

Andreas C Joerger

List of Publications by Year in descending order

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39
papers

4,881
citations

218381
26
h-index

301761
39
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40
all docs

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docs citations

40
times ranked

6512
citing authors

#	ARTICLE	IF	CITATIONS
1	Evolutionary history of the p53 family DNA-binding domain: insights from an <i>Alvinella pompejana</i> homolog. <i>Cell Death and Disease</i> , 2022, 13, 214.	2.7	10
2	Identification and functional characterization of new missense SNPs in the coding region of the TP53 gene. <i>Cell Death and Differentiation</i> , 2021, 28, 1477-1492.	5.0	26
3	Integrated analysis of Shank1 PDZ interactions with C-terminal and internal binding motifs. <i>Current Research in Structural Biology</i> , 2021, 3, 41-50.	1.1	4
4	Genotoxicity and Epigenotoxicity of Carbazole-Derived Molecules on MCF-7 Breast Cancer Cells. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3410.	1.8	4
5	Exploiting vulnerabilities of SWI/SNF chromatin remodelling complexes for cancer therapy. <i>Oncogene</i> , 2021, 40, 3637-3654.	2.6	66
6	Structure-Based Design of Selective Salt-Inducible Kinase Inhibitors. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 8142-8160.	2.9	28
7	Development of a Selective Dual Discoidin Domain Receptor (DDR)/p38 Kinase Chemical Probe. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 13451-13474.	2.9	4
8	Selective targeting of the $\hat{\pm}$ C and DFG-out pocket in p38 MAPK. <i>European Journal of Medicinal Chemistry</i> , 2020, 208, 112721.	2.6	12
9	Pan-SMARCA/PB1 Bromodomain Inhibitors and Their Role in Regulating Adipogenesis. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 14680-14699.	2.9	21
10	Characterization of a dual \langle scp \rangle BET \langle /scp \rangle / \langle scp \rangle HDAC \langle /scp \rangle inhibitor for treatment of pancreatic ductal adenocarcinoma. <i>International Journal of Cancer</i> , 2020, 147, 2847-2861.	2.3	34
11	Targeting Cavity-Creating p53 Cancer Mutations with Small-Molecule Stabilizers: the Y220X Paradigm. <i>ACS Chemical Biology</i> , 2020, 15, 657-668.	1.6	45
12	New pyrido[3,4-g]quinazoline derivatives as CLK1 and DYRK1A inhibitors: synthesis, biological evaluation and binding mode analysis. <i>European Journal of Medicinal Chemistry</i> , 2019, 166, 304-317.	2.6	32
13	A structure-guided molecular chaperone approach for restoring the transcriptional activity of the p53 cancer mutant Y220C. <i>Future Medicinal Chemistry</i> , 2019, 11, 2491-2504.	1.1	53
14	Aminobenzothiazole derivatives stabilize the thermolabile p53 cancer mutant Y220C and show anticancer activity in p53-Y220C cell lines. <i>European Journal of Medicinal Chemistry</i> , 2018, 152, 101-114.	2.6	57
15	Extending the Code of Sequence Readout by Gene Regulatory Proteins: The Role of Hoogsteen Base Pairing in p53-DNA Recognition. <i>Structure</i> , 2018, 26, 1163-1165.	1.6	3
16	The p53 Pathway: Origins, Inactivation in Cancer, and Emerging Therapeutic Approaches. <i>Annual Review of Biochemistry</i> , 2016, 85, 375-404.	5.0	483
17	2-Sulfonylpyrimidines: Mild alkylating agents with anticancer activity toward p53-compromised cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E5271-80.	3.3	88
18	Design of a molecular support for cryo-EM structure determination. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E7456-E7463.	3.3	93

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19	Harnessing Fluorine-Sulfur Contacts and Multipolar Interactions for the Design of p53 Mutant Y220C Rescue Drugs. <i>ACS Chemical Biology</i> , 2016, 11, 2265-2274.	1.6	56
20	Exploiting Transient Protein States for the Design of Small-Molecule Stabilizers of Mutant p53. <i>Structure</i> , 2015, 23, 2246-2255.	1.6	45
21	Experimental and Theoretical Evaluation of the Ethynyl Moiety as a Halogen Bioisostere. <i>ACS Chemical Biology</i> , 2015, 10, 2725-2732.	1.6	23
22	Tracing the Evolution of the p53 Tetramerization Domain. <i>Structure</i> , 2014, 22, 1301-1310.	1.6	27
23	Principles and Applications of Halogen Bonding in Medicinal Chemistry and Chemical Biology. <i>Journal of Medicinal Chemistry</i> , 2013, 56, 1363-1388.	2.9	1,002
24	Small molecule induced reactivation of mutant p53 in cancer cells. <i>Nucleic Acids Research</i> , 2013, 41, 6034-6044.	6.5	187
25	Evaluating <i>Drosophila</i> p53 as a Model System for Studying Cancer Mutations. <i>Journal of Biological Chemistry</i> , 2012, 287, 44330-44337.	1.6	13
26	Structure and Kinetic Stability of the p63 Tetramerization Domain. <i>Journal of Molecular Biology</i> , 2012, 415, 503-513.	2.0	31
27	Halogen-Enriched Fragment Libraries as Leads for Drug Rescue of Mutant p53. <i>Journal of the American Chemical Society</i> , 2012, 134, 6810-6818.	6.6	202
28	Interaction of the p53 DNA-Binding Domain with Its N-Terminal Extension Modulates the Stability of the p53 Tetramer. <i>Journal of Molecular Biology</i> , 2011, 409, 358-368.	2.0	81
29	Acetylation of lysine 120 of p53 endows DNA-binding specificity at effective physiological salt concentration. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 8251-8256.	3.3	81
30	Toward the Rational Design of p53-Stabilizing Drugs: Probing the Surface of the Oncogenic Y220C Mutant. <i>Chemistry and Biology</i> , 2010, 17, 46-56.	6.2	97
31	The Tumor Suppressor p53: From Structures to Drug Discovery. <i>Cold Spring Harbor Perspectives in Biology</i> , 2010, 2, a000919-a000919.	2.3	273
32	Structural evolution of p53, p63, and p73: Implication for heterotetramer formation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 17705-17710.	3.3	133
33	Stabilising the DNA-binding domain of p53 by rational design of its hydrophobic core. <i>Protein Engineering, Design and Selection</i> , 2009, 22, 421-430.	1.0	21
34	Structural Biology of the Tumor Suppressor p53. <i>Annual Review of Biochemistry</i> , 2008, 77, 557-582.	5.0	549
35	Targeted rescue of a destabilized mutant of p53 by an <i>in silico</i> screened drug. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 10360-10365.	3.3	319
36	Structural basis for understanding oncogenic p53 mutations and designing rescue drugs. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 15056-15061.	3.3	267

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37	Effects of Common Cancer Mutations on Stability and DNA Binding of Full-length p53 Compared with Isolated Core Domains. <i>Journal of Biological Chemistry</i> , 2006, 281, 21934-21941.	1.6	119
38	Structures of p53 Cancer Mutants and Mechanism of Rescue by Second-site Suppressor Mutations. <i>Journal of Biological Chemistry</i> , 2005, 280, 16030-16037.	1.6	144
39	Crystal Structure of a Superstable Mutant of Human p53 Core Domain. <i>Journal of Biological Chemistry</i> , 2004, 279, 1291-1296.	1.6	147