Vincenzo Cunsolo

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A new monomeric α-amylase inhibitor from the tetraploid emmer wheat is mostly active against stored product pests. Journal of Pest Science, 2022, 95, 1401-1412.	3.7	3
2	Meta-proteomic analysis of two mammoth's trunks by EVA technology and high-resolution mass spectrometry for an indirect picture of their habitat and the characterization of the collagen type I, alpha-1 and alpha-2 sequence. Amino Acids, 2022, , .	2.7	4
3	Physiactisome: A New Nanovesicle Drug Containing Heat Shock Protein 60 for Treating Muscle Wasting and Cachexia. Cells, 2022, 11, 1406.	4.1	4
4	Paleoproteomic profiling of organic residues on prehistoric pottery from Malta. Amino Acids, 2021, 53, 295-312.	2.7	18
5	Quantitative Label-Free Comparison of the Metabolic Protein Fraction in Old and Modern Italian Wheat Genotypes by a Shotgun Approach. Molecules, 2021, 26, 2596.	3.8	9
6	Meta-proteomic analysis of the Shandrin mammoth by EVA technology and high-resolution mass spectrometry: what is its gut microbiota telling us?. Amino Acids, 2021, 53, 1507-1521.	2.7	5
7	VDACs Post-Translational Modifications Discovery by Mass Spectrometry: Impact on Their Hub Function. International Journal of Molecular Sciences, 2021, 22, 12833.	4.1	8
8	Qualitative proteomic comparison of metabolic and CM-like protein fractions in old and modern wheat Italian genotypes by a shotgun approach. Journal of Proteomics, 2020, 211, 103530.	2.4	16
9	Gluten proteome comparison among durum wheat genotypes with different release date. Journal of Cereal Science, 2020, 96, 103092.	3.7	12
10	Post-Translational Modification Analysis of VDAC1 in ALS-SOD1 Model Cells Reveals Specific Asparagine and Glutamine Deamidation. Antioxidants, 2020, 9, 1218.	5.1	10
11	Mass Spectrometry and 1H-NMR Study of Schinopsis lorentzii (Quebracho) Tannins as a Source of Hypoglycemic and Antioxidant Principles. Molecules, 2020, 25, 3257.	3.8	14
12	Identification of New Antimicrobial Peptides from Mediterranean Medical Plant Charybdis pancration (Steinh.) Speta. Antibiotics, 2020, 9, 747.	3.7	10
13	A High Resolution Mass Spectrometry Study Reveals the Potential of Disulfide Formation in Human Mitochondrial Voltage-Dependent Anion Selective Channel Isoforms (hVDACs). International Journal of Molecular Sciences, 2020, 21, 1468.	4.1	14
14	Dataset of the metabolic and CM-like protein fractions in old and modern wheat Italian genotypes. Data in Brief, 2019, 27, 104730.	1.0	2
15	Polymorphism at donkey l²-lactoglobulin II locus: identification and characterization of a new genetic variant with a very low expression. Amino Acids, 2018, 50, 735-746.	2.7	3
16	Post-translational modifications of VDAC1 and VDAC2 cysteines from rat liver mitochondria. Biochimica Et Biophysica Acta - Bioenergetics, 2018, 1859, 806-816.	1.0	32
17	Proteomic Analyses on an Ancient Egyptian Cheese and Biomolecular Evidence of Brucellosis. Analytical Chemistry, 2018, 90, 9673-9676.	6.5	38
18	High resolution mass spectrometry characterization of the oxidation pattern of methionine and cysteine residues in rat liver mitochondria voltage-dependent anion selective channel 3 (VDAC3). Biochimica Et Biophysica Acta - Biomembranes, 2017, 1859, 301-311.	2.6	29

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19	Proteins and bioactive peptides from donkey milk: The molecular basis for its reduced allergenic properties. Food Research International, 2017, 99, 41-57.	6.2	55
20	Polyphemus, Odysseus and the ovine milk proteome. Journal of Proteomics, 2017, 152, 58-74.	2.4	14
21	α-Glucosidase inhibition and antioxidant activity of an oenological commercial tannin. Extraction, fractionation and analysis by HPLC/ESI-MS/MS and 1H NMR. Food Chemistry, 2017, 215, 50-60.	8.2	54
22	Site-specific glycosylation of donkey milk lactoferrin investigated by high-resolution mass spectrometry. Amino Acids, 2016, 48, 2799-2808.	2.7	10
23	Sequence characterization and glycosylation sites identification of donkey milk lactoferrin by multiple enzyme digestions and mass spectrometry. Amino Acids, 2016, 48, 1569-1580.	2.7	9
24	Zeus, Aesculapius, Amalthea and the proteome of goat milk. Journal of Proteomics, 2015, 128, 69-82.	2.4	28
25	Mass spectrometry in food proteomics: a tutorial. Journal of Mass Spectrometry, 2014, 49, 768-784.	1.6	56
26	Involvement of alanine racemase in germination of Bacillus cereus spores lacking an intact exosporium. Archives of Microbiology, 2014, 196, 79-85.	2.2	8
27	Immune mediators of sea-cucumber Holothuria tubulosa (Echinodermata) as source of novel antimicrobial and anti-staphylococcal biofilm agents. AMB Express, 2013, 3, 35.	3.0	56
28	MALDIâ€TOF mass spectrometry for the monitoring of sheâ€donkey's milk contamination or adulteration. Journal of Mass Spectrometry, 2013, 48, 148-153.	1.6	28
29	Root Protein Profiles of Two Citrus Rootstocks Grown under Iron Sufficiency/Deficiency Conditions. European Journal of Mass Spectrometry, 2013, 19, 305-324.	1.0	9
30	MSâ€based characterization of <i>α</i> _{<i>s</i>2} â€casein isoforms in donkey's milk. Journal of Mass Spectrometry, 2012, 47, 1150-1159.	1.6	19
31	Applications of liquid chromatography–mass spectrometry for food analysis. Journal of Chromatography A, 2012, 1259, 74-85.	3.7	172
32	Mass spectrometry in the proteome analysis of mature cereal kernels. Mass Spectrometry Reviews, 2012, 31, 448-465.	5.4	35
33	High Molecular Weight Glutenin Subunits in Some Durum Wheat Cultivars Investigated by Means of Mass Spectrometric Techniques. Journal of Agricultural and Food Chemistry, 2011, 59, 12226-12237.	5.2	24
34	Applications of Mass Spectrometry Techniques in the Investigation of Milk Proteome. European Journal of Mass Spectrometry, 2011, 17, 305-320.	1.0	47
35	Poppea's bath liquor: The secret proteome of she-donkey's milk. Journal of Proteomics, 2011, 74, 2083-2099.	2.4	40
36	Molecular and functional characterization of VDAC2 purified from mammal spermatozoa. Bioscience Reports, 2009, 29, 351-362.	2.4	56

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37	Sequence determination of α _{<i>s</i>1} asein isoforms from donkey by mass spectrometric methods. Journal of Mass Spectrometry, 2009, 44, 1742-1753.	1.6	29
38	Sequence and phosphorylation level determination of two donkey <i>β</i> aseins by mass spectrometry. Rapid Communications in Mass Spectrometry, 2009, 23, 1907-1916.	1.5	28
39	Proteome analysis of Citrus sinensis L. (Osbeck) flesh at ripening time. Journal of Proteomics, 2009, 73, 134-152.	2.4	75
40	Donkeys' milk protein fraction investigated by electrophoretic methods and mass spectrometric analysis. International Dairy Journal, 2009, 19, 190-197.	3.0	54
41	Characterization of the protein profile of donkey's milk whey fraction. Journal of Mass Spectrometry, 2007, 42, 1162-1174.	1.6	46
42	Detection and sequence determination of a new variant ^{î2} -lactoglobulin II from donkey. Rapid Communications in Mass Spectrometry, 2007, 21, 1438-1446.	1.5	24
43	Characterization of B- and C-type low molecular weight glutenin subunits by electrospray ionization mass spectrometry and matrix-assisted laser desorption/ ionization mass spectrometry. Proteomics, 2005, 5, 719-728.	2.2	46
44	Detection and characterization by high-performance liquid chromatography and mass spectrometry of a goatl ² -casein associated with a CSN2 null allele. Rapid Communications in Mass Spectrometry, 2005, 19, 2943-2949.	1.5	22
45	NsLTP1 and NsLTP2 Isoforms in Soft Wheat (Triticum aestivumCv. Centauro) and Farro (Triticum) Tj ETQq1 1 0.	.784314 rg	BT 10verlock
46	Identification and characterization of a newl ² -casein variant in goat milk by high-performance liquid chromatography with electrospray ionization mass spectrometry and matrix-assisted laser desorption/ionization mass spectrometry. Rapid Communications in Mass Spectrometry, 2004, 18, 1972-1982.	1.5	33
47	Structural studies of the allelic wheat glutenin subunits 1Bx7 and 1Bx20 by matrix-assisted laser desorption/ionization mass spectrometry and high-performance liquid chromatography/electrospray ionization mass spectrometry. Journal of Mass Spectrometry, 2004, 39, 66-78.	1.6	48
48	Mass Spectrometry in the Characterisation of Cereal Seed Proteins. European Journal of Mass Spectrometry, 2004, 10, 359-370.	1.0	26
49	Structural studies of glutenin subunits 1Dy10 and 1Dy12 by matrix-assisted laser desorption/ionisation mass spectrometry and high-performance liquid chromatography/electrospray ionisation mass spectrometry. Rapid Communications in Mass Spectrometry, 2003, 17, 442-454.	1.5	47
50	Investigation and correction of the gene-derived sequence of glutenin subunit 1Dx2 by matrix-assisted laser desorption/ionisation mass spectrometry. Rapid Communications in Mass Spectrometry, 2002, 16, 1911-1918.	1.5	26