

Graziano Lolli

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

1,622
citations

331670

21
h-index

345221

36
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39
all docs

39
docs citations

39
times ranked

2383
citing authors

#	ARTICLE	IF	CITATIONS
1	The structure of P-TEFb (CDK9/cyclin T1), its complex with flavopiridol and regulation by phosphorylation. <i>EMBO Journal</i> , 2008, 27, 1907-1918.	7.8	295
2	Unprecedented Selectivity and Structural Determinants of a New Class of Protein Kinase CK2 Inhibitors in Clinical Trials for the Treatment of Cancer. <i>Biochemistry</i> , 2011, 50, 8478-8488.	2.5	154
3	CAK—Cyclin-Dependent Activating Kinase: A Key Kinase in Cell Cycle Control and a Target for Drugs?. <i>Cell Cycle</i> , 2005, 4, 565-570.	2.6	151
4	Inhibition of Protein Kinase CK2 by Flavonoids and Tyrphostins. A Structural Insight. <i>Biochemistry</i> , 2012, 51, 6097-6107.	2.5	127
5	The Crystal Structure of Human CDK7 and Its Protein Recognition Properties. <i>Structure</i> , 2004, 12, 2067-2079.	3.3	124
6	CAK-Cyclin-dependent Activating Kinase: a key kinase in cell cycle control and a target for drugs?. <i>Cell Cycle</i> , 2005, 4, 572-7.	2.6	67
7	Structural Determinants of Protein Kinase CK2 Regulation by Autoinhibitory Polymerization. <i>ACS Chemical Biology</i> , 2012, 7, 1158-1163.	3.4	58
8	Cell-permeable dual inhibitors of protein kinases CK2 and PIM-1: structural features and pharmacological potential. <i>Cellular and Molecular Life Sciences</i> , 2014, 71, 3173-3185.	5.4	45
9	Structural dissection of cyclin dependent kinases regulation and protein recognition properties. <i>Cell Cycle</i> , 2010, 9, 1551-1561.	2.6	41
10	The “Gatekeeper” Residue Influences the Mode of Binding of Acetyl Indoles to Bromodomains. <i>Journal of Medicinal Chemistry</i> , 2016, 59, 3087-3097.	6.4	36
11	GroEL from the psychrophilic bacterium <i>Pseudoalteromonas haloplanktis</i> TAC 125: molecular characterization and gene cloning. <i>Extremophiles</i> , 2003, 7, 17-28.	2.3	33
12	Inhibitor affinity chromatography: Profiling the specific reactivity of the proteome with immobilized molecules. <i>Proteomics</i> , 2003, 3, 1287-1298.	2.2	33
13	Structural and functional analysis of the flexible regions of the catalytic $\hat{\pm}$ -subunit of protein kinase CK2. <i>Journal of Structural Biology</i> , 2012, 177, 382-391.	2.8	32
14	High-Throughput Fragment Docking into the BAZ2B Bromodomain: Efficient <i>in Silico</i> Screening for X-Ray Crystallography. <i>ACS Chemical Biology</i> , 2016, 11, 800-807.	3.4	32
15	Pharmacological inactivation of the prion protein by targeting a folding intermediate. <i>Communications Biology</i> , 2021, 4, 62.	4.4	30
16	Active Form of the Protein Kinase CK2 $\hat{\pm}^2$ Holoenzyme Is a Strong Complex with Symmetric Architecture. <i>ACS Chemical Biology</i> , 2014, 9, 366-371.	3.4	29
17	Screening Approaches for Targeting Ribonucleoprotein Complexes: A New Dimension for Drug Discovery. <i>SLAS Discovery</i> , 2019, 24, 314-331.	2.7	29
18	Structural and functional determinants of protein kinase CK2 $\hat{\pm}$: facts and open questions. <i>Molecular and Cellular Biochemistry</i> , 2011, 356, 67-73.	3.1	28

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19	An Inhibitor's Eye View of the ATP-Binding Site of CDKs in Different Regulatory States. <i>ACS Chemical Biology</i> , 2014, 9, 1251-1256.	3.4	27
20	The STAS domain of mammalian SLC26A5 prestin harbours an anion-binding site. <i>Biochemical Journal</i> , 2016, 473, 365-370.	3.7	24
21	Different orientations of low-molecular-weight fragments in the binding pocket of a BRD4 bromodomain. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2013, 69, 2161-2164.	2.5	23
22	Derivatives of 3-Amino-2-methylpyridine as BAZ2B Bromodomain Ligands: In Silico Discovery and in Crystallo Validation. <i>Journal of Medicinal Chemistry</i> , 2016, 59, 9919-9927.	6.4	23
23	Recognition of Cdk2 by Cdk7. <i>Proteins: Structure, Function and Bioinformatics</i> , 2007, 67, 1048-1059.	2.6	21
24	Discovery of BAZ2A bromodomain ligands. <i>European Journal of Medicinal Chemistry</i> , 2017, 139, 564-572.	5.5	21
25	Discovery of Inhibitors of Four Bromodomains by Fragment-Anchored Ligand Docking. <i>Journal of Chemical Information and Modeling</i> , 2017, 57, 2584-2597.	5.4	21
26	Binding to DNA of the RNA-polymerase II C-terminal domain allows discrimination between Cdk7 and Cdk9 phosphorylation. <i>Nucleic Acids Research</i> , 2008, 37, 1260-1268.	14.5	18
27	Characterization of the oligomeric states of the CK2 $\hat{\pm}2^2$ holoenzyme in solution. <i>Biochemical Journal</i> , 2017, 474, 2405-2416.	3.7	17
28	A novel class of selective CK2 inhibitors targeting its open hinge conformation. <i>European Journal of Medicinal Chemistry</i> , 2020, 195, 112267.	5.5	15
29	Structure and single crystal spectroscopy of Green Fluorescent Proteins. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2011, 1814, 824-833.	2.3	12
30	Structural Analysis of Small Molecule Binding to the BAZ2A and BAZ2B Bromodomains. <i>ChemMedChem</i> , 2018, 13, 1479-1487.	3.2	11
31	Identification of compounds inhibiting prion replication and toxicity by removing PrP ^C from the cell surface. <i>Journal of Neurochemistry</i> , 2020, 152, 136-150.	3.9	11
32	The lysine-specific demethylase 1 is a novel substrate of protein kinase CK2. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2014, 1844, 722-729.	2.3	9
33	Biochemical and cellular mechanism of protein kinase CK2 inhibition by deceptive curcumin. <i>FEBS Journal</i> , 2020, 287, 1850-1864.	4.7	9
34	Inhibitory Properties of ATP-Competitive Coumestrol and Boldine Are Correlated to Different Modulations of CK2 Flexibility. <i>Journal of Natural Products</i> , 2019, 82, 1014-1018.	3.0	4
35	Insight into GFPmut2 pH Dependence by Single Crystal Microspectrophotometry and X-ray Crystallography. <i>Journal of Physical Chemistry B</i> , 2018, 122, 11326-11337.	2.6	3
36	Identification of a BAZ2A Bromodomain Hit Compound by Fragment Joining. <i>ACS Bio & Med Chem Au</i> , 2021, 1, 5-10.	3.7	3

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37	Structural Basis of CK2 Regulation by Autoinhibitory Oligomerization. , 2015, , 35-47.		2