

Paola Turano

List of Publications by Year in descending order

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162
papers

7,218
citations

53660

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69108

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175
docs citations

175
times ranked

6848
citing authors

#	ARTICLE	IF	CITATIONS
1	Ferritin nanocomposites for the selective delivery of photosensitizing ruthenium-polypyridyl compounds to cancer cells. <i>Inorganic Chemistry Frontiers</i> , 2022, 9, 1070-1081.	3.0	14
2	Impact of the pre-examination phase on multicenter metabolomic studies. <i>New Biotechnology</i> , 2022, 68, 37-47.	2.4	10
3	Metabolite and lipoprotein profiles reveal sex-related oxidative stress imbalance in de novo drug-naïve Parkinson's disease patients. <i>Npj Parkinson's Disease</i> , 2022, 8, 14.	2.5	11
4	S1P Signalling Axis Is Necessary for Adiponectin-Directed Regulation of Electrophysiological Properties and Oxidative Metabolism in C2C12 Myotubes. <i>Cells</i> , 2022, 11, 713.	1.8	8
5	Serum or Plasma (and Which Plasma), That Is the Question. <i>Journal of Proteome Research</i> , 2022, 21, 1061-1072.	1.8	25
6	Serum NMR Profiling Reveals Differential Alterations in the Lipoproteome Induced by Pfizer-BioNTech Vaccine in COVID-19 Recovered Subjects and Naïve Subjects. <i>Frontiers in Molecular Biosciences</i> , 2022, 9, 839809.	1.6	11
7	Profiling metabolites and lipoproteins in COMETA, an Italian cohort of COVID-19 patients. <i>PLoS Pathogens</i> , 2022, 18, e1010443.	2.1	30
8	NMR reveals the metabolic changes induced by auranofin in A2780 cancer cells: evidence for glutathione dysregulation. <i>Dalton Transactions</i> , 2021, 50, 6349-6355.	1.6	17
9	Metabolomic/lipidomic profiling of COVID-19 and individual response to tocilizumab. <i>PLoS Pathogens</i> , 2021, 17, e1009243.	2.1	76
10	A geroscience approach for Parkinson's disease: Conceptual framework and design of PROPAG-AGEING project. <i>Mechanisms of Ageing and Development</i> , 2021, 194, 111426.	2.2	14
11	Prediagnostic circulating metabolites in female breast cancer cases with low and high mammographic breast density. <i>Scientific Reports</i> , 2021, 11, 13025.	1.6	10
12	Modelling hCDKL5 Heterologous Expression in Bacteria. <i>Metabolites</i> , 2021, 11, 491.	1.3	5
13	Metabolomic Fingerprints in Large Population Cohorts: Impact of Preanalytical Heterogeneity. <i>Clinical Chemistry</i> , 2021, 67, 1153-1155.	1.5	10
14	Iron Binding in the Ferroxidase Site of Human Mitochondrial Ferritin. <i>Chemistry - A European Journal</i> , 2021, 27, 14690-14701.	1.7	2
15	Direct detection of iron clusters in L ferritins through ESI-MS experiments. <i>Dalton Transactions</i> , 2021, 50, 16464-16467.	1.6	6
16	A framework for validating AI in precision medicine: considerations from the European ITFoC consortium. <i>BMC Medical Informatics and Decision Making</i> , 2021, 21, 274.	1.5	28
17	DNA damage response protein checkpoint kinase 2 (CHK2) links chromosomal instability to cellular metabolism in hepatocellular carcinoma (HCC). <i>Journal of Hepatology</i> , 2020, 73, S639-S640.	1.8	0
18	Distal Unfolding of Ferricytochrome c Induced by the F82K Mutation. <i>International Journal of Molecular Sciences</i> , 2020, 21, 2134.	1.8	7

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19	Plasma metabolome and cognitive skills in Down syndrome. <i>Scientific Reports</i> , 2020, 10, 10491.	1.6	23
20	Diauxie and co-utilization of carbon sources can coexist during bacterial growth in nutritionally complex environments. <i>Nature Communications</i> , 2020, 11, 3135.	5.8	51
21	Iron Biomineral Growth from the Initial Nucleation Seed in Lâ€Ferritin. <i>Chemistry - A European Journal</i> , 2020, 26, 5770-5773.	1.7	17
22	Fingerprinting Alzheimerâ€™s Disease by ¹ H Nuclear Magnetic Resonance Spectroscopy of Cerebrospinal Fluid. <i>Journal of Proteome Research</i> , 2020, 19, 1696-1705.	1.8	32
23	Effect of Estrogen Receptor Status on Circulatory Immune and Metabolomics Profiles of HER2-Positive Breast Cancer Patients Enrolled for Neoadjuvant Targeted Chemotherapy. <i>Cancers</i> , 2020, 12, 314.	1.7	22
24	Highâ€Throughput Metabolomics by 1D NMR. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 968-994.	7.2	254
25	Hochdurchsatzâ€Metabolomik mit 1Dâ€NMR. <i>Angewandte Chemie</i> , 2019, 131, 980-1007.	1.6	8
26	¹ H-NMR metabolomics reveals the Glabrescione B exacerbation of glycolytic metabolism beside the cell growth inhibitory effect in glioma. <i>Cell Communication and Signaling</i> , 2019, 17, 108.	2.7	30
27	Structural Biology of Iron-Binding Proteins by NMR Spectroscopy. <i>European Journal of Inorganic Chemistry</i> , 2019, 2019, 569-576.	1.0	4
28	Pre-analytical processes in medical diagnostics: New regulatory requirements and standards. <i>New Biotechnology</i> , 2019, 52, 121-125.	2.4	35
29	Effect of the point mutation H54N on the ferroxidase process of <i>Rana catesbeiana</i> Hâ€ ² ferritin. <i>Journal of Inorganic Biochemistry</i> , 2019, 197, 110697.	1.5	4
30	NMR for sample quality assessment in metabolomics. <i>New Biotechnology</i> , 2019, 52, 25-34.	2.4	49
31	About the use of ¹³ C- ¹³ C NOESY in bioinorganic chemistry. <i>Journal of Inorganic Biochemistry</i> , 2019, 192, 25-32.	1.5	10
32	Uniqueness of the NMR approach to metabolomics. <i>TrAC - Trends in Analytical Chemistry</i> , 2019, 120, 115300.	5.8	103
33	Plasma and urinary metabolomic profiles of Down syndrome correlate with alteration of mitochondrial metabolism. <i>Scientific Reports</i> , 2018, 8, 2977.	1.6	80
34	Cancer cell death induced by ferritins and the peculiar role of their labile iron pool. <i>Oncotarget</i> , 2018, 9, 27974-27984.	0.8	12
35	Creation and Characterization of a Genomically Hybrid Strain in the Nitrogen-Fixing Symbiotic Bacterium <i>Sinorhizobium meliloti</i> . <i>ACS Synthetic Biology</i> , 2018, 7, 2365-2378.	1.9	24
36	Chemistry at the proteinâ€mineral interface in L-ferritin assists the assembly of a functional (1/4) Tj ETQqO O O rgBT /Overlock 10 Tf 50 Academy of Sciences of the United States of America, 2017, 114, 2580-2585.	3.3	74

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37	NMR metabolomics highlights sphingosine kinase as a new molecular switch in the orchestration of aberrant metabolic phenotype in cancer cells. <i>Molecular Oncology</i> , 2017, 11, 517-533.	2.1	35
38	Structural basis of mitochondrial dysfunction in response to cytochrome <i>c</i> phosphorylation at tyrosine 48. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E3041-E3050.	3.3	53
39	Unsaturated Long-Chain Fatty Acids Are Preferred Ferritin Ligands That Enhance Iron Biom mineralization. <i>Chemistry - A European Journal</i> , 2017, 23, 9879-9887.	1.7	10
40	Quality Matters: 2016 Annual Conference of the National Infrastructures for Biobanking, Biopreservation and Biobanking, 2017, 15, 270-276.	0.5	26
41	Investigation of the Iron(II) Release Mechanism of Human H-Ferritin as a Function of pH. <i>Journal of Chemical Information and Modeling</i> , 2017, 57, 2112-2118.	2.5	22
42	NMR of Paramagnetic Species. , 2017, , 164-169.		0
43	Targeting sphingosine kinase 1 localization as novel target for ovarian cancer therapy. <i>Translational Cancer Research</i> , 2017, 6, S1277-S1280.	0.4	4
44	Solid-State NMR of PEGylated Proteins. <i>Angewandte Chemie - International Edition</i> , 2016, 55, 2446-2449.	7.2	41
45	Solid-State NMR of PEGylated Proteins. <i>Angewandte Chemie</i> , 2016, 128, 2492-2495.	1.6	12
46	Electrostatic and Structural Bases of Fe ²⁺ Translocation through Ferritin Channels. <i>Journal of Biological Chemistry</i> , 2016, 291, 25617-25628.	1.6	46
47	Insights into Interprotein Electron Transfer of Human Cytochrome <i>c</i> Variants Arranged in Multilayer Architectures by Means of an Artificial Silica Nanoparticle Matrix. <i>ACS Omega</i> , 2016, 1, 1058-1066.	1.6	11
48	Individual Human Metabolic Phenotype Analyzed by ¹ H NMR of Saliva Samples. <i>Journal of Proteome Research</i> , 2016, 15, 1787-1793.	1.8	38
49	Ferroxidase Activity in Eukaryotic Ferritin is Controlled by Accessory Iron-Binding Sites in the Catalytic Cavity. <i>Chemistry - A European Journal</i> , 2016, 22, 16213-16219.	1.7	18
50	Modulating the permeability of ferritin channels. <i>RSC Advances</i> , 2016, 6, 21219-21227.	1.7	27
51	Multi-omic profiles of human non-alcoholic fatty liver disease tissue highlight heterogenic phenotypes. <i>Scientific Data</i> , 2015, 2, 150068.	2.4	48
52	The Da Vinci European BioBank: A Metabolomics-Driven Infrastructure. <i>Journal of Personalized Medicine</i> , 2015, 5, 107-119.	1.1	9
53	COordination of Standards in MetabOlomicS (COSMOS): facilitating integrated metabolomics data access. <i>Metabolomics</i> , 2015, 11, 1587-1597.	1.4	140
54	Metabolomics profiling of pre-and post-anesthesia plasma samples of colorectal patients obtained via Ficoll separation. <i>Metabolomics</i> , 2015, 11, 1769-1778.	1.4	32

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55	Iron binding to human heavy-chain ferritin. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2015, 71, 1909-1920.	2.5	68
56	Standardizing the experimental conditions for using urine in NMR-based metabolomic studies with a particular focus on diagnostic studies: a review. <i>Metabolomics</i> , 2015, 11, 872-894.	1.4	196
57	Is His54 a gating residue for the ferritin ferroxidase site?. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2015, 1854, 1118-1122.	1.1	17
58	The impact of free or standardized lifestyle and urine sampling protocol on metabolome recognition accuracy. <i>Genes and Nutrition</i> , 2015, 10, 441.	1.2	29
59	Time-lapse anomalous X-ray diffraction shows how Fe ²⁺ substrate ions move through ferritin protein nanocages to oxidoreductase sites. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2015, 71, 941-953.	2.5	49
60	Transient iron coordination sites in proteins: Exploiting the dual nature of paramagnetic NMR. <i>Coordination Chemistry Reviews</i> , 2015, 284, 313-328.	9.5	27
61	A Relay Network of Extracellular Heme-Binding Proteins Drives <i>C. albicans</i> Iron Acquisition from Hemoglobin. <i>PLoS Pathogens</i> , 2014, 10, e1004407.	2.1	87
62	Colorectal cancer: the potential of metabolic fingerprinting. <i>Expert Review of Gastroenterology and Hepatology</i> , 2014, 8, 847-849.	1.4	13
63	Loop Electrostatics Modulates the Intersubunit Interactions in Ferritin. <i>ACS Chemical Biology</i> , 2014, 9, 2517-2525.	1.6	18
64	Coordinating subdomains of ferritin protein cages with catalysis and biomineralization viewed from the C 4 cage axes. <i>Journal of Biological Inorganic Chemistry</i> , 2014, 19, 615-622.	1.1	18
65	Superoxide Reductase: Different Interaction Modes with its Two Redox Partners. <i>ChemBioChem</i> , 2013, 14, 1858-1866.	1.3	10
66	Solution and Solid State NMR Approaches To Draw Iron Pathways in the Ferritin Nanocage. <i>Accounts of Chemical Research</i> , 2013, 46, 2676-2685.	7.6	27
67	Mechanistic insights into the superoxide- cytochrome c reaction by lysine surface scanning. <i>Journal of Biological Inorganic Chemistry</i> , 2013, 18, 429-440.	1.1	9
68	Effects of Intra- and Post-Operative Ischemia on the Metabolic Profile of Clinical Liver Tissue Specimens Monitored by NMR. <i>Journal of Proteome Research</i> , 2013, 12, 5723-5729.	1.8	39
69	Cytochrome c and superoxide: a reply. <i>Journal of Biological Inorganic Chemistry</i> , 2013, 18, 867-869.	1.1	0
70	Nuclear Magnetic Resonance as a Tool to Characterize the Interactome of Heme Proteins. <i>Handbook of Porphyrin Science</i> , 2013, , 179-219.	0.3	0
71	Role of the iron axial ligands of heme carrier HasA in heme uptake and release.. <i>Journal of Biological Chemistry</i> , 2013, 288, 2190.	1.6	1
72	Cage redesign explains assembly. <i>Nature Chemical Biology</i> , 2013, 9, 143-144.	3.9	9

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73	Electron self-exchange of cytochrome c measured via ¹³ C detected protonless NMR. <i>Journal of Porphyrins and Phthalocyanines</i> , 2013, 17, 142-149.	0.4	4
74	Soluble Variants of Human Recombinant Glutaminyl Cyclase. <i>PLoS ONE</i> , 2013, 8, e71657.	1.1	4
75	NMR as a Tool to Target Protein-Protein Interactions. , 2013, , 83-111.		0
76	A Systems Biology Approach to Deciphering the Etiology of Steatosis Employing Patient-Derived Dermal Fibroblasts and iPS Cells. <i>Frontiers in Physiology</i> , 2012, 3, 339.	1.3	22
77	Metabolomic NMR Fingerprinting to Identify and Predict Survival of Patients with Metastatic Colorectal Cancer. <i>Cancer Research</i> , 2012, 72, 356-364.	0.4	181
78	Role of the Iron Axial Ligands of Heme Carrier HasA in Heme Uptake and Release. <i>Journal of Biological Chemistry</i> , 2012, 287, 26932-26943.	1.6	32
79	What Can be Learned about the Structure and Dynamics of Biomolecules from NMR. , 2012, , 33-50.		1
80	Insights in the (un)structural organization of <i>Bacillus pasteurii</i> UreG, an intrinsically disordered GTPase enzyme. <i>Molecular BioSystems</i> , 2012, 8, 220-228.	2.9	44
81	NMR properties of sedimented solutes. <i>Physical Chemistry Chemical Physics</i> , 2012, 14, 439-447.	1.3	47
82	Structural Insights into the Ferroxidase Site of Ferritins from Higher Eukaryotes. <i>Journal of the American Chemical Society</i> , 2012, 134, 6169-6176.	6.6	90
83	Electroactive Multilayer Assemblies of Bilirubin Oxidase and Human Cytochrome C Mutants: Insight in Formation and Kinetic Behavior. <i>Langmuir</i> , 2011, 27, 4202-4211.	1.6	38
84	The Anti-Apoptotic Bcl-xL Protein, a New Piece in the Puzzle of Cytochrome C Interactome. <i>PLoS ONE</i> , 2011, 6, e18329.	1.1	44
85	Standard operating procedures for pre-analytical handling of blood and urine for metabolomic studies and biobanks. <i>Journal of Biomolecular NMR</i> , 2011, 49, 231-243.	1.6	285
86	Solid-state NMR of proteins sedimented by ultracentrifugation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 10396-10399.	3.3	163
87	Fragmenting the S100B-p53 Interaction: Combined Virtual/Biophysical Screening Approaches to Identify Ligands. <i>ChemMedChem</i> , 2010, 5, 428-435.	1.6	22
88	NMR reveals pathway for ferric mineral precursors to the central cavity of ferritin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 545-550.	3.3	143
89	Heme Acquisition by Hemophores: A Lesson from NMR. <i>Handbook of Porphyrin Science</i> , 2010, , 339-365.	0.3	1
90	Computational Study of the DNA-Binding Protein <i>Helicobacter pylori</i> NikR: The Role of Ni ²⁺ 2 Francesco Musiani and Branimir Bertoja contributed equally to the simulations presented here.. <i>Journal of Chemical Theory and Computation</i> , 2010, 6, 3503-3515.	2.3	32

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91	<i>Helicobacter pylori</i> UreE, a urease accessory protein: specific Ni ²⁺ - and Zn ²⁺ -binding properties and interaction with its cognate UreG. <i>Biochemical Journal</i> , 2009, 422, 91-100.	1.7	83
92	Zn ²⁺ -linked dimerization of UreG from <i>Helicobacter pylori</i> , a chaperone involved in nickel trafficking and urease activation. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009, 74, 222-239.	1.5	73
93	Superoxide Biosensing with Engineered Cytochrome c. <i>Procedia Chemistry</i> , 2009, 1, 1287-1290.	0.7	2
94	Mapping the Interaction between the Hemophore HasA and Its Outer Membrane Receptor HasR Using CRINEPT ⁺ -TROSY NMR Spectroscopy. <i>Journal of the American Chemical Society</i> , 2009, 131, 1736-1744.	6.6	39
95	Cytochrome c Mutants for Superoxide Biosensors. <i>Analytical Chemistry</i> , 2009, 81, 2976-2984.	3.2	42
96	Deciphering the Structural Role of Histidine 83 for Heme Binding in Hemophore HasA. <i>Journal of Biological Chemistry</i> , 2008, 283, 5960-5970.	1.6	45
97	Metalation of the Amyotrophic Lateral Sclerosis Mutant Glycine 37 to Arginine Superoxide Dismutase (SOD1) Apoprotein Restores Its Structural and Dynamical Properties in Solution to Those of Metalated Wild-Type SOD1. <i>Biochemistry</i> , 2007, 46, 9953-9962.	1.2	25
98	Cytochrome c and Organic Molecules: Solution Structure of the p-Aminophenol Adduct. <i>Biochemistry</i> , 2007, 46, 6232-6238.	1.2	10
99	A method for C ¹³ direct-detection in protonless NMR. <i>Journal of Magnetic Resonance</i> , 2007, 188, 301-310.	1.2	52
100	¹³ C- ¹³ C NOESY spectra of a 480 kDa protein: solution NMR of ferritin. <i>Journal of Biomolecular NMR</i> , 2007, 38, 237-242.	1.6	56
101	Direct-Detected ¹³ C NMR to Investigate the Iron(III) Hemophore HasA. <i>Journal of the American Chemical Society</i> , 2006, 128, 150-158.	6.6	67
102	The Nickel Site of <i>Bacillus pasteurii</i> UreE, a Urease Metallo-Chaperone, As Revealed by Metal-Binding Studies and X-ray Absorption Spectroscopy. <i>Biochemistry</i> , 2006, 45, 6495-6509.	1.2	49
103	An Italian contribution to structural genomics: Understanding metalloproteins. <i>Coordination Chemistry Reviews</i> , 2006, 250, 1419-1450.	9.5	14
104	UreG, a Chaperone in the Urease Assembly Process, Is an Intrinsically Unstructured GTPase That Specifically Binds Zn ²⁺ . <i>Journal of Biological Chemistry</i> , 2005, 280, 4684-4695.	1.6	91
105	Fully Metallated S134N Cu,Zn-Superoxide Dismutase Displays Abnormal Mobility and Intermolecular Contacts in Solution. <i>Journal of Biological Chemistry</i> , 2005, 280, 35815-35821.	1.6	56
106	Conformational variability of matrix metalloproteinases: Beyond a single 3D structure. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 5334-5339.	3.3	143
107	¹ H nuclear magnetic relaxation dispersion of Cu,Zn superoxide dismutase in the native and guanidinium-induced unfolded forms. <i>Biochemical and Biophysical Research Communications</i> , 2005, 328, 633-639.	1.0	10
108	Cytochrome c folding / unfolding: a unifying picture. <i>Journal of Porphyrins and Phthalocyanines</i> , 2004, 08, 238-245.	0.4	4

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109	The reaction of artemisinin with hemin: a further insight into the mechanism. <i>Inorganica Chimica Acta</i> , 2004, 357, 4602-4606.	1.2	4
110	The stability of the cytochrome c scaffold as revealed by NMR spectroscopy. <i>Journal of Inorganic Biochemistry</i> , 2004, 98, 814-823.	1.5	17
111	Insights into Partially Folded or Unfolded States of Metalloproteins from Nuclear Magnetic Resonance. <i>Inorganic Chemistry</i> , 2004, 43, 7945-7952.	1.9	5
112	Cytochrome c and SDS: A Molten Globule Protein with Altered Axial Ligation. <i>Journal of Molecular Biology</i> , 2004, 336, 489-496.	2.0	33
113	The Magnetic Properties of Myoglobin as Studied by NMR Spectroscopy. <i>Chemistry - A European Journal</i> , 2003, 9, 2316-2322.	1.7	45
114	Validation of paramagnetic cross correlation rates for solution structure determination of high spin iron(III) heme proteins. <i>Chemical Physics Letters</i> , 2003, 373, 460-463.	1.2	16
115	NMR studies on partially folded and unfolded states of metalloproteins. <i>Journal of Inorganic Biochemistry</i> , 2003, 96, 31.	1.5	0
116	Structural Model for an Alkaline Form of Ferricytochrome c. <i>Journal of the American Chemical Society</i> , 2003, 125, 2913-2922.	6.6	128
117	Superoxide Dismutase Folding/Unfolding Pathway: Role of the Metal Ions in Modulating Structural and Dynamical Features. <i>Journal of Molecular Biology</i> , 2003, 330, 145-158.	2.0	56
118	¹⁵ N-1H Residual Dipolar Coupling Analysis of Native and Alkaline-K79A <i>Saccharomyces cerevisiae</i> Cytochrome c. <i>Biophysical Journal</i> , 2003, 84, 3917-3923.	0.2	47
119	The metal reductase activity of some multiheme cytochromes c: NMR structural characterization of the reduction of chromium(VI) to chromium(III) by cytochrome c7. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 9750-9754.	3.3	51
120	The Unfolding of Oxidized c-Type Cytochromes: The Instructive Case of <i>Bacillus pasteurii</i> . <i>Journal of Molecular Biology</i> , 2002, 321, 693-701.	2.0	23
121	A quick solution structure determination of the fully oxidized double mutant K9-10A cytochrome c7 from <i>Desulfuromonas acetoxidans</i> and mechanistic implications. <i>Journal of Biomolecular NMR</i> , 2002, 22, 107-122.	1.6	10
122	Evidence that increases of mitochondrial immunoreactive IL-1 β by HIV-1 gp120 implicate in situ cleavage of pro-IL-1 β in the neocortex of rat. <i>Journal of Neurochemistry</i> , 2001, 78, 611-618.	2.1	29
123	A further clue to understanding the mobility of mitochondrial yeast cytochrome c. <i>FEBS Journal</i> , 2001, 268, 4468-4476.	0.2	53
124	Dimethyl propionate ester heme-containing cytochrome b 5: structure and stability. <i>Journal of Biological Inorganic Chemistry</i> , 2001, 6, 490-503.	1.1	14
125	¹⁵ N chemical shift changes in cytochrome c β 5: redox-dependent vs. guanidinium chloride-induced changes. <i>Journal of Biological Inorganic Chemistry</i> , 2000, 5, 761-764.	1.1	15
126	A proton-NMR investigation of the fully reduced cytochrome c7 from <i>Desulfuromonas acetoxidans</i> . Comparison between the reduced and the oxidized forms. <i>FEBS Journal</i> , 1999, 266, 634-643.	0.2	29

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127	Solution structure of reduced horse heart cytochrome c. <i>Journal of Biological Inorganic Chemistry</i> , 1999, 4, 21-31.	1.1	116
128	Three-dimensional solution structures of two DNA dodecamers through full relaxation matrix analysis. , 1999, 37, 564-572.		1
129	Solution structure of paramagnetic metalloproteins. <i>Pure and Applied Chemistry</i> , 1999, 71, 1717-1725.	0.9	11
130	800 MHz ¹ H NMR solution structure refinement of oxidized cytochrome c7 from <i>Desulfuromonas acetoxidans</i> . <i>FEBS Journal</i> , 1998, 256, 261-270.	0.2	36
131	Monitoring the conformational flexibility of cytochrome c at low ionic strength by ¹ H-NMR spectroscopy. <i>FEBS Journal</i> , 1998, 256, 271-278.	0.2	21
132	The Conformational Flexibility of Oxidized Cytochrome c Studied through Its Interaction with NH ₃ and at High Temperatures. <i>European Journal of Inorganic Chemistry</i> , 1998, 1998, 583-591.	1.0	25
133	Water-protein interaction in native and partially unfolded equine cytochrome c. <i>Molecular Physics</i> , 1998, 95, 797-808.	0.8	21
134	Solution Structure of Oxidized <i>Saccharomyces cerevisiae</i> iso-1-cytochrome c. <i>Biochemistry</i> , 1997, 36, 8992-9001.	1.2	125
135	Solution Structure of Oxidized Horse Heart Cytochrome c. <i>Biochemistry</i> , 1997, 36, 9867-9877.	1.2	290
136	A Molecular Dynamics Study in Explicit Water of the Reduced and Oxidized forms of Yeast Iso-1-cytochrome c. Solvation and Dynamic Properties of the two Oxidation States. <i>FEBS Journal</i> , 1997, 249, 716-723.	0.2	22
137	Pseudocontact shifts as constraints for energy minimization and molecular dynamics calculations on solution structures of paramagnetic metalloproteins. , 1997, 29, 68-76.		99
138	Pseudocontact shifts as constraints for energy minimization and molecular dynamics calculations on solution structures of paramagnetic metalloproteins. , 1997, 29, 68.		1
139	Three-Dimensional Solution Structure of <i>Saccharomyces cerevisiae</i> Reduced Iso-1-cytochrome c. <i>Biochemistry</i> , 1996, 35, 13788-13796.	1.2	89
140	NMR characterization and solution structure determination of the oxidized cytochrome c7 from <i>Desulfuromonas acetoxidans</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1996, 93, 14396-14400.	3.3	58
141	The use of pseudocontact shifts to refine solution structures of paramagnetic metalloproteins: Met80Ala cyano-cytochrome c as an example. <i>Journal of Biological Inorganic Chemistry</i> , 1996, 1, 117-126.	1.1	143
142	Can the axial ligand strength be monitored through spectroscopic measurements?. <i>Journal of Biological Inorganic Chemistry</i> , 1996, 1, 364-367.	1.1	21
143	pH, Electrolyte, and Substrate-Linked Variation in Active Site Structure of the Trp51Ala Variant of Cytochrome c Peroxidase. <i>Biochemistry</i> , 1995, 34, 13895-13905.	1.2	30
144	pH-dependent equilibria of yeast Met80Ala-iso-1-cytochrome c probed by NMR spectroscopy: a comparison with the wild-type protein. <i>Chemistry and Biology</i> , 1995, 2, 377-383.	6.2	39

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145	Paramagnetic ¹ H NMR Spectroscopy of the Cyanide Derivative of Met80Ala-iso-1-cytochrome c. <i>Journal of the American Chemical Society</i> , 1995, 117, 8067-8073.	6.6	54
146	Three-Dimensional Solution Structure of the Cyanide Adduct of a Variant of <i>Saccharomyces cerevisiae</i> Iso-1-cytochrome c Containing the Met80Ala Mutation. Identification of Ligand-Residue Interactions in the Distal Heme Cavity. <i>Biochemistry</i> , 1995, 34, 11385-11398.	1.2	65
147	The Hyperfine Coupling. , 1995, , 29-54.		3
148	Active Site Coordination Chemistry of the Cytochrome c Peroxidase Asp235Ala Variant: Spectroscopic and Functional Characterization. <i>Biochemistry</i> , 1994, 33, 7819-7829.	1.2	55
149	Frontiers in 2D NMR of paramagnetic metalloproteins. <i>Applied Magnetic Resonance</i> , 1993, 4, 461-476.	0.6	6
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