

Claudia Andreini

List of Publications by Year in descending order

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Version: 2024-02-01

47
papers

4,907
citations

159585

30
h-index

223800

46
g-index

50
all docs

50
docs citations

50
times ranked

5950
citing authors

#	ARTICLE	IF	CITATIONS
1	PDBe-KB: collaboratively defining the biological context of structural data. <i>Nucleic Acids Research</i> , 2022, 50, D534-D542.	14.5	46
2	The Intriguing Role of Iron-Sulfur Clusters in the CIAPIN1 Protein Family. <i>Inorganics</i> , 2022, 10, 52.	2.7	1
3	A New Paradigm of Multiheme Cytochrome Evolution by Grafting and Pruning Protein Modules. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	12
4	Learning to Identify Physiological and Adventitious Metal-Binding Sites in the Three-Dimensional Structures of Proteins by Following the Hints of a Deep Neural Network. <i>Journal of Chemical Information and Modeling</i> , 2022, 62, 2951-2960.	5.4	6
5	The zinc proteome of SARS-CoV-2. <i>Metallomics</i> , 2022, 14, .	2.4	6
6	Structural Bioinformatics and Deep Learning of Metalloproteins: Recent Advances and Applications. <i>International Journal of Molecular Sciences</i> , 2022, 23, 7684.	4.1	6
7	HIV-1 Tat Protein Enters Dysfunctional Endothelial Cells via Integrins and Renders Them Permissive to Virus Replication. <i>International Journal of Molecular Sciences</i> , 2021, 22, 317.	4.1	12
8	Upgraded AMBER Force Field for Zinc-Binding Residues and Ligands for Predicting Structural Properties and Binding Affinities in Zinc-Proteins. <i>ACS Omega</i> , 2020, 5, 15301-15310.	3.5	27
9	7. Basic Iron-Sulfur Centers. , 2020, 20, 199-256.		2
10	Upgrading and Validation of the AMBER Force Field for Histidine and Cysteine Zinc(II)-Binding Residues in Sites with Four Protein Ligands. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 3803-3816.	5.4	42
11	Multi-metal Restriction by Calprotectin Impacts De Novo Flavin Biosynthesis in <i>Acinetobacter baumannii</i> . <i>Cell Chemical Biology</i> , 2019, 26, 745-755.e7.	5.2	61
12	MetalPDB in 2018: a database of metal sites in biological macromolecular structures. <i>Nucleic Acids Research</i> , 2018, 46, D459-D464.	14.5	165
13	To what extent do structural changes in catalytic metal sites affect enzyme function?. <i>Journal of Inorganic Biochemistry</i> , 2018, 179, 40-53.	3.5	55
14	The cellular economy of the <i>Saccharomyces cerevisiae</i> zinc proteome. <i>Metallomics</i> , 2018, 10, 1755-1776.	2.4	66
15	The human iron-proteome. <i>Metallomics</i> , 2018, 10, 1223-1231.	2.4	106
16	Identification of the zinc, copper and cadmium metalloproteome of the protozoan <i>Tetrahymena thermophila</i> by systematic bioinformatics. <i>Archives of Microbiology</i> , 2017, 199, 1141-1149.	2.2	24
17	Differential Effects of Iron, Zinc, and Copper on <i>Dictyostelium discoideum</i> Cell Growth and Resistance to <i>Legionella pneumophila</i> . <i>Frontiers in Cellular and Infection Microbiology</i> , 2017, 7, 536.	3.9	25
18	The Relationship between Environmental Dioxygen and Iron-Sulfur Proteins Explored at the Genome Level. <i>PLoS ONE</i> , 2017, 12, e0171279.	2.5	49

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19	Minimal Functional Sites in Metalloproteins and Their Usage in Structural Bioinformatics. <i>International Journal of Molecular Sciences</i> , 2016, 17, 671.	4.1	12
20	MetalPredator: a web server to predict iron-sulfur cluster binding proteomes. <i>Bioinformatics</i> , 2016, 32, 2850-2852.	4.1	58
21	Exploiting Bacterial Operons To Illuminate Human Iron-Sulfur Proteins. <i>Journal of Proteome Research</i> , 2016, 15, 1308-1322.	3.7	42
22	Hidden relationships between metalloproteins unveiled by structural comparison of their metal sites. <i>Scientific Reports</i> , 2015, 5, 9486.	3.3	13
23	MetalS3, a database-mining tool for the identification of structurally similar metal sites. <i>Journal of Biological Inorganic Chemistry</i> , 2014, 19, 937-945.	2.6	28
24	MetalS ² : A Tool for the Structural Alignment of Minimal Functional Sites in Metal-Binding Proteins and Nucleic Acids. <i>Journal of Chemical Information and Modeling</i> , 2013, 53, 3064-3075.	5.4	16
25	MACiE: exploring the diversity of biochemical reactions. <i>Nucleic Acids Research</i> , 2012, 40, D783-D789.	14.5	73
26	FindGeo: a tool for determining metal coordination geometry. <i>Bioinformatics</i> , 2012, 28, 1658-1660.	4.1	45
27	MetalPDB: a database of metal sites in biological macromolecular structures. <i>Nucleic Acids Research</i> , 2012, 41, D312-D319.	14.5	157
28	A bioinformatics view of zinc enzymes. <i>Journal of Inorganic Biochemistry</i> , 2012, 111, 150-156.	3.5	168
29	HIV-1 Tat Promotes Integrin-Mediated HIV Transmission to Dendritic Cells by Binding Env Spikes and Competes Neutralization by Anti-HIV Antibodies. <i>PLoS ONE</i> , 2012, 7, e48781.	2.5	56
30	A Simple Protocol for the Comparative Analysis of the Structure and Occurrence of Biochemical Pathways Across Superkingdoms. <i>Journal of Chemical Information and Modeling</i> , 2011, 51, 730-738.	5.4	28
31	Minimal Functional Sites Allow a Classification of Zinc Sites in Proteins. <i>PLoS ONE</i> , 2011, 6, e26325.	2.5	113
32	Metal-MACiE: a database of metals involved in biological catalysis. <i>Bioinformatics</i> , 2009, 25, 2088-2089.	4.1	73
33	Structural Analysis of Metal Sites in Proteins: Non-heme Iron Sites as a Case Study. <i>Journal of Molecular Biology</i> , 2009, 388, 356-380.	4.2	48
34	Metalloproteomes: A Bioinformatic Approach. <i>Accounts of Chemical Research</i> , 2009, 42, 1471-1479.	15.6	281
35	Metal ions in biological catalysis: from enzyme databases to general principles. <i>Journal of Biological Inorganic Chemistry</i> , 2008, 13, 1205-1218.	2.6	868
36	Occurrence of Copper Proteins through the Three Domains of Life: A Bioinformatic Approach. <i>Journal of Proteome Research</i> , 2008, 7, 209-216.	3.7	184

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37	Mycobacterial Cells Have Dual Nickel-Cobalt Sensors. <i>Journal of Biological Chemistry</i> , 2007, 282, 32298-32310.	3.4	91
38	Non-heme iron through the three domains of life. <i>Proteins: Structure, Function and Bioinformatics</i> , 2007, 67, 317-324.	2.6	70
39	Predicting zinc binding at the proteome level. <i>BMC Bioinformatics</i> , 2007, 8, 39.	2.6	89
40	Counting the Zinc-Proteins Encoded in the Human Genome. <i>Journal of Proteome Research</i> , 2006, 5, 196-201.	3.7	887
41	Zinc through the Three Domains of Life. <i>Journal of Proteome Research</i> , 2006, 5, 3173-3178.	3.7	544
42	SPINE bioinformatics and data-management aspects of high-throughput structural biology. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2006, 62, 1184-1195.	2.5	19
43	Predicting metals sensed by ArsR&SmtB repressors: allosteric interference by a non-effector metal. <i>Molecular Microbiology</i> , 2006, 59, 1341-1356.	2.5	40
44	Comparative Analysis of the ADAM and ADAMTS Families. <i>Journal of Proteome Research</i> , 2005, 4, 881-888.	3.7	32
45	A hint to search for metalloproteins in gene banks. <i>Bioinformatics</i> , 2004, 20, 1373-1380.	4.1	120
46	Bioinformatic Comparison of Structures and Homology-Models of Matrix Metalloproteinases. <i>Journal of Proteome Research</i> , 2004, 3, 21-31.	3.7	35
47	Systematic classification of metalloproteins based on three-dimensional structural similarity of their metal sites. <i>Protocol Exchange</i> , 0, , .	0.3	1