

Andr s Asz di

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2537708/publications.pdf>

Version: 2024-02-01

23
papers

991
citations

759233

12
h-index

677142

22
g-index

23
all docs

23
docs citations

23
times ranked

1459
citing authors

#	ARTICLE	IF	CITATIONS
1	MULTOVL: fast multiple overlaps of genomic regions. <i>Bioinformatics</i> , 2012, 28, 3318-3319.	4.1	36
2	Intergenic Polycomb target sites are dynamically marked by non-coding transcription during lineage commitment. <i>RNA Biology</i> , 2012, 9, 314-325.	3.1	8
3	H3K27me3 forms BLOCs over silent genes and intergenic regions and specifies a histone banding pattern on a mouse autosomal chromosome. <i>Genome Research</i> , 2009, 19, 221-233.	5.5	212
4	Analyzing the performance of conformational search programs on compound databases. <i>Journal of Molecular Graphics and Modelling</i> , 2007, 25, 700-710.	2.4	18
5	SH2 Binding Site Comparison: A New Application of the SURFCOMP Method. <i>Journal of Chemical Information and Modeling</i> , 2005, 45, 414-421.	5.4	4
6	mRNA Openers and Closers: Modulating AU-Rich Element-Controlled mRNA Stability by a Molecular Switch in mRNA Secondary Structure. <i>ChemBioChem</i> , 2004, 5, 1432-1447.	2.6	112
7	SURFCOMP: A Novel Graph-Based Approach to Molecular Surface Comparison.. <i>ChemInform</i> , 2004, 35, no.	0.0	0
8	Percutaneous absorption of drugs used in atopic eczema: pimecrolimus permeates less through skin than corticosteroids and tacrolimus. <i>International Journal of Pharmaceutics</i> , 2004, 269, 29-35.	5.2	168
9	SURFCOMP: A Novel Graph-Based Approach to Molecular Surface Comparison. <i>Journal of Chemical Information and Computer Sciences</i> , 2004, 44, 837-847.	2.8	44
10	Parallel Global Optimization of High-Dimensional Problems. <i>Lecture Notes in Computer Science</i> , 2002, , 148-155.	1.3	3
11	Protein structure: geometry, topology and classification. <i>Reports on Progress in Physics</i> , 2001, 64, 517-590.	20.1	52
12	Distance geometry based comparative modelling. <i>Folding & Design</i> , 1997, 2, S3-S6.	4.5	11
13	Hierarchic inertial projection: A fast distance matrix embedding algorithm. <i>Computers & Chemistry</i> , 1997, 21, 13-23.	1.2	10
14	Protein modeling by multiple sequence threading and distance geometry. <i>Proteins: Structure, Function and Bioinformatics</i> , 1997, 29, 38-42.	2.6	10
15	Estimating polypeptide-carbon distances from multiple sequence alignments. <i>Journal of Mathematical Chemistry</i> , 1995, 17, 167-184.	1.5	9
16	Global Fold Determination from a Small Number of Distance Restraints. <i>Journal of Molecular Biology</i> , 1995, 251, 308-326.	4.2	128
17	Secondary structure formation in model polypeptide chains. <i>Protein Engineering, Design and Selection</i> , 1994, 7, 633-644.	2.1	42
18	Calpeptin, a calpain inhibitor, promotes neurite elongation in differentiating PC12 cells. <i>Neuroscience Letters</i> , 1994, 170, 91-93.	2.1	19

#	ARTICLE	IF	CITATIONS
19	Calcium-dependent proteolysis and isopeptide bond formation: Calpains and transglutaminases. <i>Pure and Applied Chemistry</i> , 1992, 64, 1093-1097.	1.9	2
20	Detecting changes in neuronal activities induced by N-methyl-d-aspartate receptor blockade using non-linear dynamics techniques. <i>Neuroscience</i> , 1992, 46, 785-791.	2.3	2
21	MAP2: a sensitive cross-linker and adjustable spacer in dendritic architecture. <i>FEBS Letters</i> , 1991, 295, 5-9.	2.8	61
22	Signal convergence on protein kinase A as a molecular correlate of learning.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1991, 88, 5832-5836.	7.1	32
23	Molecular kinetic modelling of associative learning. <i>Neuroscience</i> , 1987, 22, 37-48.	2.3	8