Fragments. ChemMedChem, 2016, 11, 1259-1263.	1.6	18
48	Protein-ligand docking. Frontiers in Bioscience - Landmark, 2011, 16, 2289.	3.0	17
49	Role of phosphorylated Thr160 for the activation of the CDK2/Cyclin A complex. Proteins: Structure, Function and Bioinformatics, 2005, 62, 89-98.	1.5	16
50	Synthesis, Biological Evaluation, and 3D QSAR Study of 2-Methyl-4-oxo-3-oxetanylcarbamic Acid Esters as N-Acylethanolamine Acid Amidase (NAAA) Inhibitors. Journal of Medicinal Chemistry, 2014, 57, 10101-10111.	2.9	13
51	Anandamide transport inhibition by <scp>ARN</scp> 272 attenuates nauseaâ€induced behaviour in rats, and vomiting in shrews (<i><scp>S</scp>uncus murinus</i>). British Journal of Pharmacology, 2013, 170, 1130-1136.	2.7	12
52	Describing the Conformational Landscape of Small Organic Molecules through Gaussian Mixtures in Dihedral Space. Journal of Chemical Theory and Computation, 2014, 10, 2557-2568.	2.3	10
53	Application of Conformational Clustering in Protein–Ligand Docking. Methods in Molecular Biology, 2012, 819, 169-186.	0.4	7
54	Modeling lipid raft domains containing a mono-unsaturated phosphatidylethanolamine species. RSC Advances, 2015, 5, 37102-37111.	1.7	6

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55	Discovery and SAR Evolution of Pyrazole Azabicyclo [3.2.1] octane Sulfonamides as a Novel Class of Non-Covalent N-Acylethanolamine-Hydrolyzing Acid Amidase (NAAA) Inhibitors for Oral Administration. Journal of Medicinal Chemistry, 2021, 64, 13327-13355.	2.9	6
56	Pyrazole-Based Acid Ceramidase Inhibitors: Design, Synthesis, and Structure–Activity Relationships. Synthesis, 2016, 48, 2739-2756.	1.2	4
57	Aryl and heteroaryl <i>N</i> -[4-[4-(2,3-substituted-phenyl)piperazine-1-yl]alkyl]carbamates with improved physico-chemical properties as dual modulators of dopamine D3 receptor and fatty acid amide hydrolase. MedChemComm, 2016, 7, 537-541.	3.5	4
58	Multitarget Compounds for Bipolar Disorder: From Rational Design to Preliminary Pharmacokinetic Evaluation. ChemMedChem, 2020, 15, 949-954.	1.6	4
59	The multitarget FAAH inhibitor/D3 partial agonist ARN15381 decreases nicotine self-administration in male rats. European Journal of Pharmacology, 2022, 928, 175088.	1.7	4
60	Computational Methods in Multitarget Drug Discovery. , 2017, , 239-258.		3
61	Dopamine D3 receptor ligands: a patent review (2014–2020). Expert Opinion on Therapeutic Patents, 2022, 32, 605-627.	2.4	1
62	(38) Computational approaches to the study of dual-site and peripheral site binding ache inhibitors. Chemico-Biological Interactions, 2005, 157-158, 414-415.	1.7	0