

Alessandro Desideri

List of Publications by Year in descending order

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264
papers

6,069
citations

81889

39
h-index

155644

55
g-index

268
all docs

268
docs citations

268
times ranked

6690
citing authors

#	ARTICLE	IF	CITATIONS
1	Dysbiosis of gut microbiota in a selected population of Parkinson's patients. <i>Parkinsonism and Related Disorders</i> , 2019, 65, 124-130.	2.2	144
2	Temperature-Controlled Encapsulation and Release of an Active Enzyme in the Cavity of a Self-Assembled DNA Nanocage. <i>ACS Nano</i> , 2013, 7, 9724-9734.	14.6	132
3	Crystal structure of yeast Cu,Zn superoxide dismutase. <i>Journal of Molecular Biology</i> , 1992, 225, 791-809.	4.2	121
4	Synergistic antitumor effect between vorinostat and topotecan in small cell lung cancer cells is mediated by generation of reactive oxygen species and DNA damage-induced apoptosis. <i>Molecular Cancer Therapeutics</i> , 2009, 8, 3075-3087.	4.1	104
5	Unique structural features of the monomeric Cu,Zn superoxide dismutase from <i>Escherichia coli</i> , revealed by X-ray crystallography. <i>Journal of Molecular Biology</i> , 1997, 274, 408-420.	4.2	83
6	Evolutionary conservativeness of electric field in the Cu,Zn superoxide dismutase active site. <i>Journal of Molecular Biology</i> , 1992, 223, 337-342.	4.2	81
7	Single Mutation in the Linker Domain Confers Protein Flexibility and Camptothecin Resistance to Human Topoisomerase I. <i>Journal of Biological Chemistry</i> , 2003, 278, 43268-43275.	3.4	81
8	Droplet Microfluidics Platform for Highly Sensitive and Quantitative Detection of Malaria-Causing <i>Plasmodium</i> Parasites Based on Enzyme Activity Measurement. <i>ACS Nano</i> , 2012, 6, 10676-10683.	14.6	81
9	Molecular dynamics simulation of solvated azurin: Correlation between surface solvent accessibility and water residence times. <i>Proteins: Structure, Function and Bioinformatics</i> , 2000, 39, 56-67.	2.6	80
10	Receptor-Mediated Entry of Pristine Octahedral DNA Nanocages in Mammalian Cells. <i>ACS Nano</i> , 2016, 10, 5971-5979.	14.6	76
11	An electrochemical multienzymatic biosensor for determination of cholesterol. <i>Bioelectrochemistry</i> , 2001, 54, 17-22.	4.6	73
12	Impaired copper binding by the H46R mutant of human Cu,Zn superoxide dismutase, involved in amyotrophic lateral sclerosis. <i>FEBS Letters</i> , 1994, 356, 314-316.	2.8	71
13	Role of the Dimeric Structure in Cu,Zn Superoxide Dismutase. <i>Journal of Biological Chemistry</i> , 1998, 273, 5655-5661.	3.4	68
14	Oxidovanadium(IV) complexes with chrysin and silibinin: anticancer activity and mechanisms of action in a human colon adenocarcinoma model. <i>Journal of Biological Inorganic Chemistry</i> , 2015, 20, 1175-1191.	2.6	65
15	A Comparative Genomic Analysis Provides Novel Insights Into the Ecological Success of the Monophasic <i>Salmonella</i> Serovar 4,[5],12:i:-. <i>Frontiers in Microbiology</i> , 2018, 9, 715.	3.5	65
16	Evolutionary constraints for dimer formation in prokaryotic Cu,Zn superoxide dismutase 1 Edited by R. Huber. <i>Journal of Molecular Biology</i> , 1999, 285, 283-296.	4.2	63
17	Selective targeting and degradation of doxorubicin-loaded folate-functionalized DNA nanocages. <i>Nanomedicine: Nanotechnology, Biology, and Medicine</i> , 2018, 14, 1181-1190.	3.3	59
18	Anion Size Modulates the Structure of the A State of Cytochrome c. <i>Biochemistry</i> , 2000, 39, 12632-12638.	2.5	58

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19	Evidence of Domain Formation in Cardiolipin~Glycerophospholipid Mixed Monolayers. A Thermodynamic and AFM Study. <i>Journal of Physical Chemistry B</i> , 2005, 109, 15950-15957.	2.6	58
20	Crystal structure solution and refinement of the semisynthetic cobalt-substituted bovine erythrocyte superoxide dismutase at 2.0 Å... resolution. <i>Journal of Molecular Biology</i> , 1992, 226, 227-238.	4.2	53
21	Functional Analysis and Molecular Dynamics Simulation of LOX-1 K167N Polymorphism Reveal Alteration of Receptor Activity. <i>PLoS ONE</i> , 2009, 4, e4648.	2.5	53
22	Targeting Tumor Cells through Chitosan-Folate Modified Microcapsules Loaded with Camptothecin. <i>Bioconjugate Chemistry</i> , 2011, 22, 1066-1072.	3.6	52
23	Tuning the isoelectric point of graphene by electrochemical functionalization. <i>Scientific Reports</i> , 2015, 5, 11794.	3.3	50
24	Thr729 in human topoisomerase I modulates anti-cancer drug resistance by altering protein domain communications as suggested by molecular dynamics simulations. <i>Nucleic Acids Research</i> , 2008, 36, 5645-5651.	14.5	49
25	Nitric oxide binding to ferrous native horse heart cytochrome c and to its carboxymethylated derivative: A spectroscopic and thermodynamic Study. <i>Journal of Inorganic Biochemistry</i> , 1994, 53, 273-280.	3.5	48
26	Modulation of the Catalytic Rate of Cu,Zn Superoxide Dismutase in Single and Double Mutants of Conserved Positively and Negatively Charged Residues. <i>Biochemistry</i> , 1995, 34, 6043-6049.	2.5	48
27	Functional and crystallographic characterization of <i>Salmonella typhimurium</i> Cu,Zn superoxide dismutase coded by the sodCI virulence gene 1 Edited by R. Huber. <i>Journal of Molecular Biology</i> , 2000, 302, 465-478.	4.2	47
28	Effect on DNA relaxation of the single Thr718Ala mutation in human topoisomerase I: a functional and molecular dynamics study. <i>Nucleic Acids Research</i> , 2005, 33, 3339-3350.	14.5	47
29	Structure of Nanoscale Truncated Octahedral DNA Cages: Variation of Single-Stranded Linker Regions and Influence on Assembly Yields. <i>ACS Nano</i> , 2010, 4, 1367-1376.	14.6	47
30	Protein concerted motions in the DNA-human topoisomerase I complex. <i>Nucleic Acids Research</i> , 2003, 31, 1525-1535.	14.5	44
31	Interaction between natural compounds and human topoisomerase I. <i>Biological Chemistry</i> , 2012, 393, 1327-1340.	2.5	44
32	New Hints on the pH-Driven Tautomeric Equilibria of the Topotecan Anticancer Drug in Aqueous Solutions from an Integrated Spectroscopic and Quantum-Mechanical Approach. <i>Journal of the American Chemical Society</i> , 2005, 127, 15429-15436.	13.7	43
33	Gimatecan and other camptothecin derivatives poison <i>Leishmania</i> DNA-topoisomerase IB leading to a strong leishmanicidal effect. <i>Biochemical Pharmacology</i> , 2013, 85, 1433-1440.	4.4	43
34	Real-Time Label-Free Direct Electronic Monitoring of Topoisomerase Enzyme Binding Kinetics on Graphene. <i>ACS Nano</i> , 2015, 9, 11166-11176.	14.6	43
35	Azide, cyanide, fluoride, imidazole and pyridine binding to ferric and ferrous native horse heart cytochrome c and to its carboxymethylated derivative: A comparative study. <i>Journal of Inorganic Biochemistry</i> , 1996, 62, 213-222.	3.5	42
36	Structure of calmodulin complexed with an olfactory CNG channel fragment and role of the central linker: Residual dipolar coupling to evaluate calmodulin binding modes outside the kinase family. <i>Journal of Biomolecular NMR</i> , 2005, 31, 185-199.	2.8	42

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37	UV-Vis Spectra of the Anticancer Camptothecin Family Drugs in Aqueous Solution: Specific Spectroscopic Signatures Unraveled by a Combined Computational and Experimental Study. <i>Journal of Physical Chemistry B</i> , 2009, 113, 5369-5375.	2.6	42
38	Cellular uptake of covalent and non-covalent DNA nanostructures with different sizes and geometries. <i>Nanoscale</i> , 2019, 11, 10808-10818.	5.6	42
39	Effects of dutasteride on the expression of genes related to androgen metabolism and related pathway in human prostate cancer cell lines. <i>Investigational New Drugs</i> , 2007, 25, 491-497.	2.6	41
40	Real-time detection of TDP1 activity using a fluorophore-quencher coupled DNA-biosensor. <i>Biosensors and Bioelectronics</i> , 2013, 48, 230-237.	10.1	41
41	Effect of oxindolimine copper and zinc complexes on human topoisomerase I activity. <i>Metallomics</i> , 2014, 6, 117-125.	2.4	41
42	Structural Evolution and Dynamics of the p53 Proteins. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2017, 7, a028308.	6.2	41
43	Valproic Acid Induces Neuroendocrine Differentiation and UGT2B7 Up-Regulation in Human Prostate Carcinoma Cell Line. <i>Drug Metabolism and Disposition</i> , 2007, 35, 968-972.	3.3	40
44	Erybraedin C, a natural compound from the plant <i>Bituminaria bituminosa</i> , inhibits both the cleavage and religation activities of human topoisomerase I. <i>Biochemical Journal</i> , 2010, 425, 531-539.	3.7	40
45	Detection of quasispecies variants predicted to use CXCR4 by ultra-deep pyrosequencing during early HIV infection. <i>Aids</i> , 2011, 25, 611-617.	2.2	40
46	The p53 tetramer shows an induced-fit interaction of the C-terminal domain with the DNA-binding domain. <i>Oncogene</i> , 2016, 35, 3272-3281.	5.9	40
47	Effect of Low-Protein Diet and Inulin on Microbiota and Clinical Parameters in Patients with Chronic Kidney Disease. <i>Nutrients</i> , 2019, 11, 3006.	4.1	40
48	Quercetin pentaacetate inhibits in vitro human respiratory syncytial virus adhesion. <i>Virus Research</i> , 2020, 276, 197805.	2.2	40
49	Removal of non-blue copper from ascorbate oxidase. <i>FEBS Letters</i> , 1979, 100, 318-320.	2.8	37
50	Magnetic susceptibility studies of the native cupro-zinc superoxide dismutase and its cobalt-substituted derivatives. Antiferromagnetic coupling in the imidazolate-bridged copper(II)-cobalt(II) pair. <i>Journal of the American Chemical Society</i> , 1986, 108, 300-302.	13.7	37
51	Molecular mechanism of statin-mediated LOX-1 inhibition. <i>Cell Cycle</i> , 2015, 14, 1583-1595.	2.6	36
52	Crystal structure of the cyanide-inhibited <i>Xenopus laevis</i> Cu,Zn superoxide dismutase at 98 K. <i>FEBS Letters</i> , 1994, 349, 93-98.	2.8	35
53	Role of the electrostatic loop charged residues in Cu, Zn superoxide dismutase. <i>Protein Science</i> , 1998, 7, 2354-2358.	7.6	35
54	Role of the Tertiary and Quaternary Structures in the Stability of Dimeric Copper,Zinc Superoxide Dismutases. <i>Archives of Biochemistry and Biophysics</i> , 2000, 377, 215-218.	3.0	35

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55	Palladium(II) complexes with thiosemicarbazones derived from pyrene as topoisomerase IB inhibitors. <i>Dalton Transactions</i> , 2019, 48, 16509-16517.	3.3	34
56	Reconstitution of stellacyanin as a case of direct Cu(I) transfer between yeast copper thionein and a blue copper apoprotein. <i>FEBS Letters</i> , 1983, 152, 94-96.	2.8	32
57	Crystallographic Study of Azide-inhibited Bovine Cu,Zn Superoxide Dismutase. <i>Journal of Molecular Biology</i> , 1994, 240, 179-183.	4.2	32
58	Structural and Dynamical Effects Induced by the Anticancer Drug Topotecan on the Human Topoisomerase I-DNA Complex. <i>PLoS ONE</i> , 2010, 5, e10934.	2.5	32
59	Prokaryotic Cu,Zn superoxide dismutases. <i>Biochemical Society Transactions</i> , 2003, 31, 1322-1325.	3.4	31
60	Inhibition of glutathione transferase γ from human placenta by 1-chloro-2,4-dinitrobenzene occurs because of covalent reaction with cysteine 47. <i>Archives of Biochemistry and Biophysics</i> , 1992, 297, 119-122.	3.0	30
61	Spectroscopic Characterization of Recombinant Cu,Zn Superoxide Dismutase from <i>Photobacterium leiognathi</i> Expressed in <i>Escherichia coli</i> : Evidence for a Novel Catalytic Copper Binding Site. <i>Biochemistry</i> , 1997, 36, 7109-7113.	2.5	30
62	Cu,Zn Superoxide Dismutase from <i>Photobacterium leiognathii</i> is an Hyperefficient Enzyme. <i>Biochemistry</i> , 1998, 37, 12287-12292.	2.5	30
63	Conjugated eicosapentaenoic acid inhibits human topoisomerase IB with a mechanism different from camptothecin. <i>Archives of Biochemistry and Biophysics</i> , 2009, 486, 103-110.	3.0	30
64	Entry, fate and degradation of DNA nanocages in mammalian cells: a matter of receptors. <i>Nanoscale</i> , 2018, 10, 12078-12086.	5.6	30
65	A Novel Co(II) binding site in copper-free superoxide dismutase. <i>FEBS Letters</i> , 1979, 106, 142-144.	2.8	29
66	Structural dynamics of the mitochondrial ADP/ATP carrier revealed by molecular dynamics simulation studies. <i>Proteins: Structure, Function and Bioinformatics</i> , 2006, 65, 681-691.	2.6	29
67	Evidence of the crucial role of the linker domain on the catalytic activity of human topoisomerase I by experimental and simulative characterization of the Lys681Ala mutant. <i>Nucleic Acids Research</i> , 2009, 37, 6849-6858.	14.5	29
68	Design of a novel LOX-1 receptor antagonist mimicking the natural substrate. <i>Biochemical and Biophysical Research Communications</i> , 2013, 438, 340-345.	2.1	29
69	Molecular mechanism of the camptothecin resistance of Glu710Gly topoisomerase IB mutant analyzed in vitro and in silico. <i>Molecular Cancer</i> , 2013, 12, 100.	19.2	29
70	Simulations of DNA topoisomerase IB bound to supercoiled DNA reveal changes in the flexibility pattern of the enzyme and a secondary protein-DNA binding site. <i>Nucleic Acids Research</i> , 2014, 42, 9304-9312.	14.5	29
71	Iron entry route in horse spleen apoferritin. <i>FEBS Letters</i> , 1991, 287, 10-14.	2.8	28
72	Simulation of superoxide-superoxide dismutase association rate for six natural variants. Comparison with the experimental catalytic rate. <i>The Journal of Physical Chemistry</i> , 1994, 98, 10554-10557.	2.9	28

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73	Fourier Transform Infrared Analysis of the Interaction of Azide with the Active Site of Oxidized and Reduced Bovine Cu,Zn Superoxide Dismutase. <i>Biochemistry</i> , 1998, 37, 4459-4464.	2.5	28
74	A Novel Heme Protein, the Cu,Zn-Superoxide Dismutase from <i>Haemophilus ducreyi</i> . <i>Journal of Biological Chemistry</i> , 2001, 276, 30326-30334.	3.4	28
75	DNA hairpins promote temperature controlled cargo encapsulation in a truncated octahedral nanocage structure family. <i>Nanoscale</i> , 2016, 8, 13333-13341.	5.6	28
76	Inhibition of human DNA topoisomerase IB by nonmutagenic ruthenium(II)-based compounds with antitumoral activity. <i>Metallomics</i> , 2016, 8, 179-192.	2.4	28
77	Effective binding force calculation in a dimeric protein by molecular dynamics simulation. <i>Journal of Chemical Physics</i> , 2002, 116, 6329-6338.	3.0	27
78	Molecular Dynamics Simulation of the RNA Complex of a Double-Stranded RNA-Binding Domain Reveals Dynamic Features of the Intermolecular Interface and Its Hydration. <i>Biophysical Journal</i> , 2002, 83, 3542-3552.	0.5	27
79	Inhibition of human DNA topoisomerase IB by a Cyclometalated Gold III compound: Analysis on the different steps of the enzyme catalytic cycle. <i>Archives of Biochemistry and Biophysics</i> , 2011, 516, 108-112.	3.0	27
80	Molecular dynamics of the full-length p53 monomer. <i>Cell Cycle</i> , 2013, 12, 3098-3108.	2.6	27
81	Reptile Heme Protein Structure: X-ray Crystallographic Study of the Aquo-met and Cyano-met Derivatives of the Loggerhead Sea Turtle (<i>Caretta caretta</i>) Myoglobin at 2.0 Å... Resolution. <i>Journal of Molecular Biology</i> , 1995, 247, 459-465.	4.2	26
82	Conserved Enzyme-Substrate Electrostatic Attraction in Prokaryotic Cu,Zn Superoxide Dismutases. <i>Biochemical and Biophysical Research Communications</i> , 1998, 244, 908-911.	2.1	26
83	Structure and Hydration of the DNA-Human Topoisomerase I Covalent Complex. <i>Biophysical Journal</i> , 2001, 81, 490-500.	0.5	26
84	Structure and Stability of the Insulin Dimer Investigated by Molecular Dynamics Simulation. <i>Journal of Biomolecular Structure and Dynamics</i> , 2001, 18, 761-772.	3.5	26
85	Role of the protein in the DNA sequence specificity of the cleavage site stabilized by the camptothecin topoisomerase IB inhibitor: a metadynamics study. <i>Nucleic Acids Research</i> , 2013, 41, 9977-9986.	14.5	26
86	Human topoisomerase inhibition and DNA/BSA binding of Ru(II)-SCAR complexes as potential anticancer candidates for oral application. <i>BioMetals</i> , 2017, 30, 321-334.	4.1	26
87	.beta.-Bungarotoxin-mediated liposome fusion: spectroscopic characterization by fluorescence and ESR. <i>Biochemistry</i> , 1990, 29, 9644-9651.	2.5	25
88	Reduced sensitivity of O ₂ transport to allosteric effectors and temperature in loggerhead sea turtle hemoglobin: functional and spectroscopic study. <i>BBA - Proteins and Proteomics</i> , 1992, 1159, 129-133.	2.1	25
89	Role of the Linker Domain and the 203-214 N-Terminal Residues in the Human Topoisomerase I DNA Complex Dynamics. <i>Biophysical Journal</i> , 2004, 87, 4087-4097.	0.5	25
90	Molecular dynamics simulation of human LOX-1 provides an explanation for the lack of OxLDL binding to the Trp150Ala mutant. <i>BMC Structural Biology</i> , 2007, 7, 73.	2.3	25

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91	Deciphering the Structural Properties That Confer Stability to a DNA Nanocage. <i>ACS Nano</i> , 2009, 3, 1813-1822.	14.6	25
92	Role of 13-(di)phenylalkyl berberine derivatives in the modulation of the activity of human topoisomerase IB. <i>International Journal of Biological Macromolecules</i> , 2015, 77, 68-75.	7.5	25
93	Evidence of His61 Imidazolate Bridge Rupture in Reduced Crystalline Cu,Zn Superoxide Dismutase. <i>Biochemical and Biophysical Research Communications</i> , 1997, 241, 119-121.	2.1	24
94	Efficacy of a Binuclear Cyclopalladated Compound Therapy for Cutaneous Leishmaniasis in the Murine Model of Infection with <i>Leishmania amazonensis</i> and Its Inhibitory Effect on Topoisomerase IB. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	3.2	24
95	pH-induced cleavage of the proximal histidine to iron bond in the nitric oxide derivative of ferrous monomeric hemosproteins and of the α -chelated α ™ protoheme model compound. <i>BBA - Proteins and Proteomics</i> , 1985, 829, 299-302.	2.1	23
96	Cooperative effect of inositol hexakisphosphate, bezafibrate, and clofibrac acid on the spectroscopic properties of the nitric oxide derivative of ferrous human hemoglobin. <i>Journal of Inorganic Biochemistry</i> , 1993, 50, 263-272.	3.5	23
97	Mutation of Lys-120 and Lys-134 drastically reduces the catalytic rate of Cu,Zn superoxide dismutase. <i>FEBS Letters</i> , 1994, 352, 76-78.	2.8	23
98	Membrane-perturbing activity of Viperidae myotoxins: an electrostatic surface potential approach to a puzzling problem. , 2000, 13, 14-19.		23
99	High-Density ZnO Nanowires as a Reversible Myogenic Differentiation Switch. <i>ACS Applied Materials & Interfaces</i> , 2018, 10, 14097-14107.	8.0	23
100	Comparative stability studies on the iron and manganese forms of the cambialistic superoxide dismutase from <i>Propionibacterium shermanii</i> . <i>FEBS Letters</i> , 1997, 414, 122-124.	2.8	22
101	X-ray Absorption Investigation of a Unique Protein Domain Able To Bind both Copper(I) and Copper(II) at Adjacent Sites of the N-Terminus of <i>Haemophilus ducreyi</i> Cu,Zn Superoxide Dismutase. <i>Biochemistry</i> , 2005, 44, 13144-13150.	2.5	22
102	Human topoisomerase IB is a target of a thiosemicarbazone copper(II) complex. <i>Archives of Biochemistry and Biophysics</i> , 2016, 606, 34-40.	3.0	22
103	Simulative and Experimental Characterization of a pH-Dependent Clamp-like DNA Triple-Helix Nanoswitch. <i>Journal of the American Chemical Society</i> , 2017, 139, 5321-5329.	13.7	22
104	Can Gut Microbiota Be a Good Predictor for Parkinson's Disease? A Machine Learning Approach. <i>Brain Sciences</i> , 2020, 10, 242.	2.3	22
105	Combined and selective miR-21 silencing and doxorubicin delivery in cancer cells using tailored DNA nanostructures. <i>Cell Death and Disease</i> , 2021, 12, 7.	6.3	22
106	Reaction of N,N-diethyldithiocarbamate and other bidentate ligands with Zn, Co and Cu bovine carbonic anhydrases. <i>BBA - Proteins and Proteomics</i> , 1983, 746, 168-175.	2.1	21
107	Formation of a molten-globule-like state of cytochrome c induced by high concentrations of glycerol. <i>Biochimie</i> , 1999, 81, 745-750.	2.6	21
108	The different cleavage DNA sequence specificity explains the camptothecin resistance of the human topoisomerase I Glu418Lys mutant. <i>Nucleic Acids Research</i> , 2006, 34, 5093-5100.	14.5	21

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109	The open state of human topoisomerase I as probed by molecular dynamics simulation. <i>Nucleic Acids Research</i> , 2007, 35, 3032-3038.	14.5	21
110	The sterile alpha-motif (SAM) domain of p63 binds in vitro monoasialoganglioside (GM1) micelles. <i>Biochemical Pharmacology</i> , 2011, 82, 1262-1268.	4.4	21
111	AS1411 Aptamer Linked to DNA Nanostructures Diverts Its Traffic Inside Cancer Cells and Improves Its Therapeutic Efficacy. <i>Pharmaceutics</i> , 2021, 13, 1671.	4.5	21
112	Intestinal Taxa Abundance and Diversity in Inflammatory Bowel Disease Patients: An Analysis including Covariates and Confounders. <i>Nutrients</i> , 2022, 14, 260.	4.1	21
113	An X-ray absorption study of the reconstitution process of bovine Cu,Zn superoxide dismutase by Cu(I)-glutathione complex. <i>FEBS Letters</i> , 1993, 322, 165-167.	2.8	20
114	Low-Temperature Optical Spectroscopy of Native and Azide-Reacted Bovine Cu,Zn Superoxide Dismutase. A Structural Dynamics Study. <i>Biochemistry</i> , 1994, 33, 15103-15109.	2.5	20
115	Flexibility in monomeric Cu,Zn superoxide dismutase detected by limited proteolysis and molecular dynamics simulation. <i>Proteins: Structure, Function and Bioinformatics</i> , 2002, 47, 513-520.	2.6	20
116	Cost-effectiveness analysis of noninvasive strategies to evaluate patients with chest pain. <i>Journal of the American Society of Echocardiography</i> , 2003, 16, 1287-1291.	2.8	20
117	Hif1 α down-regulation is associated with transposition of great arteries in mice treated with a retinoic acid antagonist. <i>BMC Genomics</i> , 2010, 11, 497.	2.8	20
118	A Simple and Fast Semiautomatic Procedure for the Atomistic Modeling of Complex DNA Polyhedra. <i>Journal of Chemical Information and Modeling</i> , 2016, 56, 941-949.	5.4	20
119	Formate as an NMR probe of anion binding to copper-zinc and copper-cobalt bovine erythrocyte superoxide dismutase. <i>Biochemistry</i> , 1992, 31, 12410-12415.	2.5	19
120	Substrate-induced conformational changes of the mitochondrial oxoglutarate carrier: a spectroscopic and molecular modelling study. <i>Molecular Membrane Biology</i> , 2005, 22, 443-452.	2.0	19
121	NMR Structure of the p63 SAM Domain and Dynamical Properties of G534V and T537P Pathological Mutants, Identified in the AEC Syndrome. <i>Cell Biochemistry and Biophysics</i> , 2006, 44, 475-489.	1.8	19
122	Importance of V3 Loop Flexibility and Net Charge in the Context of Co-Receptor Recognition. A Molecular Dynamics Study on HIV gp120. <i>Journal of Biomolecular Structure and Dynamics</i> , 2012, 29, 879-891.	3.5	19
123	Metal complexes of 3-(4-bromophenyl)-1-pyridin-2-ylprop-2-en-1-one thiosemicarbazone: cytotoxic activity and investigation on the mode of action of the gold(III) complex. <i>BioMetals</i> , 2016, 29, 515-526.	4.1	19
124	Ru/Fe bimetallic complexes: Synthesis, characterization, cytotoxicity and study of their interactions with DNA/HSA and human topoisomerase IB. <i>Archives of Biochemistry and Biophysics</i> , 2017, 636, 28-41.	3.0	19
125	Development of Derivatives of 3, 3- α -Diindolylmethane as Potent <i>Leishmania donovani</i> Bi-Subunit Topoisomerase IB Poisons. <i>PLoS ONE</i> , 2011, 6, e28493.	2.5	19
126	Topoisomerase 1B as a Target Against Leishmaniasis. <i>Mini-Reviews in Medicinal Chemistry</i> , 2015, 15, 203-210.	2.4	19

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127	A Natural Anticancer Agent Thapsigargin Targets Human Topoisomerase II α . <i>Anti-Cancer Agents in Medicinal Chemistry</i> , 2013, 13, 356-363.	1.7	19
128	Static and dynamic water molecules in Cu,Zn superoxide dismutase. <i>Proteins: Structure, Function and Bioinformatics</i> , 2003, 51, 607-615.	2.6	18
129	Active-site Copper and Zinc Ions Modulate the Quaternary Structure of Prokaryotic Cu,Zn Superoxide Dismutase. <i>Journal of Molecular Biology</i> , 2003, 326, 1351-1360.	4.2	18
130	Non-mutagenic Ru(II) complexes: cytotoxicity, topoisomerase II α inhibition, DNA and HSA binding. <i>Dalton Transactions</i> , 2019, 48, 14885-14897.	3.3	18
131	Electron paramagnetic resonance properties of liver fluke (<i>Dicrocoelium dendriticum</i>) nitrosyl hemoglobin. <i>FEBS Letters</i> , 1984, 166, 378-380.	2.8	17
132	Toward the Engineering of a Super Efficient Enzyme. <i>Biochemical and Biophysical Research Communications</i> , 1999, 256, 425-428.	2.1	17
133	Single mutation at the intersubunit interface confers extra efficiency to Cu,Zn superoxide dismutase. <i>FEBS Letters</i> , 2000, 483, 17-20.	2.8	17
134	Simulative and experimental investigation on the cleavage site that generates the soluble human LOX-1. <i>Archives of Biochemistry and Biophysics</i> , 2013, 540, 9-18.	3.0	17
135	Design, selection and optimization of an anti-TRAIL-R2/anti-CD3 bispecific antibody able to educate T cells to recognize and destroy cancer cells. <i>MAbs</i> , 2018, 10, 1084-1097.	5.2	17
136	Diving behaviour and haemoglobin function: the primary structure of the α - and β -chains of the sea turtle (<i>Caretta caretta</i>) and its functional implications. <i>Biochemical Journal</i> , 1996, 316, 959-965.	3.7	16
137	The Heme-Containing N-Fragment (Residues 1-56) of Cytochrome c Is a Bis-histidine Functional System. <i>Archives of Biochemistry and Biophysics</i> , 2000, 379, 331-336.	3.0	16
138	Single Mutation Induces a Metal-Dependent Subunit Association in Dimeric Cu,Zn Superoxide Dismutase. <i>Biochemical and Biophysical Research Communications</i> , 2000, 272, 81-83.	2.1	16
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