

Athanassios S Galanis

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7197266/publications.pdf>

Version: 2024-02-01

11
papers

530
citations

1305906

8
h-index

1427216

11
g-index

12
all docs

12
docs citations

12
times ranked

925
citing authors

#	ARTICLE	IF	CITATIONS
1	Structural analysis of metastable pharmaceutical loratadine form II, by 3D electron diffraction and DFT+D energy minimisation. <i>CrystEngComm</i> , 2020, 22, 7490-7499.	1.3	13
2	Fast electron diffraction tomography. <i>Journal of Applied Crystallography</i> , 2015, 48, 718-727.	1.9	134
3	Synthetic Peptides as Structural Maquettes of Angiotensin-I Converting Enzyme Catalytic Sites. <i>Bioinorganic Chemistry and Applications</i> , 2010, 2010, 1-13.	1.8	3
4	Enhanced microwave-assisted method for on-bead disulfide bond formation: Synthesis of Î±-conotoxin MII. <i>Biopolymers</i> , 2009, 92, 23-34.	1.2	29
5	Manufacturing peptides as active pharmaceutical ingredients. <i>Future Medicinal Chemistry</i> , 2009, 1, 361-377.	1.1	151
6	Solid-Phase Peptide Synthesis in Water Using Microwave-Assisted Heating. <i>Organic Letters</i> , 2009, 11, 4488-4491.	2.4	77
7	[^{99m} Tc]Demotensin 5 and 6 in the NTS1-R-targeted imaging of tumours: synthesis and preclinical results. <i>European Journal of Nuclear Medicine and Molecular Imaging</i> , 2007, 34, 1804-1814.	3.3	37
8	Amyloid fibril formation propensity is inherent into the hexapeptide tandemly repeating sequence of the central domain of silkmoth chorion proteins of the A-family. <i>Journal of Structural Biology</i> , 2006, 156, 480-488.	1.3	39
9	Structural Features of Angiotensin-I Converting Enzyme Catalytic Sites: Conformational Studies in Solution, Homology Models and Comparison with Other Zinc Metallopeptidases. <i>Current Topics in Medicinal Chemistry</i> , 2004, 4, 403-429.	1.0	27
10	Solid-phase synthesis and conformational properties of angiotensin converting enzyme catalytic-site peptides: The basis for a structural study on the enzyme-substrate interaction. <i>Biopolymers</i> , 2004, 76, 512-526.	1.2	7
11	Zinc binding in peptide models of angiotensin-I converting enzyme active sites studied through ¹ H-NMR and chemical shift perturbation mapping. <i>Biopolymers</i> , 2003, 69, 244-252.	1.2	8