Joaquin Abian

List of Publications by Year in descending order

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101384 123241 4,569 132 36 61 citations g-index h-index papers 134 134 134 6173 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Complexes of Iron with Phenolic Compounds from Soybean Nodules and Other Legume Tissues: Prooxidant and Antioxidant Properties. Free Radical Biology and Medicine, 1997, 22, 861-870.	1.3	315
2	Comparison of conventional, narrow-bore and capillary liquid chromatography/mass spectrometry for electrospray ionization mass spectrometry: practical considerations. Journal of Mass Spectrometry, 1999, 34, 244-254.	0.7	170
3	General Statistical Framework for Quantitative Proteomics by Stable Isotope Labeling. Journal of Proteome Research, 2014, 13, 1234-1247.	1.8	165
4	Proteomics-based identification of novelCandida albicans antigens for diagnosis of systemic candidiasis in patients with underlying hematological malignancies. Proteomics, 2004, 4, 3084-3106.	1.3	150
5	The defense response of germinating maize embryos against fungal infection: A proteomics approach. Proteomics, 2004, 4, 383-396.	1.3	144
6	Preparation and Evaluation of Immunosorbents for Selective Trace Enrichment of Phenylurea and Triazine Herbicides in Environmental Waters. Analytical Chemistry, 1995, 67, 2451-2460.	3.2	122
7	The Biliverdin Chromophore Binds Covalently to a Conserved Cysteine Residue in the N-Terminus of Agrobacterium Phytochrome Agp 1â€. Biochemistry, 2004, 43, 3659-3669.	1.2	121
8	The coupling of gas and liquid chromatography with mass spectrometry. Journal of Mass Spectrometry, 1999, 34, 157-168.	0.7	119
9	Roles of hnRNP A1, SR Proteins, and p68 Helicase in c-H- ras Alternative Splicing Regulation. Molecular and Cellular Biology, 2003, 23, 2927-2941.	1.1	116
10	Fish proteome analysis: Model organisms and nonâ€sequenced species. Proteomics, 2010, 10, 858-872.	1.3	113
11	rBAT-b ^{0,+} AT heterodimer is the main apical reabsorption system for cystine in the kidney. American Journal of Physiology - Renal Physiology, 2002, 283, F540-F548.	1.3	91
12	Diethylmaleate activates the transcription factor Pap1 by covalent modification of critical cysteine residues. Molecular Microbiology, 2002, 45, 243-254.	1.2	87
13	Leghemoglobin green derivatives with nitrated hemes evidence production of highly reactive nitrogen species during aging of legume nodules. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 2660-2665.	3.3	81
14	Quantitative peptide bioanalysis using column-switching nano liquid chromatography/mass spectrometry. , 1998, 33, 976-983.		76
15	Comparative photodegradation rates of alachlor and bentazone in natural water and determination of breakdown products. Environmental Toxicology and Chemistry, 1995, 14, 1287-1298.	2.2	73
16	Isotope dilution mass spectrometry for absolute quantification in proteomics: Concepts and strategies. Journal of Proteomics, 2014, 96, 184-199.	1.2	73
17	Analysis of chlorotriazines and their degradation products in environmental samples by selecting various operating modes in thermospray HPLC/MS/MS. Journal of Agricultural and Food Chemistry, 1993, 41, 1264-1273.	2.4	72
18	Urine Proteomics to Detect Biomarkers for Chronic Allograft Dysfunction. Journal of the American Society of Nephrology: JASN, 2009, 20, 428-435.	3.0	70

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19	Leghemoglobin is nitrated in functional legume nodules in a tyrosine residue within the heme cavity by a nitrite/peroxideâ€dependent mechanism. Plant Journal, 2015, 81, 723-735.	2.8	70
20	High-performance liquid chromatographyâ€"thermospray mass spectrometry of ten sulfonamide antibiotics. Journal of Chromatography A, 1993, 629, 267-276.	1.8	68
21	Phosphorylation Analysis of Primary Human T Lymphocytes Using Sequential IMAC and Titanium Oxide Enrichment. Journal of Proteome Research, 2008, 7, 5167-5176.	1.8	68
22	Electrospray ionization mass spectrometry of oligosaccharides derived from fucoidan of Ascophyllum nodosum. Carbohydrate Research, 2007, 342, 826-834.	1.1	67
23	Flavin excretion from roots of iron-deficient sugar beet (Beta vulgaris L.). Planta, 1994, 193, 514-519.	1.6	62
24	The 2-oxoglutarate carrier promotes liver cancer by sustaining mitochondrial GSH despite cholesterol loading. Redox Biology, 2018, 14, 164-177.	3.9	59
25	Characterization of the Human Plasma Phosphoproteome Using Linear Ion Trap Mass Spectrometry and Multiple Search Engines. Journal of Proteome Research, 2010, 9, 876-884.	1.8	54
26	Absolute and Site-Specific Quantification of Protein Phosphorylation Using Integrated Elemental and Molecular Mass Spectrometry:  Its Potential To Assess Phosphopeptide Enrichment Procedures. Analytical Chemistry, 2008, 80, 1777-1787.	3.2	53
27	Utility of proteomics to assess pollutant response of clams from the Doñana bank of Guadalquivir Estuary (SW Spain). Proteomics, 2006, 6, S245-S255.	1.3	52
28	Proteomic analysis of plasma from patients with systemic lupus erythematosus: Increased presence of haptoglobin $\hat{l}\pm 2$ polypeptide chains over the $\hat{l}\pm 1$ isoforms. Proteomics, 2006, 6, S282-S292.	1.3	51
29	Thyroglobulin Peptides Associate In Vivo to HLA-DR in Autoimmune Thyroid Glands. Journal of Immunology, 2008, 181, 795-807.	0.4	48
30	Light-Induced Conformational Changes of Cyanobacterial Phytochrome Cph1 Probed by Limited Proteolysis and Autophosphorylation. Biochemistry, 2005, 44, 450-461.	1.2	47
31	Cell viability and proteomic analysis in cultured neurons exposed to methylmercury. Human and Experimental Toxicology, 2007, 26, 263-272.	1.1	47
32	Dissection of the HLA-DR4 Peptide Repertoire in Endocrine Epithelial Cells: Strong Influence of Invariant Chain and HLA-DM Expression on the Nature of Ligands. Journal of Immunology, 2004, 173, 1085-1093.	0.4	46
33	Determination of protein markers in human serum: Analysis of protein expression in toxic oil syndrome studies. Proteomics, 2004, 4, 303-315.	1.3	44
34	Functional Characterization of an Unusual Phytochelatin Synthase, LjPCS3, of <i>Lotus japonicus</i> Plant Physiology, 2008, 148, 536-545.	2.3	41
35	Preparation of porous n-type silicon sample plates for desorption/ionization on silicon mass spectrometry (DIOS-MS). Lab on A Chip, 2002, 2, 247-253.	3.1	40
36	Two-Dimensional Difference Gel Electrophoresis Urinary Proteomic Profile in the Search of Nonimmune Chronic Allograft Dysfunction Biomarkers. Transplantation, 2010, 89, 548-558.	0.5	40

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37	Biochemical characterization of recombinant and circulating human Spalpha. Tissue Antigens, 2004, 63, 335-344.	1.0	38
38	Composition of the HLAâ€DRâ€associated human thymus peptidome. European Journal of Immunology, 2013, 43, 2273-2282.	1.6	38
39	Increased CD38 expression in T cells and circulating anti-CD38 IgG autoantibodies differentially correlate with distinct cytokine profiles and disease activity in systemic lupus erythematosus patients. Cytokine, 2013, 62, 232-243.	1.4	37
40	Methylmercury disrupts the balance between phosphorylated and non-phosphorylated cofilin in primary cultures of mice cerebellar granule cells A proteomic study. Toxicology and Applied Pharmacology, 2010, 242, 109-118.	1.3	36
41	DrosophilaMTN: a metazoan copper-thionein related to fungal forms. FEBS Letters, 2000, 467, 189-194.	1.3	33
42	F2 isoprostane is already increased at the onset of type 1 diabetes mellitus: Effect of glycemic control. Metabolism: Clinical and Experimental, 2004, 53, 1118-1120.	1.5	33
43	Effect of Abscisic Acid on the Linoleic Acid Metabolism in Developing Maize Embryos. Plant Physiology, 1991, 95, 1277-1283.	2.3	31
44	Storage time and deodorization temperature influence the formation of aniline-derived compounds in denatured rapeseed oils. Food and Chemical Toxicology, 2001, 39, 91-96.	1.8	31
45	HLA-DR4 Molecules in Neuroendocrine Epithelial Cells Associate to a Heterogeneous Repertoire of Cytoplasmic and Surface Self Peptides. Journal of Immunology, 2002, 169, 5052-5060.	0.4	31
46	Determination of d-fagomine in buckwheat and mulberry by cation exchange HPLC/ESI–Q-MS. Analytical and Bioanalytical Chemistry, 2012, 402, 1953-1960.	1.9	31
47	Differential Expression of Proteins From Cultured Endothelial Cells Exposed to Uremic Versus Normal Serum. American Journal of Kidney Diseases, 2008, 51, 603-612.	2.1	30
48	2â€D DIGE analysis of Senegalese sole (<i>Solea senegalensis</i>) testis proteome in wildâ€caught and hormoneâ€treated F1 fish. Proteomics, 2009, 9, 2171-2181.	1.3	30
49	Quantitative electrospray LC–MS and LC–MS/MS in biomedicine. Journal of Pharmaceutical and Biomedical Analysis, 1998, 17, 1129-1138.	1.4	29
50	Transcriptional and proteomic profiling of flatfish (<i>Solea senegalensis</i>) spermatogenesis. Proteomics, 2011, 11, 2195-2211.	1.3	29
51	Cigarette smoke concentrate increases 8-epi-PGF2\$alpha; and TGF\$beta;1 secretion in rat mesangial cells. Life Sciences, 2004, 75, 611-621.	2.0	26
52	Surfing Transcriptomic Landscapes. A Step beyond the Annotation of Chromosome 16 Proteome. Journal of Proteome Research, 2014, 13, 158-172.	1.8	26
53	Characterization of phenolic glucosides from soybean root nodules by ion-exchange high performance liquid chromatography, ultraviolet spectroscopy and electrospray mass spectrometry. Phytochemical Analysis, 1998, 9, 171-176.	1.2	24
54	Lipoprotein lipase is nitrated in vivo after lipopolysaccharide challenge. Free Radical Biology and Medicine, 2009, 47, 1553-1560.	1.3	24

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55	Proteomic study of neuron and astrocyte cultures from senescenceâ€accelerated mouse SAMP8 reveals degenerative changes. Journal of Neurochemistry, 2009, 111, 945-955.	2.1	24
56	The presence of d-fagomine in the human diet from buckwheat-based foodstuffs. Food Chemistry, 2013, 136, 1316-1321.	4.2	24
57	Urinary Proteome Analysis Identified Neprilysin and VCAM as Proteins Involved in Diabetic Nephropathy. Journal of Diabetes Research, 2018, 2018, 1-12.	1.0	24
58	Formation of the adduct 6-(deoxyguanosin-N2-yl)-3-amino-benzo[a]pyrene from the mutagenic environmental contaminant 3-nitrobenzo[a]pyrene. Carcinogenesis, 1993, 14, 1065-1067.	1.3	23
59	Direct analysis of the major human seminal prostaglandins by thermospray high-performance liquid chromatography—mass spectrometry. Journal of Chromatography A, 1987, 394, 147-153.	1.8	22
60	Determination of Aniline Derivatives in Oils Related to the Toxic Oil Syndrome by Atmospheric Pressure Ionization-Tandem Mass Spectrometry. Analytical Chemistry, 2001, 73, 3828-3837.	3.2	22
61	Characterization of peptides and proteins in commercial HSA solutions. Proteomics, 2010, 10, 172-181.	1.3	22
62	A Comprehensive Tyrosine Phosphoproteomic Analysis Reveals Novel Components of the Platelet CLEC-2 Signaling Cascade. Thrombosis and Haemostasis, 2020, 120, 262-276.	1.8	22
63	Thermospray liquid chromatography/mass spectrometry of prostaglandin methyl ester derivatives: Application to the determination of prostaglandins E2 and D2 in rat gastric mucosa. Biological Mass Spectrometry, 1988, 16, 215-219.	0.5	21
64	Increased expression and phosphorylation of the two S100A9 isoforms in mononuclear cells from patients with systemic lupus erythematosus: A proteomic signature for circulating low-density granulocytes. Journal of Proteomics, 2012, 75, 1778-1791.	1.2	21
65	A Candida albicans PeptideAtlas. Journal of Proteomics, 2014, 97, 62-68.	1.2	21
66	Non-aqueous capillary electrophoresis of the positional isomers of a sulfated monosaccharide. Journal of Chromatography A, 2003, 987, 467-476.	1.8	20
67	The effects of smoking and its cessation on 8-epi-PGF2α and transforming growth factor-beta 1 in Type 1 diabetes mellitus. Diabetic Medicine, 2004, 21, 285-289.	1.2	19
68	The peptide-binding motif of HLA-DR8 shares important structural features with other type 1 diabetes-associated alleles. Genes and Immunity, 2011, 12, 504-512.	2.2	19
69	The phosphoproteome of human Jurkat T cell clones upon costimulation with anti-CD3/anti-CD28 antibodies. Journal of Proteomics, 2016, 131, 190-198.	1.2	19
70	Transcriptomic and proteomic analysis of liver and muscle alterations caused by surgical stress in rats. Clinical Science, 2005, 108, 167-178.	1.8	18
71	Expression of differential antennal proteins in males and females of an important crop pest, Sesamia nonagrioides. Insect Biochemistry and Molecular Biology, 2009, 39, 11-19.	1.2	18
72	Identification of p21Cip1 binding proteins by gel electrophoresis and capillary liquid chromatography microelectrospray tandem mass spectrometry. Proteomics, 2002, 2, 455.	1.3	17

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73	UTP affects the Schwannoma cell line proteome through P2Y receptors leading to cytoskeletal reorganisation. Proteomics, 2012, 12, 145-156.	1.3	17
74	Spanish Human Proteome Project: Dissection of Chromosome 16. Journal of Proteome Research, 2013, 12, 112-122.	1.8	17
75	Comparative study of different thermospray interfaces with carbamate pesticides: Influence of the ion source geometry. Journal of the American Society for Mass Spectrometry, 1995, 6, 656-667.	1.2	16
76	A Combination of Proteomic Approaches Identifies A Panel of Circulating Extracellular Vesicle Proteins Related to the Risk of Suffering Cardiovascular Disease in Obese Patients. Proteomics, 2019, 19, e1800248.	1.3	16
77	LymPHOS 2.0: an update of a phosphosite database of primary human T cells. Database: the Journal of Biological Databases and Curation, 2015, 2015, bav115.	1.4	15
78	Inflammatory capacity of exosomes released in the early stages of acute pancreatitis predicts the severity of the disease. Journal of Pathology, 2022, 256, 83-92.	2.1	15
79	Thermospray high-performance liquid chromatography/mass spectrometric determination of cyclosporins. Rapid Communications in Mass Spectrometry, 1992, 6, 684-689.	0.7	14
80	Proteomic Analysis of Polypeptides Captured from Blood during Extracorporeal Albumin Dialysis in Patients with Cholestasis and Resistant Pruritus. PLoS ONE, 2011, 6, e21850.	1.1	14
81	The Exposed Proteomes of Brachyspira hyodysenteriae and B. pilosicoli. Frontiers in Microbiology, 2016, 7, 1103.	1.5	14
82	On the origin of some controversial ions (m/z 59, 60, 77, and 119) in the thermospray reagent plasma from ammonium acetate. Journal of the American Society for Mass Spectrometry, 1994, 5, 186-193.	1.2	13
83	Identification of the Autoantigen HB as the Barrier-to-Autointegration Factor. Journal of Biological Chemistry, 2003, 278, 50641-50644.	1.6	13
84	LymPHOS: Design of a phosphosite database of primary human T cells. Proteomics, 2009, 9, 3741-3751.	1.3	13
85	Discovery of lipoprotein lipase pl isoforms and contributions to their characterization. Journal of Proteomics, 2009, 72, 1031-1039.	1.2	13
86	Large-Scale Filter-Aided Sample Preparation Method for the Analysis of the Ubiquitinome. Analytical Chemistry, 2017, 89, 3840-3846.	3.2	13
87	Discovery of large molecules as new biomarkers in wastewater using environmental proteomics and suitable polymer probes. Science of the Total Environment, 2020, 747, 141145.	3.9	13
88	On-line preconcentration microliquid chromatography tandem mass spectrometric method for bradykinin analysis in plasma. Journal of Separation Science, 2001, 13, 265-274.	1.0	12
89	Phosphoproteomic Analysis of Platelets in Severe Obesity Uncovers Platelet Reactivity and Signaling Pathways Alterations. Arteriosclerosis, Thrombosis, and Vascular Biology, 2021, 41, 478-490.	1.1	12
90	Application of thermospray/high-performance liquid chromatography/mass spectrometry to the identification of glutathione conjugates derived from bioactive epoxides. Biological Mass Spectrometry, 1988, 16, 339-344.	0.5	11

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91	Detection and isolation of pentachlorophenol in oil samples associated with the Spanish toxic oil syndrome. Bulletin of Environmental Contamination and Toxicology, 1990, 45, 181-188.	1.3	11
92	The primary structure of a chondrichthyan protamine: A new apparent contradiction in protamine evolution. Journal of Molecular Evolution, 1996, 43, 528-535.	0.8	11
93	A collection of open source applications for mass spectrometry data mining. Proteomics, 2014, 14, 2275-2279.	1.3	11
94	Brachyspira hyodysenteriae and B. pilosicoli Proteins Recognized by Sera of Challenged Pigs. Frontiers in Microbiology, 2017, 8, 723.	1.5	11
95	Mitochondrial Proteome of Affected Glutamatergic Neurons in a Mouse Model of Leigh Syndrome. Frontiers in Cell and Developmental Biology, 2020, 8, 660.	1.8	11
96	LC-MS ion maps for the characterization of aniline derivatives of fatty acids and triglycerides in laboratory-denatured rapeseed oil. Journal of Mass Spectrometry, 2007, 42, 527-541.	0.7	10
97	Synthesis of N-[ring-G-3H]phenyllinoleamide and N-phenyl[1-14C]linoleamide as labelled standards for spanish toxic oil syndrome studies. Journal of Labelled Compounds and Radiopharmaceuticals, 1986, 23, 1029-1033.	0.5	9
98	Determination of oxidation products of N-phenyllinoleamide: Spanish toxic oil syndrome studies. Biomedical Applications, 1988, 426, 83-91.	1.7	9
99	Gas chromatographic/mass spectrometric analysis of high-performance liquid chromatographic fractions reflecting arachidonic acid metabolism in mouse peritoneal macrophages. Biological Mass Spectrometry, 1992, 21, 69-79.	0.5	9
100	Absorption and effects of 3-(N-phenylamino)-1,2-propanediol esters in relation to toxic oil syndrome. Lipids, 2001, 36, 1125-1133.	0.7	9
101	On the Generation and Outcome of 3-(N-Phenylamino)propane-1,2-diol Derivatives in Deodorized Model Oils Related to Toxic Oil Syndrome. Chemical Research in Toxicology, 2005, 18, 665-674.	1.7	9
102	Proteotyping of human haptoglobin by MALDI-TOF profiling: Phenotype distribution in a population of toxic oil syndrome patients. Proteomics, 2006, 6, S272-S281.	1.3	9
103	Application of Proteomic Tools To Detect the Nonspecificity of a Polyclonal Antibody against Lipoprotein Lipase. Journal of Proteome Research, 2008, 7, 4173-4177.	1.8	9
104	A multicentric study to evaluate the use of relative retention times in targeted proteomics. Journal of Proteomics, 2017, 152, 138-149.	1.2	9
105	Effects of sample matrix and high performance liquid chromatography eluent composition on the thermospray response for polar compounds. Rapid Communications in Mass Spectrometry, 1988, 2, 232-235.	0.7	8
106	Mass Spectrometric Determination of the Cleavage Sites in Escherichia coli Dihydroorotase Induced by a Cysteine-specific Reagent. Journal of Biological Chemistry, 1997, 272, 26934-26939.	1.6	8
107	A comprehensive Candida albicans PeptideAtlas build enables deep proteome coverage. Journal of Proteomics, 2016, 131, 122-130.	1.2	8
108	Characteristic fragmentation of thromboxane B2 in thermospray high-performance liquid chromatographyâ€"mass spectrometry. Biomedical Applications, 1991, 562, 153-168.	1.7	7

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109	Metabolism of N-phenyllinoleamide by rat liver. Biomedical Applications, 1993, 615, 191-196.	1.7	7
110	Products formed from the in vitro reaction of metabolites of 3-aminochrysene with calf thymus DNA. Chemico-Biological Interactions, 1993, 86, 1-15.	1.7	7
111	Thermospray and electrospray mass spectrometry of flavocoenzymes. Analysis of riboflavin sulphates from sugar beet. Analytica Chimica Acta, 1995, 302, 215-223.	2.6	7
112	Automated Strong Cation Exchange Extraction of Fatty Acid Esters of 3-(N-Phenylamino)-1,2-propanediol from Oil Samples for Routine Quantification by HPLC-APCI/MS/MS. Journal of Agricultural and Food Chemistry, 2001, 49, 5085-5091.	2.4	7
113	Analysis of the HLA class I associated peptide repertoire in a hepatocellular carcinoma cell line reveals tumor-specific peptides as putative targets for immunotherapy. Proteomics - Clinical Applications, 2007, 1, 286-298.	0.8	7
114	Platelet membrane lipid rafts protein composition varies following GPVI and CLEC-2 receptors activation. Journal of Proteomics, 2019, 195, 88-97.	1.2	6
115	Manganese-induced neurotoxicity in cerebellar granule neurons due to perturbation of cell network pathways with potential implications for neurodegenerative disorders. Metallomics, 2020, 12, 1656-1678.	1.0	6
116	Spontaneous changes in brain striatal dopamine synthesis and storage dynamics ex vivo reveal end-product feedback-inhibition of tyrosine hydroxylase. Neuropharmacology, 2022, 212, 109058.	2.0	6
117	Use of methyl oxime derivatives to enhance structural information in thermospray high-performance liquid chromatography-mass spectrometry. Journal of Chromatography A, 1991, 554, 155-173.	1.8	5
118	High-performance liquid chromatography-thermospray mass spectrometry of gibberellins. Journal of Chromatography A, 1992, 603, 157-164.	1.8	5
119	Mass spectrometric identification of N-phenyllinoleamide metabolites in mouse peritoneal macrophages. Rapid Communications in Mass Spectrometry, 1995, 9, 753-760.	0.7	5
120	Isolation of HLA-DR-naturally presented peptides identifies T-cell epitopes for rheumatoid arthritis. Annals of the Rheumatic Diseases, 2022, , annrheumdis-2021-220371.	0.5	5
121	Epoxidation of 6,7- and 10,11-oxidosqualenes by the squalene epoxidase present in rat liver microsomes. Bioorganic and Medicinal Chemistry Letters, 1993, 3, 2581-2586.	1.0	4
122	Proteomics of toxic oil syndrome in humans: Phenotype distribution in a population of patients. Chemico-Biological Interactions, 2011, 192, 129-135.	1.7	4
123	Lipoprotein lipase isoelectric point isoforms in humans. Biochemical and Biophysical Research Communications, 2014, 445, 480-485.	1.0	4
124	Isoform-speciff quantiffation of endothelins in HUVEC culture supernatants by on-line high-performance liquid chromatography/electrospray mass spectrometry. Biomedical Chromatography, 2004, 18, 388-395.	0.8	3
125	Determination of pesticide epitopic density in protein immunoconjugates by electrospray mass spectrometry. International Journal of Mass Spectrometry and Ion Processes, 1997, 160, 395-407.	1.9	2
126	Capillary Separations. , 2004, 251, 143-164.		2

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127	Chapter 9 LC-MS. II: Applications for pesticide food analysis. Comprehensive Analytical Chemistry, 2005, , 403-437.	0.7	2
128	Prostaglandin levels in infertile patients affected by asthenozoospermia and prostatitis. Prostaglandins Leukotrienes and Essential Fatty Acids, 1988, 31, 41-44.	1.0	1
129	High-performance liquid chromatography/thermospray mass spectrometry of some prostaglandins of the F series. Journal of Mass Spectrometry, 1995, 30, 608-616.	0.7	1
130	P2P proteomics – data sharing for enhanced protein identification. Automated Experimentation, 2012, 4, 1.	2.0	1
131	TCellXTalk facilitates the detection of co-modified peptides for the study of protein post-translational modification cross-talk in T cells. Bioinformatics, 2019, 35, 1404-1413.	1.8	1
132	The Primary Structure of a Chondrichthyan Protamine: A New Apparent Contradiction in Protamine Evolution. Journal of Molecular Evolution, 1996, 43, 528-535.	0.8	0