Anna Maria Timperio

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

91 2,260 25 44 g-index

102 2,574 4.1 4.65 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
91	Detection and Comparison of Bioactive Compounds in Different Extracts of Two Hazelnut Skin Varieties, Tonda Gentile Romana and Tonda Di Giffoni, Using a Metabolomics Approach. Metabolites, 2021, 11,	5.6	4
90	PGE2 Released by Pancreatic Cancer Cells Undergoing ER Stress Transfers the Stress to DCs Impairing Their Immune Function. <i>Molecular Cancer Therapeutics</i> , 2021 , 20, 934-945	6.1	5
89	The Potential of Lisosan G as a Possible Treatment for Glaucoma. <i>Frontiers in Pharmacology</i> , 2021 , 12, 719951	5.6	1
88	Specific adaptations are selected in opposite sun exposed Antarctic cryptoendolithic communities as revealed by untargeted metabolomics. <i>PLoS ONE</i> , 2020 , 15, e0233805	3.7	7
87	Multidisciplinary characterization of melanin pigments from the black fungus Cryomyces antarcticus. <i>Applied Microbiology and Biotechnology</i> , 2020 , 104, 6385-6395	5.7	16
86	Meteorite-Assisted Phosphorylation of Adenosine Under Proton Irradiation Conditions. <i>ChemSystemsChem</i> , 2020 , 2, e1900039	3.1	4
85	Urine Metabolome during Parturition. <i>Metabolites</i> , 2020 , 10,	5.6	2
84	Docosahexaenoic Acid Reverted the All- Retinoic Acid-Induced Cellular Proliferation of T24 Bladder Cancer Cell Line. <i>Journal of Clinical Medicine</i> , 2020 , 9,	5.1	1
83	Brain protein changes in Mecp2 mouse mutant models: Effects on disease progression of Mecp2 brain specific gene reactivation. <i>Journal of Proteomics</i> , 2020 , 210, 103537	3.9	4
82	GLYI4 Plays A Role in Methylglyoxal Detoxification and Jasmonate-Mediated Stress Responses in. <i>Biomolecules</i> , 2019 , 9,	5.9	9
81	Tartary buckwheat malt as ingredient of gluten-free cookies. <i>Journal of Cereal Science</i> , 2018 , 80, 37-43	3.8	37
80	Proteomic analysis of the Rett syndrome experimental model mecp2 mutant zebrafish. <i>Journal of Proteomics</i> , 2017 , 154, 128-133	3.9	10
79	Nonenzymatic Oligomerization of 3RSRCyclic CMP Induced by Proton and UV Irradiation Hints at a Nonfastidious Origin of RNA. <i>ChemBioChem</i> , 2017 , 18, 1535-1543	3.8	12
78	Evaluation of MALDI-TOF mass spectrometry and MALDI BioTyper in comparison to 16S rDNA sequencing for the identification of bacteria isolated from Arctic sea water. <i>PLoS ONE</i> , 2017 , 12, e01818	8ହୁଣ୍ଡ	37
77	Persistent Unresolved Inflammation in the -308 Female Mutated Mouse Model of Rett Syndrome. <i>Mediators of Inflammation</i> , 2017 , 2017, 9467819	4.3	12
76	Proton irradiation: a key to the challenge of N-glycosidic bond formation in a prebiotic context. <i>Scientific Reports</i> , 2017 , 7, 14709	4.9	24
75	Erectile dysfunction and diabetes: Association with the impairment of lipid metabolism and oxidative stress. <i>Clinical Biochemistry</i> , 2016 , 49, 70-8	3.5	6

74	Non-Enzymatic Oligomerization of 3R 5RCyclic AMP. PLoS ONE, 2016, 11, e0165723	3.7	10
73	Expression and oxidative modifications of plasma proteins in autism spectrum disorders: Interplay between inflammatory response and lipid peroxidation. <i>Proteomics - Clinical Applications</i> , 2016 , 10, 110	3 ³ 1 ¹ 112	25
72	Proteome and metabolome profiling of wild-type and YCA1-knock-out yeast cells during acetic acid-induced programmed cell death. <i>Journal of Proteomics</i> , 2015 , 128, 173-88	3.9	21
71	Unraveling the seed endosperm proteome of the lotus (Nelumbo nucifera Gaertn.) utilizing 1DE and 2DE separation in conjunction with tandem mass spectrometry. <i>Proteomics</i> , 2015 , 15, 1717-35	4.8	6
70	Differential proteome-metabolome profiling of YCA1-knock-out and wild type cells reveals novel metabolic pathways and cellular processes dependent on the yeast metacaspase. <i>Molecular BioSystems</i> , 2015 , 11, 1573-83		6
69	Proteomic and metabolic profiles of Cakile maritima Scop. Sea Rocket grown in the presence of cadmium. <i>Molecular BioSystems</i> , 2015 , 11, 1096-109		11
68	Analysis of the mitochondrial proteome of cybrid cells harbouring a truncative mitochondrial DNA mutation in respiratory complex I. <i>Molecular BioSystems</i> , 2014 , 10, 1313-9		7
67	Hsp10 nuclear localization and changes in lung cells response to cigarette smoke suggest novel roles for this chaperonin. <i>Open Biology</i> , 2014 , 4,	7	11
66	Na+/K+-ATPase II-subunit is recruited in Na-K-2Cl co-transporter isoform 2 multiprotein complexes in rat kidneys: possible role in blood pressure regulation. <i>Journal of Hypertension</i> , 2014 , 32, 1842-53	1.9	5
65	Systems biology: a new tool for farm animal science. Current Protein and Peptide Science, 2014, 15, 100-	17 .8	16
64	Cadmium stress responses in Brassica juncea: hints from proteomics and metabolomics. <i>Journal of Proteome Research</i> , 2013 , 12, 4979-97	5.6	72
63	Crosstalk between salicylic acid and jasmonate in Arabidopsis investigated by an integrated proteomic and transcriptomic approach. <i>Molecular BioSystems</i> , 2013 , 9, 1169-87		40
62	Biomarker discovery and applications for foods and beverages: proteomics to nanoproteomics. Journal of Proteomics, 2013 , 93, 74-92	3.9	36
61	The mitochondrial Italian Human Proteome Project initiative (mt-HPP). <i>Molecular BioSystems</i> , 2013 , 9, 1984-92		8
60	Analysis of TAp73-dependent signaling via omics technologies. <i>Journal of Proteome Research</i> , 2013 , 12, 4207-20	5.6	16
59	Red Blood Cell Lipidomics analysis through HPLC-ESI-qTOF: application to red blood cell storage. Journal of Integrated OMICS, 2013, 3,	0.5	2
58	Production of the phytoalexins trans-resveratrol and delta-viniferin in two economy-relevant grape cultivars upon infection with Botrytis cinerea in field conditions. <i>Plant Physiology and Biochemistry</i> , 2012 , 50, 65-71	5.4	33
57	Acclimation to intense light implies changes at the level of trimeric subunits involved in the structural organization of the main light-harvesting complex of photosystem II (LHCII) and their isoforms. <i>Plant Physiology and Biochemistry</i> , 2012 , 50, 8-14	5.4	7

56	Murine macrophages response to iron. <i>Journal of Proteomics</i> , 2012 , 76 Spec No., 10-27	3.9	20
55	A monoclonal antibody for the CD45 receptor in the teleost fish Dicentrarchus labrax. <i>Developmental and Comparative Immunology</i> , 2012 , 37, 342-53	3.2	6
54	Identification of moesin as NKCC2-interacting protein and analysis of its functional role in the NKCC2 apical trafficking. <i>Biology of the Cell</i> , 2012 , 104, 658-76	3.5	10
53	Analysis of the cattle liver proteome by high-sensitive liquid chromatography coupled with mass spectrometry method. <i>Methods in Molecular Biology</i> , 2012 , 909, 43-62	1.4	4
52	Combinatorial peptide ligand libraries to discover liver disease biomarkers in plasma samples. <i>Methods in Molecular Biology</i> , 2012 , 909, 311-9	1.4	1
51	Clinical metabolomics: the next stage of clinical biochemistry. <i>Blood Transfusion</i> , 2012 , 10 Suppl 2, s19-2	2 4 .6	19
50	Rat liver mitochondrial proteome: changes associated with aging and acetyl-L-carnitine treatment. <i>Journal of Proteomics</i> , 2011 , 74, 2536-47	3.9	26
49	Docosohaexanoic acid-supplemented PACA44 cell lines and over-activation of Krebs cycle: an integrated proteomic, metabolomic and interactomic overview. <i>Journal of Proteomics</i> , 2011 , 74, 2138-5	8 ^{3.9}	14
48	Vascular endothelial growth factor up-regulation in the mouse hippocampus and its role in the control of epileptiform activity. <i>European Journal of Neuroscience</i> , 2011 , 33, 482-98	3.5	18
47	Female urinary proteomics: New insight into exogenous and physiological hormone-dependent changes. <i>Proteomics - Clinical Applications</i> , 2011 , 5, 343-53	3.1	13
46	Comparative proteomic analysis of hemocyanins in Dinocras cephalotes and Perla marginata (Plecoptera). <i>Environmental Entomology</i> , 2011 , 40, 167-71	2.1	3
45	Fibroblasts from PS1 mutated pre-symptomatic subjects and Alzheimerß disease patients share a unique protein levels profile. <i>Journal of Alzheimerß Disease</i> , 2010 , 21, 431-44	4.3	4
44	Proteomics and transcriptomics investigation on longissimus muscles in Large White and Casertana pig breeds. <i>Journal of Proteome Research</i> , 2010 , 9, 6450-66	5.6	49
43	Recombinant clotting factor VIII concentrates: Heterogeneity and high-purity evaluation. <i>Electrophoresis</i> , 2010 , 31, 2730-9	3.6	14
42	Comparison among plasma-derived clotting factor VIII by using monodimensional gel electrophoresis and mass spectrometry. <i>Blood Transfusion</i> , 2010 , 8 Suppl 3, s98-104	3.6	1
41	Comparison of milk fat globule membrane (MFGM) proteins of Chianina and Holstein cattle breed milk samples through proteomics methods. <i>Nutrients</i> , 2009 , 1, 302-15	6.7	25
40	Accumulation of overoxidized Peroxiredoxin III in aged rat liver mitochondria. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2009 , 1787, 890-6	4.6	51
39	Functional effects of somatostatin receptor 1 activation on synaptic transmission in the mouse hippocampus. <i>Journal of Neurochemistry</i> , 2009 , 111, 1466-77	6	11

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38	Comparative proteomics and transcriptomics analyses of livers from two different Bos taurus breeds: "Chianina and Holstein Friesian". <i>Journal of Proteomics</i> , 2009 , 73, 309-22	3.9	35
37	High resolution preparation of monocyte-derived macrophages (MDM) protein fractions for clinical proteomics. <i>Proteome Science</i> , 2009 , 7, 4	2.6	8
36	Proteomic analysis of multiprotein complexes in the thylakoid membrane upon cadmium treatment. <i>Journal of Proteome Research</i> , 2009 , 8, 310-26	5.6	77
35	Liquid-chromatography-mass spectrometry of thylakoid membrane proteins. <i>Methods in Molecular Biology</i> , 2009 , 492, 113-30	1.4	3
34	Modulation of the neuronal response to ischaemia by somatostatin analogues in wild-type and knock-out mouse retinas. <i>Journal of Neurochemistry</i> , 2008 , 106, 2224-35	6	43
33	Coupling of native liquid phase isoelectrofocusing and blue native polyacrylamide gel electrophoresis: a potent tool for native membrane multiprotein complex separation. <i>Journal of Proteome Research</i> , 2008 , 7, 1326-40	5.6	32
32	Induction of apoptosis in Jeko-1 mantle cell lymphoma cell line by resveratrol: a proteomic analysis. Journal of Proteome Research, 2008 , 7, 2670-80	5.6	19
31	Proteomics applied on plant abiotic stresses: role of heat shock proteins (HSP). <i>Journal of Proteomics</i> , 2008 , 71, 391-411	3.9	350
30	A proteomic approach for evaluating the cell response to a novel histone deacetylase inhibitor in colon cancer cells. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2008 , 1784, 1702-10	4	12
29	Exploring the platelet proteome via combinatorial, hexapeptide ligand libraries. <i>Journal of Proteome Research</i> , 2007 , 6, 4290-303	5.6	79
28	Chemically enhanced liquid chromatography/tandem mass spectrometry determination of glutamic acid in the diffusion medium of retinal cells. <i>Biomedical Chromatography</i> , 2007 , 21, 1069-76	1.7	25
27	Proteomic analysis of photosystem I components from different plant species. <i>Proteomics</i> , 2007 , 7, 1860	64786	16
26	Changes in neuronal response to ischemia in retinas with genetic alterations of somatostatin receptor expression. <i>European Journal of Neuroscience</i> , 2007 , 25, 1447-59	3.5	41
25	Proteomics, pigment composition, and organization of thylakoid membranes in iron-deficient spinach leaves. <i>Journal of Experimental Botany</i> , 2007 , 58, 3695-710	7	90
24	Assay of ochratoxin A in grape by high-pressure liquid chromatography coupled on line with an ESI-mass spectrometry. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2006 , 832, 127-33	3.2	23
23	Involvement of active oxygen species in protein and oligonucleotide degradation induced by nitrofurans. <i>Biochemistry and Cell Biology</i> , 2005 , 83, 166-75	3.6	14
22	Hydrazide derivatives produce active oxygen species as hydrazine. <i>Bioorganic Chemistry</i> , 2005 , 33, 459-6	9.1	9
21	Fingerprinting of Antenna Proteins of Photosystem I by Reversed Phase High Performance Liquid Chromatography. <i>Chromatographia</i> , 2005 , 61, 1-7	2.1	1

20	Investigation of the lateral light-induced migration of photosystem II light-harvesting proteins by nano-high performance liquid chromatography electrospray ionization mass spectrometry. <i>Journal of Biological Chemistry</i> , 2005 , 280, 28858-66	5.4	11
19	Separation and identification of the light harvesting proteins contained in grana and stroma thylakoid membrane fractions. <i>Journal of Chromatography A</i> , 2004 , 1040, 73-81	4.5	13
18	Multidimensional proteomic analysis of photosynthetic membrane proteins by liquid extraction-ultracentrifugation-liquid chromatography-mass spectrometry. <i>Proteomics</i> , 2004 , 4, 3909-20	4.8	44
17	Intact mass measurements for unequivocal identification of hydrophobic photosynthetic photosystems I and II antenna proteins. <i>Electrophoresis</i> , 2004 , 25, 1353-66	3.6	13
16	Separation and identification of photosynthetic antenna membrane proteins by high- performance liquid chromatography electrospray ionization mass spectrometry. <i>European Journal of Mass Spectrometry</i> , 2004 , 10, 321-33	1.1	3
15	Identification of a furazolidone metabolite responsible for the inhibition of amino oxidases. <i>Xenobiotica</i> , 2003 , 33, 153-67	2	21
14	Characterization of a variant of the spinach PSII type I light-harvesting protein using kinetically controlled digestion and RP-HPLC-ESI-MS. <i>Analytical Chemistry</i> , 2003 , 75, 6775-80	7.8	17
13	Proteomics of light-harvesting proteins in different plant species. Analysis and comparison by liquid chromatography-electrospray ionization mass spectrometry. Photosystem II. <i>Plant Physiology</i> , 2003 , 131, 198-214	6.6	49
12	Proteomics of light-harvesting proteins in different plant species. Analysis and comparison by liquid chromatography-electrospray ionization mass spectrometry. Photosystem I. <i>Plant Physiology</i> , 2002 , 130, 1938-50	6.6	39
11	Isoforms of photosystem II antenna proteins in different plant species revealed by liquid chromatography-electrospray ionization mass spectrometry. <i>Journal of Biological Chemistry</i> , 2001 , 276, 45755-61	5.4	29
10	High-performance liquid chromatography-electrospray ionization mass spectrometry using monolithic capillary columns for proteomic studies. <i>Analytical Chemistry</i> , 2001 , 73, 2390-6	7.8	194
9	High performance liquid chromatography-electrospray mass spectrometry for the simultaneous resolution and identification of intrinsic thylakoid membrane proteins. <i>Proteins: Structure, Function and Bioinformatics</i> , 2000 , 41, 398-406	4.2	21
8	Resolution and identification of the protein components of the photosystem II antenna system of higher plants by reversed-phase liquid chromatography with electrospray-mass spectrometric detection. <i>Journal of Chromatography A</i> , 2000 , 886, 111-21	4.5	46
7	Metal binding to Pseudomonas aeruginosa azurin: a kinetic investigation. <i>Zeitschrift Fur Naturforschung - Section C Journal of Biosciences</i> , 2000 , 55, 347-54	1.7	5
6	Isolation and characterization of chloroplast Photosystem II antenna of spinach by reversed-phase liquid chromatography. <i>Photosynthesis Research</i> , 1999 , 61, 281-290	3.7	16
5	Rapid resolution by reversed-phase high-performance liquid chromatography of the thylakoid membrane proteins of the photosystem II light-harvesting complex. <i>Journal of Chromatography A</i> , 1997 , 779, 131-138	4.5	13
4	Random amplified polymorphic DNA (RAPD) interpretation requires a sensitive method for the detection of amplified DNA. <i>Electrophoresis</i> , 1996 , 17, 1553-4	3.6	15
3	Capillary electrophoresis of closely related intrinsic thylakoid membrane proteins of the photosystem II light-harvesting complex (LHC II). <i>Electrophoresis</i> , 1996 , 17, 1597-601	3.6	10

2 Metabolic responses in opposite sun-exposed Antarctic cryptoendolithic communities

2

From Targeted Quantification to Untargeted Metabolomics

1