

Pietro Pucci

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5212575/publications.pdf>

Version: 2024-02-01

244
papers

7,949
citations

50244

46
h-index

91828

69
g-index

251
all docs

251
docs citations

251
times ranked

12050
citing authors

#	ARTICLE	IF	CITATIONS
1	From untargeted metabolomics to the multiple reaction monitoring-based quantification of polyphenols in chocolates from different geographical areas. <i>Journal of Mass Spectrometry</i> , 2021, 56, e4651.	0.7	15
2	Tyrosine Phosphorylation Modulates Peroxiredoxin-2 Activity in Normal and Diseased Red Cells. <i>Antioxidants</i> , 2021, 10, 206.	2.2	4
3	An integrated transcriptomic and proteomic approach to identify the main <i>Torymus sinensis</i> venom components. <i>Scientific Reports</i> , 2021, 11, 5032.	1.6	18
4	Hemoglobin Yamagata [¹³² (H10)Lys→Asn; (HBB: c.399A>T)]: a mosaic to be put together. <i>Clinical Chemistry and Laboratory Medicine</i> , 2021, 59, 1670-1679.	1.4	1
5	Identification of SARS-CoV-2 Proteins from Nasopharyngeal Swabs Probed by Multiple Reaction Monitoring Tandem Mass Spectrometry. <i>ACS Omega</i> , 2021, 6, 34945-34953.	1.6	10
6	A Hyperthermoactive-Cas9 Editing Tool Reveals the Role of a Unique Arsenite Methyltransferase in the Arsenic Resistance System of <i>Thermus thermophilus</i> HB27. <i>MBio</i> , 2021, 12, e0281321.	1.8	8
7	TRIM8 interacts with KIF11 and KIFC1 and controls bipolar spindle formation and chromosomal stability. <i>Cancer Letters</i> , 2020, 473, 98-106.	3.2	16
8	Fyn specifically Regulates the activity of red cell glucose-6-phosphate-dehydrogenase. <i>Redox Biology</i> , 2020, 36, 101639.	3.9	14
9	Heat-Induced Brain Vitrification from the Vesuvius Eruption in <i>c.e.</i> 79. <i>New England Journal of Medicine</i> , 2020, 382, 383-384.	13.9	12
10	Venomomics of the ectoparasitoid wasp <i>Bracon nigricans</i> . <i>BMC Genomics</i> , 2020, 21, 34.	1.2	20
11	The interaction between the F55 virus-encoded transcription regulator and the RadA host recombinase reveals a common strategy in Archaea and Bacteria to sense the UV-induced damage to the host DNA. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2020, 1863, 194493.	0.9	15
12	The TRAPP complex mediates secretion arrest induced by stress granule assembly. <i>EMBO Journal</i> , 2019, 38, e101704.	3.5	20
13	Lanthionine and Other Relevant Sulfur Amino Acid Metabolites: Detection of Prospective Uremic Toxins in Serum by Multiple Reaction Monitoring Tandem Mass Spectrometry. <i>Methods in Molecular Biology</i> , 2019, 2007, 9-17.	0.4	5
14	Ultra-Rapid Glutathionylation of Ribonuclease: Is this the Real Incipit of its Oxidative Folding?. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5440.	1.8	6
15	Hb Vanvitelli: A new unstable β -globin chain variant causes undiagnosed chronic haemolytic anaemia when co-inherited with deletion β^0 . <i>Clinical Biochemistry</i> , 2019, 74, 80-85.	0.8	1
16	Evolution of an insect immune barrier through horizontal gene transfer mediated by a parasitic wasp. <i>PLoS Genetics</i> , 2019, 15, e1007998.	1.5	32
17	The complex CBX7-PRMT1 has a critical role in regulating E-cadherin gene expression and cell migration. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2019, 1862, 509-521.	0.9	18
18	Genome-wide mapping of 8-oxo-7,8-dihydro-2'-deoxyguanosine reveals accumulation of oxidatively-generated damage at DNA replication origins within transcribed long genes of mammalian cells. <i>Nucleic Acids Research</i> , 2019, 47, 221-236.	6.5	94

#	ARTICLE	IF	CITATIONS
19	TRIM8-driven transcriptomic profile of neural stem cells identified glioma-related nodal genes and pathways. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2019, 1863, 491-501.	1.1	22
20	Identification of proteinaceous binders in paintings: A targeted proteomic approach for cultural heritage. <i>Microchemical Journal</i> , 2019, 144, 319-328.	2.3	18
21	Fyn Specifically Regulates the Activity of Red Cell Glucose-6-Phosphate-Dehydrogenase. <i>Blood</i> , 2019, 134, 3527-3527.	0.6	0
22	S-glutathionylation exerts opposing roles in the regulation of STAT1 and STAT3 signaling in reactive microglia. <i>Free Radical Biology and Medicine</i> , 2018, 117, 191-201.	1.3	29
23	New insights on the functional role of URG7 in the cellular response to ER stress. <i>Biology of the Cell</i> , 2018, 110, 147-158.	0.7	12
24	Multiple Reaction Monitoring Tandem Mass Spectrometry Approach for the Identification of Biological Fluids at Crime Scene Investigations. <i>Analytical Chemistry</i> , 2018, 90, 5627-5636.	3.2	21
25	A hypothesis of sudden body fluid vaporization in the 79 AD victims of Vesuvius. <i>PLoS ONE</i> , 2018, 13, e0203210.	1.1	17
26	New perspectives in cancer: Modulation of lipid metabolism and inflammation resolution. <i>Pharmacological Research</i> , 2018, 128, 80-87.	3.1	31
27	The centrosomal OFD1 protein interacts with the translation machinery and regulates the synthesis of specific targets. <i>Scientific Reports</i> , 2017, 7, 1224.	1.6	36
28	Formyl peptide receptor 1 suppresses gastric cancer angiogenesis and growth by exploiting inflammation resolution pathways. <i>Oncolmmunology</i> , 2017, 6, e1293213.	2.1	43
29	The multifunctional polydnavirus TnBVANK1 protein: impact on host apoptotic pathway. <i>Scientific Reports</i> , 2017, 7, 11775.	1.6	23
30	PRUNE is crucial for normal brain development and mutated in microcephaly with neurodevelopmental impairment. <i>Brain</i> , 2017, 140, 940-952.	3.7	62
31	Quantitative determination of free D-Asp, L-Asp and N-methyl-D-aspartate in mouse brain tissues by chiral separation and Multiple Reaction Monitoring tandem mass spectrometry. <i>PLoS ONE</i> , 2017, 12, e0179748.	1.1	13
32	A Novel Pathogenic BRCA1 Splicing Variant Produces Partial Intron Retention in the Mature Messenger RNA. <i>International Journal of Molecular Sciences</i> , 2016, 17, 2145.	1.8	9
33	Identification of major Toxoneuron nigriceps venom proteins using an integrated transcriptomic/proteomic approach. <i>Insect Biochemistry and Molecular Biology</i> , 2016, 76, 49-61.	1.2	44
34	Proteome analysis of human amniotic mesenchymal stem cells (hA-MSCs) reveals impaired antioxidant ability, cytoskeleton and metabolic functionality in maternal obesity. <i>Scientific Reports</i> , 2016, 6, 25270.	1.6	27
35	Divergent behavior of hydrogen sulfide pools and of the sulfur metabolite lanthionine, a novel uremic toxin, in dialysis patients. <i>Biochimie</i> , 2016, 126, 97-107.	1.3	37
36	Intermolecular disulfide bond influences unphosphorylated STAT3 dimerization and function. <i>Biochemical Journal</i> , 2016, 473, 3205-3219.	1.7	24

#	ARTICLE	IF	CITATIONS
37	Spectroscopic investigation of auranofin binding to zinc finger HIV-2 nucleocapsid peptides. <i>Inorganica Chimica Acta</i> , 2016, 453, 330-338.	1.2	7
38	Regulating levels of the neuromodulator <i>serine</i> in human brain: structural insight into pLG72 and <i>amino acid oxidase</i> interaction. <i>FEBS Journal</i> , 2016, 283, 3353-3370.	2.2	15
39	A new hexapeptide from the leader peptide of rMnSOD enters cells through the oestrogen receptor to deliver therapeutic molecules. <i>Scientific Reports</i> , 2016, 6, 18691.	1.6	7
40	Identification of p38 MAPK and JNK as new targets for correction of Wilson disease-causing ATP7B mutants. <i>Hepatology</i> , 2016, 63, 1842-1859.	3.6	42
41	Proteomic strategies for cultural heritage: From bones to paintings. <i>Microchemical Journal</i> , 2016, 126, 341-348.	2.3	60
42	Protein conformational perturbations in hereditary amyloidosis: Differential impact of single point mutations in ApoA1 amyloidogenic variants. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2016, 1860, 434-444.	1.1	23
43	β -Thalassemia Associated with Hb Instability: A Tale of Two Features. The Case of Hb Rogliano or β 108(G15)Thr \rightarrow Asn and Hb Policoro or β 124(H7)Ser \rightarrow Pro.. <i>PLoS ONE</i> , 2015, 10, e0115738.	1.1	9
44	Deglycosylation Step to Improve the Identification of Egg Proteins in Art Samples. <i>Analytical Chemistry</i> , 2015, 87, 10178-10182.	3.2	22
45	Proteolytic cleavage of Ser52Pro variant transthyretin triggers its amyloid fibrillogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 1539-1544.	3.3	91
46	<i>Xanthomonas campestris</i> lipooligosaccharides trigger innate immunity and oxidative burst in <i>Arabidopsis</i> . <i>Plant Physiology and Biochemistry</i> , 2014, 85, 51-62.	2.8	12
47	Phosphorylation-Regulated Degradation of the Tumor-Suppressor Form of PED by Chaperone-Mediated Autophagy in Lung Cancer Cells. <i>Journal of Cellular Physiology</i> , 2014, 229, 1359-1368.	2.0	42
48	The role of copper(II) in the aggregation of human amylin. <i>Metallomics</i> , 2014, 6, 1841-1852.	1.0	51
49	S-Glutathionylation at Cys328 and Cys542 Impairs STAT3 Phosphorylation. <i>ACS Chemical Biology</i> , 2014, 9, 1885-1893.	1.6	66
50	Wilson Disease Protein ATP7B Utilizes Lysosomal Exocytosis to Maintain Copper Homeostasis. <i>Developmental Cell</i> , 2014, 29, 686-700.	3.1	203
51	HDAC6 mediates the acetylation of TRIM50. <i>Cellular Signalling</i> , 2014, 26, 363-369.	1.7	17
52	Innate immunity probed by lipopolysaccharides affinity strategy and proteomics. <i>Analytical and Bioanalytical Chemistry</i> , 2013, 405, 775-784.	1.9	9
53	A new anti-infective strategy to reduce the spreading of antibiotic resistance by the action on adhesion-mediated virulence factors in <i>Staphylococcus aureus</i> . <i>Microbial Pathogenesis</i> , 2013, 63, 44-53.	1.3	39
54	Vesicular and non-vesicular transport feed distinct glycosylation pathways in the Golgi. <i>Nature</i> , 2013, 501, 116-120.	13.7	136

#	ARTICLE	IF	CITATIONS
55	Molecular and Functional Analysis of the Large 5' Promoter Region of CFTR Gene Revealed Pathogenic Mutations in CF and CFTR-Related Disorders. <i>Journal of Molecular Diagnostics</i> , 2013, 15, 331-340.	1.2	27
56	A simple and reliable methodology to detect egg white in art samples. <i>Journal of Biosciences</i> , 2013, 38, 397-408.	0.5	29
57	The side chain of glutamine 13 is the acyl-donor amino acid modified by type 2 transglutaminase in subunit T of the native rabbit skeletal muscle troponin complex. <i>Amino Acids</i> , 2013, 44, 227-234.	1.2	1
58	Structural characterization and biological properties of human gastrokine 1. <i>Molecular BioSystems</i> , 2013, 9, 412.	2.9	18
59	Role of GALNT2 in the modulation of ENPP1 expression, and insulin signaling and action. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2013, 1833, 1388-1395.	1.9	23
60	Comparison of the action of different proteases on virulence properties related to the staphylococcal surface. <i>Journal of Applied Microbiology</i> , 2013, 114, 266-277.	1.4	42
61	The Mitochondrial Italian Human Proteome Project Initiative (mt-HPP). <i>Molecular BioSystems</i> , 2013, 9, 1984-92.	2.9	10
62	W-F Substitutions in Apomyoglobin Increase the Local Flexibility of the N-terminal Region Causing Amyloid Aggregation: A H/D Exchange Study. <i>Protein and Peptide Letters</i> , 2013, 20, 898-904.	0.4	6
63	A complex of α_6 integrin and E-cadherin drives liver metastasis of colorectal cancer cells through hepatic angiopoietin-like 6. <i>EMBO Molecular Medicine</i> , 2012, 4, 1156-1175.	3.3	44
64	Functional amyloids in insect immune response. <i>Insect Biochemistry and Molecular Biology</i> , 2012, 42, 203-211.	1.2	42
65	The E3-Ubiquitin Ligase TRIM50 Interacts with HDAC6 and p62, and Promotes the Sequestration and Clearance of Ubiquitinated Proteins into the Aggresome. <i>PLoS ONE</i> , 2012, 7, e40440.	1.1	76
66	Resolution of the effects induced by W7F substitutions on the conformation and dynamics of the amyloid-forming apomyoglobin mutant W7FW14F. <i>European Biophysics Journal</i> , 2012, 41, 615-627.	1.2	13
67	Deamidation at Asparagine and Glutamine As a Major Modification upon Deterioration/Aging of Proteinaceous Binders in Mural Paintings. <i>Analytical Chemistry</i> , 2011, 83, 2056-2064.	3.2	86
68	Effects of the Known Pathogenic Mutations on the Aggregation Pathway of the Amyloidogenic Peptide of Apolipoprotein A-I. <i>Journal of Molecular Biology</i> , 2011, 407, 465-476.	2.0	48
69	A novel ErbB2 epitope targeted by human antitumor immunoagents. <i>FEBS Journal</i> , 2011, 278, 1156-1166.	2.2	12
70	Insights into the fate of the N-terminal amyloidogenic polypeptide of ApoA-I in cultured target cells. <i>Journal of Cellular and Molecular Medicine</i> , 2011, 15, 2652-2663.	1.6	24
71	Plasma nitroproteome of kidney disease patients. <i>Amino Acids</i> , 2011, 40, 653-667.	1.2	48
72	Polymerization of hemoglobins in Arctic fish: <i>Lycodes reticulatus</i> and <i>Gadus morhua</i> . <i>IUBMB Life</i> , 2011, 63, 346-354.	1.5	8

#	ARTICLE	IF	CITATIONS
73	The MicroRNA 15a/16 ¹ Cluster Down-regulates Protein Repair Isoaspartyl Methyltransferase in Hepatoma Cells. <i>Journal of Biological Chemistry</i> , 2011, 286, 43690-43700.	1.6	17
74	Effects of a lipid environment on the fibrillogenic pathway of the N-terminal polypeptide of human apolipoprotein A-I, responsible for in vivo amyloid fibril formation. <i>European Biophysics Journal</i> , 2010, 39, 1289-1299.	1.2	22
75	PED interacts with Rac1 and regulates cell migration/invasion processes in human non-small cell lung cancer cells. <i>Journal of Cellular Physiology</i> , 2010, 225, 63-72.	2.0	18
76	Mitochondrial Chaperone Trap1 and the Calcium Binding Protein Sorcin Interact and Protect Cells against Apoptosis Induced by Antitubercular Agents. <i>Cancer Research</i> , 2010, 70, 6577-6586.	0.4	120
77	Glycoproteome Study in Myocardial Lesions Serum by Integrated Mass Spectrometry Approach: Preliminary Insights. <i>European Journal of Mass Spectrometry</i> , 2010, 16, 123-149.	0.5	8
78	The co-chaperone BAG3 interacts with the cytosolic chaperonin CCT: New hints for actin folding. <i>International Journal of Biochemistry and Cell Biology</i> , 2010, 42, 641-650.	1.2	44
79	hnRNP H1 and intronic G runs in the splicing control of the human rpl3 gene. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2010, 1799, 419-428.	0.9	35
80	Stoichiometry and Topology of the Complex of the Endogenous ATP Synthase Inhibitor Protein IF ₁ with Calmodulin. <i>Biochemistry</i> , 2010, 49, 7542-7552.	1.2	7
81	Puzzle of protein complexes in vivo: a present and future challenge for functional proteomics. <i>Expert Review of Proteomics</i> , 2009, 6, 159-169.	1.3	21
82	Chromobox Protein Homologue 7 Protein, with Decreased Expression in Human Carcinomas, Positively Regulates E-Cadherin Expression by Interacting with the Histone Deacetylase 2 Protein. <i>Cancer Research</i> , 2009, 69, 7079-7087.	0.4	72
83	Enzymatically active fibrils generated by the self-assembly of the ApoA-I fibrillogenic domain functionalized with a catalytic moiety. <i>Biomaterials</i> , 2009, 30, 829-835.	5.7	19
84	Proteomic strategies for the identification of proteinaceous binders in paintings. <i>Analytical and Bioanalytical Chemistry</i> , 2009, 395, 2269-2280.	1.9	75
85	Different carbon sources affect lifespan and protein redox state during <i>Saccharomyces cerevisiae</i> chronological ageing. <i>Cellular and Molecular Life Sciences</i> , 2009, 66, 933-947.	2.4	28
86	Description of the topographical changes associated to the different stages of the DsbA catalytic cycle. <i>Protein Science</i> , 2009, 11, 1600-1612.	3.1	11
87	Technical advances in proteomics mass spectrometry: identification of post-translational modifications. <i>Clinical Chemistry and Laboratory Medicine</i> , 2009, 47, 647-65.	1.4	16
88	The molecular chaperone Hsp90 is a component of the cap-binding complex and interacts with the translational repressor Cup during <i>Drosophila</i> oogenesis. <i>Gene</i> , 2009, 432, 67-74.	1.0	39
89	The effect of prime-site occupancy on the hepatitis C virus NS3 protease structure. <i>Protein Science</i> , 2009, 11, 2102-2112.	3.1	5
90	Topological investigation of amyloid fibrils obtained from β 2-microglobulin. <i>Protein Science</i> , 2009, 11, 2362-2369.	3.1	53

#	ARTICLE	IF	CITATIONS
91	The peculiar structural features of kiwi fruit pectin methylesterase: Amino acid sequence, oligosaccharides structure, and modeling of the interaction with its natural proteinaceous inhibitor. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008, 71, 195-206.	1.5	39
92	Biophysical and biochemical characterization of a liposarcoma-derived recombinant MnSOD protein acting as an anticancer agent. <i>International Journal of Cancer</i> , 2008, 123, 2684-2695.	2.3	23
93	Peptidoglycan and Muropeptides from Pathogens <i>Agrobacterium</i> and <i>Xanthomonas</i> Elicit Plant Innate Immunity: Structure and Activity. <i>Chemistry and Biology</i> , 2008, 15, 438-448.	6.2	129
94	The different forms of PNS myelin P0 protein within and outside lipid rafts. <i>Journal of Neurochemistry</i> , 2008, 107, 291-301.	2.1	15
95	Hb Southern Italy: coexistence of two missense mutations (the Hb Sun Prairie \pm 2 130 Ala $\hat{+}$) Tj ETQq1 1 0.784314 rgB / Haematology, 2008, 143, 138-142.	1.2	4
96	In HspA from <i>Helicobacter pylori</i> vicinal disulfide bridges are a key determinant of domain B structure. <i>FEBS Letters</i> , 2008, 582, 3537-3541.	1.3	12
97	A Rapid and Selective Mass Spectrometric Method for the Identification of Nitrated Proteins. <i>Methods in Molecular Biology</i> , 2008, 477, 15-29.	0.4	9
98	Protease treatment affects both invasion ability and biofilm formation in <i>Listeria monocytogenes</i> . <i>Microbial Pathogenesis</i> , 2008, 45, 45-52.	1.3	81
99	cis-acting sequences and trans-acting factors in the localization of mRNA for mitochondrial ribosomal proteins. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2008, 1779, 820-829.	0.9	31
100	Multistep, sequential control of the trafficking and function of the multiple sulfatase deficiency gene product, SUMF1 by PDI, ERGIC-53 and ERp44. <i>Human Molecular Genetics</i> , 2008, 17, 2610-2621.	1.4	62
101	Hb Foggia or $\hat{A}117(GH5)Phe \rightarrow Ser$: a new $\hat{A}2$ globin allele affecting the $\hat{A}Hb-AHSP$ interaction. <i>Haematologica</i> , 2008, 93, 141-142.	1.7	23
102	Hb J-CAPE TOWN [$\hat{I}92(FG4)Arg \hat{+} Gln (\hat{I}1)$, CGG $\hat{+}$ CAG] in Southern Italy Found in a Patient with Erythrocytosis. <i>Hemoglobin</i> , 2007, 31, 113-120.	0.4	1
103	Bidimensional Tandem Mass Spectrometry for Selective Identification of Nitration Sites in Proteins. <i>Analytical Chemistry</i> , 2007, 79, 2109-2117.	3.2	51
104	Sulfatase modifying factor 1 trafficking through the cells: from endoplasmic reticulum to the endoplasmic reticulum. <i>EMBO Journal</i> , 2007, 26, 2443-2453.	3.5	42
105	Functional proteomics: protein-protein interactions in vivo. <i>Italian Journal of Biochemistry</i> , 2007, 56, 310-4.	0.3	5
106	Structure and Function of the Long Pentraxin PTX3 Glycosidic Moiety: Fine-Tuning of the Interaction with C1q and Complement Activation. <i>Biochemistry</i> , 2006, 45, 11540-11551.	1.2	113
107	Pancreatic cancer-derived S-100A8 N-terminal peptide: A diabetes cause?. <i>Clinica Chimica Acta</i> , 2006, 372, 120-128.	0.5	75
108	Recombinant amyloidogenic domain of ApoA-I: Analysis of its fibrillogenic potential. <i>Biochemical and Biophysical Research Communications</i> , 2006, 351, 223-228.	1.0	18

#	ARTICLE	IF	CITATIONS
109	Tubulin nitration in human gliomas. <i>Neuroscience Letters</i> , 2006, 394, 57-62.	1.0	25
110	Partial purification and MALDI-TOF MS analysis of UN1, a tumor antigen membrane glycoprotein. <i>International Journal of Biological Macromolecules</i> , 2006, 39, 122-126.	3.6	7
111	Lysine 58-cleaved beta2-microglobulin is not detectable by 2D electrophoresis in ex vivo amyloid fibrils of two patients affected by dialysis-related amyloidosis. <i>Protein Science</i> , 2006, 16, 343-349.	3.1	24
112	H-prune-nm23-H1 protein complex and correlation to pathways in cancer metastasis. <i>Journal of Bioenergetics and Biomembranes</i> , 2006, 38, 205-213.	1.0	33
113	Indole-3-acetic acid improves <i>Escherichia coli</i> ™s defences to stress. <i>Archives of Microbiology</i> , 2006, 185, 373-382.	1.0	129
114	Exploring the Mechanism of Formation of Native-like and Precursor Amyloid Oligomers for the Native Acylphosphatase from <i>Sulfolobus solfataricus</i> . <i>Structure</i> , 2006, 14, 993-1001.	1.6	36
115	The gene of an archaeal Î±-L-fucosidase is expressed by translational frameshifting. <i>Nucleic Acids Research</i> , 2006, 34, 4258-4268.	6.5	22
116	Ribosomal protein L7a binds RNA through two distinct RNA-binding domains. <i>Biochemical Journal</i> , 2005, 385, 289-299.	1.7	30
117	Proteomics of Î²2-microglobulin amyloid fibrils. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2005, 1753, 23-33.	1.1	36
118	Direct interactions among Ret, GDNF and GFRÎ±1 molecules reveal new insights into the assembly of a functional three-protein complex. <i>Cellular Signalling</i> , 2005, 17, 717-727.	1.7	43
119	Liquid crystalline elastomers based on diglycidyl terminated rigid monomers and aliphatic acids. Part 1. Synthesis and characterization. <i>Polymer</i> , 2005, 46, 2105-2121.	1.8	32
120	Interaction Proteomics. <i>Bioscience Reports</i> , 2005, 25, 45-56.	1.1	48
121	Biological properties of a human compact anti-ErbB2 antibody. <i>Carcinogenesis</i> , 2005, 26, 1890-1895.	1.3	37
122	Functional proteomics. <i>Clinica Chimica Acta</i> , 2005, 357, 140-150.	0.5	42
123	Pancreatic cancer-associated diabetes mellitus: An open field for proteomic applications. <i>Clinica Chimica Acta</i> , 2005, 357, 184-189.	0.5	33
124	Identification of proteins interacting with the RNAPII FCP1 phosphatase: FCP1 forms a complex with arginine methyltransferase PRMT5 and it is a substrate for PRMT5-mediated methylation. <i>FEBS Letters</i> , 2005, 579, 683-689.	1.3	62
125	CysMap and CysJoin: Database and tools for protein disulphides localisation. <i>FEBS Letters</i> , 2005, 579, 3048-3054.	1.3	3
126	Thermal Stability and Aggregation of <i>Sulfolobus solfataricus</i> Î²-Glycosidase Are Dependent upon the N-Î±-Methylation of Specific Lysyl Residues. <i>Journal of Biological Chemistry</i> , 2004, 279, 10185-10194.	1.6	36

#	ARTICLE	IF	CITATIONS
127	The Role of the Conserved Residues His-246, His-199, and Tyr-255 in the Catalysis of Catechol 2,3-Dioxygenase from <i>Pseudomonas stutzeri</i> OX1. <i>Journal of Biological Chemistry</i> , 2004, 279, 48630-48639.	1.6	51
128	Hierarchical Formation of Disulfide Bonds in the Immunoglobulin Fc Fragment Is Assisted by Protein-disulfide Isomerase. <i>Journal of Biological Chemistry</i> , 2004, 279, 15059-15066.	1.6	10
129	Conformational analysis of HAMLET, the folding variant of human α_1 -lactalbumin associated with apoptosis. <i>Protein Science</i> , 2004, 13, 1322-1330.	3.1	57
130	Tuber borchii fruit body: 2-dimensional profile and protein identification. <i>Phytochemistry</i> , 2004, 65, 813-820.	1.4	19
131	Hb Cardarelli [β 286(F2)Ala \rightarrow Pro]: A New Unstable and Hyperaffine Variant in Association with β -thalassemia. <i>Hemoglobin</i> , 2004, 28, 103-115.	0.4	8
132	Phenol Hydroxylase and Toluene/ o -Xylene Monooxygenase from <i>Pseudomonas stutzeri</i> OX1: Interplay between Two Enzymes. <i>Applied and Environmental Microbiology</i> , 2004, 70, 2211-2219.	1.4	113
133	The Regions of the Sequence Most Exposed to the Solvent Within the Amyloidogenic State of a Protein Initiate the Aggregation Process. <i>Journal of Molecular Biology</i> , 2004, 336, 253-262.	2.0	34
134	Purification and characterization of a small (7.3 kDa) putative lipid transfer protein from maize seeds. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2003, 794, 109-114.	1.2	26
135	Hexafluoroisopropanol and Acid Destabilized Forms of Apomyoglobin Exhibit Structural Differences. <i>Biochemistry</i> , 2003, 42, 312-319.	1.2	25
136	A novel zinc finger transcriptional repressor, ZNF224, interacts with the negative regulatory element (AldA-NRE) and inhibits gene expression. <i>FEBS Letters</i> , 2003, 534, 93-100.	1.3	29
137	The FCP1 phosphatase interacts with RNA polymerase II and with MEP50 a component of the methylosome complex involved in the assembly of snRNP. <i>Nucleic Acids Research</i> , 2003, 31, 999-1005.	6.5	20
138	Assignment of Disulphide Bridges in Par j 2.0101, a Major Allergen of <i>Parietaria judaica</i> Pollen. <i>Biological Chemistry</i> , 2003, 384, 1165-1172.	1.2	14
139	Structural and biochemical characterization of a new type of lectin isolated from carp eggs. <i>Biochemical Journal</i> , 2003, 376, 433-440.	1.7	40
140	Structural Characterization of the M* Partly Folded Intermediate of Wild Type and P138A Aspartate Aminotransferase from <i>Escherichia coli</i> . <i>Journal of Biological Chemistry</i> , 2002, 277, 17428-17437.	1.6	11
141	Hb VILA REAL [β 36(C2)Pro \rightarrow His] IN ITALY: CHARACTERIZATION OF THE AMINO ACID SUBSTITUTION AND THE DNA MUTATION. <i>Hemoglobin</i> , 2002, 26, 21-31.	0.4	2
142	Binding and Relaxometric Properties of Heme Complexes with Cyanogen Bromide Fragments of Human Serum Albumin. <i>Biophysical Journal</i> , 2002, 83, 2248-2258.	0.2	17
143	Molecular Basis of Phospholipase A2 Inhibition by Petrosaspongiolide M. <i>ChemBioChem</i> , 2002, 3, 664.	1.3	34
144	Expression and purification of the recombinant subunits of toluene/o -xylene monooxygenase and reconstitution of the active complex. <i>FEBS Journal</i> , 2002, 269, 5689-5699.	0.2	67

#	ARTICLE	IF	CITATIONS
145	Slow Folding of Three-Fingered Toxins Is Associated with the Accumulation of Native Disulfide-Bonded Intermediates. <i>Biochemistry</i> , 2001, 40, 15257-15266.	1.2	17
146	A nucleotide insertion and frameshift cause albumin K ⁶⁷ N, an extended and O-glycosylated mutant of human serum albumin with two additional disulfide bridges. <i>FEBS Journal</i> , 2001, 268, 344-352.	0.2	19
147	Conformational analysis of putative regulatory subunit D of the toluene/o-xylene-monooxygenase complex from <i>Pseudomonas stutzeri</i> OX1. <i>Protein Science</i> , 2001, 10, 482-490.	3.1	12
148	Assignment of the Complete Disulphide Bridge Pattern in the Human Recombinant Follitropin β^2 -Chain. <i>Biological Chemistry</i> , 2001, 382, 961-8.	1.2	6
149	IDENTIFICATION OF Hb VILLEJUIF [¹²³ (H1)Thr ⁶⁷ Ile] IN SOUTHERN ITALY. <i>Hemoglobin</i> , 2001, 25, 67-78.	0.4	1
150	Structural Studies of the Complex Between Decapeptide Inhibitors and the Serine Protease NS3/4A of Hepatitis C Virus. , 2001, , 545-546.		0
151	Structural characterization of the oligosaccharide chains of human β_2 -microglobulin from urine and amniotic fluid. <i>FEBS Journal</i> , 2000, 267, 2105-2112.	0.2	19
152	Mass spectrometry study of ecto-5 α -nucleotidase from bull seminal plasma. <i>FEBS Journal</i> , 2000, 267, 4978-4987.	0.2	21
153	Characterization of five new low-molecular-mass trypsin inhibitors from white mustard (<i>Sinapis Tj</i> ETQq1 1 0.784314 rgBT / Overlock 10	0.2	18
154	Structural and Functional Features of Modified Heat-Stable Toxins Produced by Enteropathogenic <i>Klebsiella</i> Cells. <i>Pediatric Research</i> , 2000, 48, 685-690.	1.1	9
155	Trypsin Sheds Light on the Singular Case of Seminal RNase, a Dimer with Two Quaternary Conformations. <i>Journal of Biological Chemistry</i> , 2000, 275, 8000-8006.	1.6	10
156	Modern Mass Spectrometric Methodologies in Monitoring Milk Quality. <i>Analytical Chemistry</i> , 2000, 72, 408-415.	3.2	93
157	Binding of β -Actinin to Titin: Implications for Z-Disk Assembly. <i>Biochemistry</i> , 2000, 39, 5255-5264.	1.2	47
158	Early intermediates in the PDI ¹ -assisted folding of ribonuclease A. <i>Protein Science</i> , 2000, 9, 525-535.	3.1	22
159	Novel Autocrine and Paracrine Loops of the Stem Cell Factor/Chymase Network. <i>International Archives of Allergy and Immunology</i> , 1999, 118, 422-425.	0.9	17
160	Conformational changes in the NS3 protease from hepatitis C virus strain Bk monitored by limited proteolysis and mass spectrometry. <i>Protein Science</i> , 1999, 8, 1445-1454.	3.1	23
161	Characterization of low-molecular-mass trypsin isoinhibitors from oil-rape (<i>Brassica napus</i> var.) Tj ETQq1 1 0.784314 rgBT / Overlock 10	0.2	31
162	Structural and membrane-binding properties of saposin D. <i>FEBS Journal</i> , 1999, 263, 486-494.	0.2	24

#	ARTICLE	IF	CITATIONS
163	Topology of the Thyroid Transcription Factor 1 Homeodomain-DNA Complex. <i>Biochemistry</i> , 1999, 38, 64-72.	1.2	25
164	Multiple Determinants Influence Complex Formation of the Hepatitis C Virus NS3 Protease Domain with Its NS4A Cofactor Peptide. <i>Biochemistry</i> , 1999, 38, 5206-5215.	1.2	31
165	Conformational Changes in Human Hepatitis C Virus NS3 Protease upon Binding of Product-Based Inhibitors. <i>Biochemistry</i> , 1999, 38, 13844-13852.	1.2	33
166	Analysis of human serum albumin variants by mass spectrometric procedures. <i>BBA - Proteins and Proteomics</i> , 1998, 1384, 79-92.	2.1	8
167	Axinellins A and B: New Proline-Containing Antiproliferative Cyclopeptides from the Vanuatu Sponge <i>Axinella carteri</i> . <i>European Journal of Organic Chemistry</i> , 1998, 1998, 2659-2665.	1.2	57
168	Selective and asymmetric action of trypsin on the dimeric forms of seminal RNase. <i>Protein Science</i> , 1998, 7, 2653-2658.	3.1	6
169	The Length of a Single Turn Controls the Overall Folding Rate of α -Three-Fingered-Snake Toxins. <i>Biochemistry</i> , 1998, 37, 16060-16068.	1.2	15
170	Thetbf-1 Gene from the White Truffle <i>Tuber borchii</i> Codes for a Structural Cell Wall Protein Specifically Expressed in Fruitbody1. <i>Fungal Genetics and Biology</i> , 1998, 25, 87-99.	0.9	28
171	Topology of the calmodulin-melittin complex 1 Edited by P.E. Wright. <i>Journal of Molecular Biology</i> , 1998, 277, 945-958.	2.0	90
172	Structural characterization and independent folding of a chimeric glycoprotein comprising granulocyte-macrophage colony stimulating factor and erythropoietin sequences. <i>Glycobiology</i> , 1998, 8, 779-790.	1.3	4
173	Amino Acid Sequence, S-S Bridge Arrangement and Distribution in Plant Tissues of Thionins from <i>Viscum album</i> . <i>Biological Chemistry</i> , 1997, 378, 989-996.	1.2	37
174	Effect of Glutaredoxin and Protein Disulfide Isomerase on the Glutathione-Dependent Folding of Ribonuclease A. <i>Biochemistry</i> , 1997, 36, 12259-12267.	1.2	61
175	Microheterogeneity of Odorant-Binding Proteins in the Porcupine Revealed by N-Terminal Sequencing and Mass Spectrometry. <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 1997, 117, 287-291.	0.7	23
176	Identification of the prion protein allotypes which accumulate in the brain of sporadic and familial Creutzfeldt-Jakob disease patients. <i>Nature Medicine</i> , 1997, 3, 521-525.	15.2	58
177	Structural Characterization of Four Genetic Variants of Human Serum Albumin Associated with Alloalbuminemia in Italy. <i>FEBS Journal</i> , 1997, 247, 476-482.	0.2	13
178	Surface topology of Minibody by selective chemical modifications and mass spectrometry. <i>Protein Science</i> , 1997, 6, 1901-1909.	3.1	52
179	Glutathione-Dependent Pathways of Refolding of RNase T1 by Oxidation and Disulfide Isomerization: Catalysis by Protein Disulfide Isomerase. <i>Biochemistry</i> , 1996, 35, 13636-13646.	1.2	42
180	Structure of fuscopeptins, phytotoxic metabolites of <i>Pseudomonas fuscovaginae</i> . <i>FEBS Letters</i> , 1996, 381, 213-216.	1.3	55

#	ARTICLE	IF	CITATIONS
181	Assignment of protein disulphides by a computer method using mass spectrometric data. FEBS Letters, 1996, 393, 241-247.	1.3	12
182	Amino acid sequence and molecular modelling of glycoprotein IIb-IIIa and fibronectin receptor iso-antagonists from <i>Trimeresurus elegans</i> venom. Biochemical Journal, 1996, 319, 775-782.	1.7	32
183	The Gene, Protein and Glycan Structures of Laccase from <i>Pleurotus ostreatus</i> . FEBS Journal, 1996, 235, 508-515.	0.2	93
184	Structural Characterisation of Human Recombinant Glycophormones Follitropin, Lutropin and Choriogonadotropin Expressed in Chinese Hamster Ovary Cells. FEBS Journal, 1996, 242, 608-618.	0.2	47
185	Identification of disulphide bonds in the refolding of bovine pancreatic RNase A. Folding & Design, 1996, 1, 381-390.	4.5	18
186	Probing the tertiary structure of proteins by limited proteolysis and mass spectrometry: The case of minibody. Protein Science, 1996, 5, 802-813.	3.1	62
187	Transglutaminase from Rat Coagulating Gland Secretion. Journal of Biological Chemistry, 1996, 271, 27416-27423.	1.6	39
188	Amino Acid Sequence and Disulphide-bridge Pattern of three gamma-Thionins from <i>Sorghum bicolor</i> . FEBS Journal, 1995, 228, 250-256.	0.2	10
189	Human β -fetoprotein produced from hep G2 cell line: Structure and heterogeneity of the oligosaccharide moiety. Journal of Mass Spectrometry, 1995, 30, 632-638.	0.7	9
190	Structural Analysis of Saposin C and B. Journal of Biological Chemistry, 1995, 270, 9953-9960.	1.6	79
191	Neurokinin Receptors Could Be Differentiated by Their Capacity to Respond to the Transglutaminase-synthesized β -(Glutamyl ⁵)Spermine Derivative of Substance P. Journal of Neurochemistry, 1995, 65, 420-426.	2.1	22
192	Amino Acid Sequence and Disulphide-bridge Pattern of three gamma-Thionins from <i>Sorghum bicolor</i> . FEBS Journal, 1995, 228, 250-256.	0.2	42
193	Hb G-Miwlo [β 64(E13)ASP β 'ASN] Observed in a Caucasian Family. Hemoglobin, 1994, 18, 53-56.	0.4	5
194	Relevance of chlorine-substituent for the antifungal activity of syringomycin and syringotoxin, metabolites of the phytopathogenic bacterium <i>Pseudomonas syringae</i> pv. <i>syringae</i> . Experientia, 1994, 50, 130-133.	1.2	48
195	Post-translational modifications in aspartate aminotransferase from <i>Sulfolobus solfataricus</i> . Detection of N-e-methyllysines by mass spectrometry. FEBS Journal, 1994, 222, 761-767.	0.2	15
196	[d-Leu ²]Deltorpin, a 17 amino acid opioid peptide from the skin of the Brazilian hylid frog, <i>Phyllomedusa burmeisteri</i> . Peptides, 1994, 15, 199-202.	1.2	35
197	Novel bioactive lipodepsipeptides from <i>Pseudomonas syringae</i> : The pseudomycins. FEBS Letters, 1994, 355, 96-100.	1.3	86
198	Production and structural characterization of amino terminally histidine tagged human oncostatin M in <i>E. Coli</i> . Cytokine, 1994, 6, 255-264.	1.4	14

#	ARTICLE	IF	CITATIONS
199	Hb F-Sassari: A Novel G ^β Variant with a Threonine Residue at Position ^β 75, Characterized by Mass Spectrometry Techniques. Hemoglobin, 1994, 18, 307-315.	0.4	12
200	[4] Structural characterization of hemoglobin variants using capillary electrophoresis and fast atom bombardment mass spectrometry. Methods in Enzymology, 1994, 231, 45-65.	0.4	5
201	Human-immunodeficiency-virus transmembrane glycoprotein gp41 is an amino acceptor and donor substrate for transglutaminase in vitro. FEBS Journal, 1993, 215, 99-104.	0.2	22
202	Structural characterization of a biologically active human lipocortin 1 expressed in Escherichia coli. FEBS Journal, 1993, 211, 347-355.	0.2	20
203	Hemoglobin Ozieri: a new β -chain variant (β 71(E20)Ala \rightarrow Val). Characterization using FAB- and electrospray-mass spectrometric techniques. BBA - Proteins and Proteomics, 1993, 1162, 203-208.	2.1	16
204	Disulfide isoform intermediates in the reoxidation of recombinant human basic fibroblast growth factor. Biochemistry, 1993, 32, 4991-4996.	1.2	4
205	HB O-Arab [β 121(GH4)GLU \rightarrow LYS]: Association with DNA Polymorphisms of African Ancestry in two Mediterranean Families. Hemoglobin, 1993, 17, 523-535.	0.4	2
206	HB City of Hope [β 69(E13)GLY \rightarrow SER] in Italy: Association of the Gene with Haplotype IX. Hemoglobin, 1992, 16, 27-34.	0.4	4
207	Substance P inactivation by transglutaminase in vitro. Peptides, 1992, 13, 151-154.	1.2	16
208	Enzyme catalysed lactonization of 3,5 dihydroxy esters: Enantioselective synthesis of naturally occurring 3-hydroxy-5-decanolide, (β)-massoialactone, and 3-hydroxy-5-icosanolide.. Tetrahedron: Asymmetry, 1992, 3, 29-32.	1.8	27
209	River buffalo (Bubalus bubalis L.) AA phenotype haemoglobins: characterization by immobilized polyacrylamide gel electrophoresis and high performance liquid chromatography and determination of the primary structure of the constitutive chains by mass spectrometry. Comparative Biochemistry and Physiology Part B: Comparative Biochemistry, 1992, 101, 91-98.	0.2	4
210	Characterization by mass spectrometry of a recombinant hepatitis delta virus antigen and its proteolytic products. FEBS Journal, 1992, 204, 515-521.	0.2	2
211	Stabilization of recombinant human basic fibroblast growth factor by chemical modifications of cysteine residues. FEBS Journal, 1992, 204, 649-655.	0.2	31
212	Factors affecting the fast atom bombardment mass spectrometric analysis of proteolytic digests of proteins. Biological Mass Spectrometry, 1992, 21, 22-26.	0.5	14
213	A remarkable short synthesis of optically active mevinic acid analogs by biocatalytic lactonization of syn-3,5-dihydroxy esters. Journal of Organic Chemistry, 1991, 56, 4050-4052.	1.7	35
214	A third instance of the high oxygen affinity variant, HB heathrow [β 103(G5)PHE \rightarrow LEU]: Identification of the mutation by mass spectrometry and by DNA analysis. Hemoglobin, 1991, 15, 43-51.	0.4	19
215	Human α -fetoprotein primary structure: a mass spectrometric study. Biochemistry, 1991, 30, 5061-5066.	1.2	39
216	Mass spectrometric identification of the amino donor and acceptor sites in a transglutaminase protein substrate secreted from rat seminal vesicles. Biochemistry, 1991, 30, 3114-3120.	1.2	51

#	ARTICLE	IF	CITATIONS
217	Syringopeptins, new phytotoxic lipodepsipeptides of <i>Pseudomonas syringae</i> pv. <i>syringae</i> . <i>FEBS Letters</i> , 1991, 291, 109-112.	1.3	126
218	Single-step purification and structural characterization of human interleukin-6 produced in <i>Escherichia coli</i> From a T7 RNA polymerase expression vector. <i>FEBS Journal</i> , 1991, 198, 541-547.	0.2	128
219	Assignment of the five disulfide bridges in an alpha-amylase inhibitor from wheat kernel by fast-atom-bombardment mass spectrometry and Edman degradation. <i>FEBS Journal</i> , 1991, 199, 595-600.	0.2	39
220	FAB overlapping: a strategy for sequencing homologous proteins. <i>International Journal of Mass Spectrometry and Ion Processes</i> , 1991, 111, 287-300.	1.9	5
221	Capillary zone electrophoresis and mass spectrometry for the characterization of genetic variants of human hemoglobin. <i>Analytical Biochemistry</i> , 1991, 194, 1-8.	1.1	54
222	Structure of syringotoxin, a bioactive metabolite of <i>Pseudomonas syringae</i> pv. <i>syringae</i> . <i>FEBS Letters</i> , 1990, 269, 377-380.	1.3	86
223	Certain N-terminal peptides inhibit uptake of mature aspartate aminotransferase by isolated mitochondria. <i>Biochemical and Biophysical Research Communications</i> , 1990, 170, 609-615.	1.0	3
224	Synthesis and conformational studies of peptides encompassing the carboxy-terminal helix of thermolysin. <i>International Journal of Peptide and Protein Research</i> , 1990, 35, 396-405.	0.1	3
225	Synthesis and characterization of a recombinant fragment of human β_2 -fetoprotein with antigenic selectivity versus albumin. <i>Protein Engineering, Design and Selection</i> , 1989, 2, 605-610.	1.0	5
226	Analysis by fast atom bombardment mass spectrometry of 4,4-dicarboxy-5-(pyridoxyl-5-phosphate)-proline, of 4-carboxy-5-(pyridoxyl-5-phosphate)-proline and 4,4-dicarboxy-5-pyridoxylproline. <i>Biomedical & Environmental Mass Spectrometry</i> , 1989, 18, 995-999.	1.6	2
227	The structure of syringomycins A1, E and G. <i>FEBS Letters</i> , 1989, 255, 27-31.	1.3	158
228	Substance P as a transglutaminase substrate: Identification of the reaction products by fast atom bombardment mass spectrometry. <i>Analytical Biochemistry</i> , 1988, 172, 499-503.	1.1	27
229	β_2 -Endorphin modification by transglutaminase in vitro: Identification by FABMS of glutamine-11 and lysine-29 as acyl donor and acceptor sites. <i>Biochemical and Biophysical Research Communications</i> , 1988, 154, 735-740.	1.0	29
230	Multiple forms of syringomycin. <i>Physiological and Molecular Plant Pathology</i> , 1988, 33, 493-496.	1.3	46
231	Identification by Fast Atom Bombardment Mass Spectrometry of HB Indianapolis [125 I(C14)CYS β 'ARG] in a Family from Naples, Italy. <i>Hemoglobin</i> , 1988, 12, 323-336.	0.4	30
232	Enzymatic methyl esterification of synthetic tripeptides: structural requirements of the peptide substrate. Detection of the reaction products by fast-atom-bombardment mass spectrometry. <i>FEBS Journal</i> , 1988, 177, 233-239.	0.2	17
233	Digestion by pancreatic juice of a beta-casein-containing fragment of buffalo beta-casein. <i>International Journal of Peptide and Protein Research</i> , 1987, 29, 504-508.	0.1	19
234	Assignment of phosphorylation sites in buffalo β_2 -casein by fast atom bombardment mass spectrometry. <i>Biochemical and Biophysical Research Communications</i> , 1986, 140, 28-37.	1.0	25

#	ARTICLE	IF	CITATIONS
235	Sequence determination of cyanogen bromide-peptides by combined use of high-performance liquid chromatography fast atom bombardment mass spectrometry. <i>Journal of Chromatography A</i> , 1985, 331, 425-431.	1.8	11
236	A new method for rapid assignment of S-S bridges in proteins. <i>Biochemical and Biophysical Research Communications</i> , 1985, 126, 1122-1128.	1.0	150
237	Protein fingerprint by fast atom bombardment mass spectrometry: Characterization of normal and variant human haemoglobins. <i>Biochemical and Biophysical Research Communications</i> , 1985, 130, 84-90.	1.0	66
238	Isolation and characterization of dipeptidyl peptidase IV from human meconium. <i>FEBS Letters</i> , 1985, 184, 273-277.	1.3	26
239	Modified calmodulin calcium binding domain III.. <i>International Journal of Peptide and Protein Research</i> , 1984, 23, 454-461.	0.1	9
240	Separation of phenylthiohydantoin ^α amino acids by high-performance liquid chromatography. <i>Journal of Chromatography A</i> , 1983, 270, 371-377.	1.8	41
241	Studies on gliadin related peptides. I. Synthesis, purification and 1H n.m.r. characterization of the pentapeptide H ^α Tyr ^β (Gln) ³ Pro ^γ OH. <i>International Journal of Peptide and Protein Research</i> , 1983, 22, 482-488. ^{0.1}		2
242	Reactivity of sulphhydryl groups of cytosolic and mitochondrial bovine aspartate aminotransferases. <i>Molecular and Cellular Biochemistry</i> , 1981, 35, 121-128.	1.4	8
243	Mitochondrial bovine aspartate aminotransferase. <i>FEBS Letters</i> , 1979, 101, 351-354.	1.3	19
244	Limited Proteolysis Mass Spectrometry of Protein Complexes. , 0, , 63-82.		3