

Luis Javier González

List of Publications by Year in descending order

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63
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1,331
citations

361045

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h-index

414034

32
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67
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67
docs citations

67
times ranked

1658
citing authors

#	ARTICLE	IF	CITATIONS
1	A COVID-19 vaccine candidate composed of the SARS-CoV-2 RBD dimer and <i>Neisseria meningitidis</i> outer membrane vesicles. <i>RSC Chemical Biology</i> , 2022, 3, 242-249.	2.0	15
2	Chimeric Antigen by the Fusion of SARS-CoV-2 Receptor Binding Domain with the Extracellular Domain of Human CD154: A Promising Improved Vaccine Candidate. <i>Vaccines</i> , 2022, 10, 897.	2.1	5
3	Mass spectrometric and kinetics characterization of modified species of Growth Hormone Releasing Hexapeptide generated under thermal stress in different pH and buffers. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2021, 194, 113776.	1.4	1
4	SARS-CoV-2 RBD-Tetanus Toxoid Conjugate Vaccine Induces a Strong Neutralizing Immunity in Preclinical Studies. <i>ACS Chemical Biology</i> , 2021, 16, 1223-1233.	1.6	57
5	Synthesis, LC-MS/MS analysis, and biological evaluation of two vaccine candidates against ticks based on the antigenic PO peptide from <i>R. sanguineus</i> linked to the p64K carrier protein from <i>Neisseria meningitidis</i> . <i>Analytical and Bioanalytical Chemistry</i> , 2021, 413, 5885-5900.	1.9	3
6	In-solution buffer-free digestion allows full-sequence coverage and complete characterization of post-translational modifications of the receptor-binding domain of SARS-CoV-2 in a single ESI-MS spectrum. <i>Analytical and Bioanalytical Chemistry</i> , 2021, 413, 7559-7585.	1.9	11
7	Functional and Mass Spectrometric Evaluation of an Anti-Tick Antigen Based on the PO Peptide Conjugated to Bm86 Protein. <i>Pathogens</i> , 2020, 9, 513.	1.2	21
8	CIGB-814, an altered peptide ligand derived from human heat-shock protein 60, decreases anti-cyclic citrullinated peptides antibodies in patients with rheumatoid arthritis. <i>Clinical Rheumatology</i> , 2019, 38, 955-960.	1.0	21
9	An on-matrix digestion procedure for AP-MS experiments dissects the interplay between complex-conserved and serotype-specific reactivities in Dengue virus-human plasma interactome. <i>Journal of Proteomics</i> , 2019, 193, 71-84.	1.2	3
10	Growth hormone-releasing peptide 6 prevents cutaneous hypertrophic scarring: early mechanistic data from a proteome study. <i>International Wound Journal</i> , 2018, 15, 538-546.	1.3	5
11	Characterization of low-abundance species in the active pharmaceutical ingredient of CIGB-300: A clinical-grade anticancer synthetic peptide. <i>Journal of Peptide Science</i> , 2018, 24, e3081.	0.8	9
12	Phase I Clinical Trial with a Novel Altered Peptide Ligand Derived from Human Heat-Shock Protein 60 for Treatment of Rheumatoid Arthritis: Safety, Pharmacokinetics and Preliminary Therapeutic Effects. <i>Journal of Clinical Trials</i> , 2018, 08, .	0.1	10
13	Antimycobacterial Activity: A New Pharmacological Target for Conotoxins Found in the First Reported Conotoxin from <i>Conasprella ximenes</i> . <i>Toxins</i> , 2018, 10, 51.	1.5	16
14	A deeper mining on the protein composition of VA-MENGOC-BC®: An OMV-based vaccine against <i>N. meningitidis</i> serogroup B and C. <i>Human Vaccines and Immunotherapeutics</i> , 2017, 13, 2548-2560.	1.4	11
15	Development and validation of a bioanalytical method based on LC-MS/MS analysis for the quantitation of CIGB-814 peptide in plasma from Rheumatoid Arthritis patients. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2017, 143, 130-140.	1.4	9
16	Protein content of the <i>Hylesia metabus</i> egg nest setae (Cramer [1775]) (Lepidoptera: Saturniidae) and its association with the parental investment for the reproductive success and lepidopterism. <i>Journal of Proteomics</i> , 2017, 150, 183-200.	1.2	9
17	A New Member of Gamma-Conotoxin Family Isolated from <i>Conus princeps</i> Displays a Novel Molecular Target. <i>Toxins</i> , 2016, 8, 39.	1.5	7
18	Comparative proteomic analysis of growth hormone secretagogue A233 treatment of murine macrophage cells J774A.2 indicates it has a role in antiviral innate response. <i>Biochemistry and Biophysics Reports</i> , 2016, 5, 379-387.	0.7	7

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19	Comparative proteomics analysis of the antitumor effect of CIGB-552 peptide in HT-29 colon adenocarcinoma cells. <i>Journal of Proteomics</i> , 2015, 126, 163-171.	1.2	13
20	Structural characterization and biological implications of sulfated N-glycans in a serine protease from the neotropical moth <i>Hylesia metabus</i> (Cramer [1775]) (Lepidoptera: Saturniidae). <i>Glycobiology</i> , 2015, 26, cwv096.	1.3	18
21	Data for comparative proteomics analysis of the antitumor effect of CIGB-552 peptide in HT-29 colon adenocarcinoma cells. <i>Data in Brief</i> , 2015, 4, 468-473.	0.5	0
22	Bio-analytical method based on MALDI-MS analysis for the quantification of CIGB-300 anti-tumor peptide in human plasma. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2015, 105, 107-114.	1.4	5
23	Characterization of Protein Complexes using Targeted Proteomics. <i>Current Topics in Medicinal Chemistry</i> , 2014, 14, 344-350.	1.0	6
24	SCX charge state selective separation of tryptic peptides combined with 2D-RP-HPLC allows for detailed proteome mapping. <i>Journal of Proteomics</i> , 2013, 91, 164-171.	1.2	27
25	Computational proteomics pitfalls and challenges: Havana Bioinfo 2012 Workshop report. <i>Journal of Proteomics</i> , 2013, 87, 134-138.	1.2	19
26	Pharmacokinetic study of Growth Hormone-Releasing Peptide 6 (GHRP-6) in nine male healthy volunteers. <i>European Journal of Pharmaceutical Sciences</i> , 2013, 48, 40-46.	1.9	8
27	HI-Bone: A Scoring System for Identifying Phenylisothiocyanate-Derivatized Peptides Based on Precursor Mass and High Intensity Fragment Ions. <i>Analytical Chemistry</i> , 2013, 85, 3515-3520.	3.2	7
28	Periplasmic proteins encoded by VCA0261 and VC2216 genes together with copA and cueR products are required for copper tolerance but not for virulence in <i>Vibrio cholerae</i> . <i>Microbiology (United Kingdom)</i> , 2012, 158, 2005-2016.	0.7	25
29	Selective Isolation of Multiply Charged Peptides: A Confident Strategy for Protein Identification Using a Linear Trap Quadrupole Mass Spectrometer. <i>European Journal of Mass Spectrometry</i> , 2012, 18, 505-508.	0.5	0
30	Introducing an Asp-Pro Linker in the Synthesis of Random One-Bead-One-Compound Hexapeptide Libraries Compatible with ESI-MS Analysis. <i>ACS Combinatorial Science</i> , 2012, 14, 145-149.	3.8	7
31	Development and validation of a bioanalytical LC-MS method for the quantification of GHRP-6 in human plasma. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2012, 60, 19-25.	1.4	24
32	CIGB-300, a synthetic peptide-based drug that targets the CK2 phosphoacceptor domain. Translational and clinical research. <i>Molecular and Cellular Biochemistry</i> , 2011, 356, 45-50.	1.4	41
33	Peptide fractionation by acid pH SDS-free electrophoresis. <i>Electrophoresis</i> , 2011, 32, 1323-1326.	1.3	13
34	Selective Isolation and Detection of Two Different Positively Charged Peptides Groups by Strong Cation Exchange Chromatography and Matrix-Assisted Laser Desorption/Ionization Mass Spectrometry: Application to Proteomics Studies. <i>European Journal of Mass Spectrometry</i> , 2010, 16, 693-702.	0.5	4
35	Evaluation of Phenylthiocarbonyl-Derivatized Peptides by Electrospray Ionization Mass Spectrometry: Selective Isolation and Analysis of Modified Multiply Charged Peptides for Liquid Chromatography-Tandem Mass Spectrometry Experiments. <i>Analytical Chemistry</i> , 2010, 82, 8492-8501.	3.2	10
36	Proteomic Profile Regulated by the Anticancer Peptide CIGB-300 in Non-Small Cell Lung Cancer (NSCLC) Cells. <i>Journal of Proteome Research</i> , 2010, 9, 5473-5483.	1.8	26

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37	Proteomic study via a non-gel based approach of meningococcal outer membrane vesicle vaccine obtained from strain CU385: A road map for discovering new antigens. <i>Hum Vaccin</i> , 2009, 5, 347-356.	2.4	20
38	Anaerobic growth promotes synthesis of colonization factors encoded at the <i>Vibrio</i> pathogenicity island in <i>Vibrio cholerae</i> El Tor. <i>Research in Microbiology</i> , 2009, 160, 48-56.	1.0	27
39	CIGB-300, a novel proapoptotic peptide that impairs the CK2 phosphorylation and exhibits anticancer properties both in <i>in vitro</i> and <i>in vivo</i> . <i>Molecular and Cellular Biochemistry</i> , 2008, 316, 163-167.	1.4	86
40	Differential soluble protein expression between <i>Trichomonas vaginalis</i> isolates exhibiting low and high virulence phenotypes. <i>Journal of Proteomics</i> , 2008, 71, 109-122.	1.2	30
41	Proteomics Based on Peptide Fractionation by SDS-Free PAGE. <i>Journal of Proteome Research</i> , 2008, 7, 2427-2434.	1.8	18
42	Letter: Specific Isotope Labeling for the Identification of Free N-Terminal Peptides of Proteins Separated by Polyacrylamide Gel Electrophoresis. <i>European Journal of Mass Spectrometry</i> , 2007, 13, 307-309.	0.5	1
43	Application of two-dimensional electrophoresis and matrix-assisted laser desorption/ionization time-of-flight mass spectrometry for proteomic analysis of the sexually transmitted parasite <i>Trichomonas vaginalis</i> . <i>Journal of Mass Spectrometry</i> , 2007, 42, 1463-1473.	0.7	23
44	Double acylation for identification of amino-terminal peptides of proteins isolated by polyacrylamide gel electrophoresis. <i>Rapid Communications in Mass Spectrometry</i> , 2007, 21, 2237-2244.	0.7	6
45	Proteome analysis of <i>Leishmania (Viannia) braziliensis</i> by two-dimensional gel electrophoresis and mass spectrometry. <i>Molecular and Biochemical Parasitology</i> , 2007, 154, 6-21.	0.5	41
46	Increased antiviral activity of microscale-purified HuIFN γ 8 (human interferon γ 8) over HuIFN γ 2b in Hep-2 cells challenged with Mengo virus. <i>Biotechnology and Applied Biochemistry</i> , 2007, 48, 159.	1.4	4
47	Selective Isolation of Lysine-Free Tryptic Peptides Delimited by Arginine Residues: A New Tool for Proteome Analysis. <i>Journal of Proteome Research</i> , 2006, 5, 1204-1213.	1.8	10
48	Cloning, expression and growth promoting action of Red tilapia (<i>Oreochromis</i> sp.) neuropeptide Y. <i>Peptides</i> , 2006, 27, 710-718.	1.2	27
49	Selective isolation of multiple positively charged peptides for 2-DE-free quantitative proteomics. <i>Proteomics</i> , 2006, 6, 4444-4455.	1.3	18
50	A multivalent recombinant antibody fragment specific for carcinoembryonic antigen. <i>Biotechnology and Applied Biochemistry</i> , 2006, 43, 39.	1.4	17
51	SCAPE: A New Tool for the Selective CAPture of PEptides in Protein Identification. <i>Journal of Proteome Research</i> , 2005, 4, 491-496.	1.8	24
52	Effect of high concentration of Co (II) on <i>Enterobacter liquefaciens</i> strain C-1: A bacterium highly resistant to heavy metals with an unknown genome. <i>Proteomics</i> , 2004, 4, 1265-1279.	1.3	21
53	Automated interpretation of mass spectra of complex mixtures by matching of isotope peak distributions. <i>Rapid Communications in Mass Spectrometry</i> , 2004, 18, 2465-2472.	0.7	32
54	Isotopica: a tool for the calculation and viewing of complex isotopic envelopes. <i>Nucleic Acids Research</i> , 2004, 32, W674-W678.	6.5	27

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55	Identification of nuclear proteins of small cell lung cancer cell line H82: An improved procedure for the analysis of silver-stained proteins. <i>Electrophoresis</i> , 2003, 24, 237-252.	1.3	26
56	Isolation and characterization of modified species of a mutated (Cys125→Ala) recombinant human interleukin-2. <i>Journal of Chromatography A</i> , 2002, 971, 129-142.	1.8	15
57	Primary structure of two cytolysin isoforms from <i>Stichodactyla helianthus</i> differing in their hemolytic activity.. <i>Toxicon</i> , 2001, 39, 1253-1256.	0.8	57
58	Differentiating β - and γ -aspartic acids by electrospray ionization and low-energy tandem mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2000, 14, 2092-2102.	0.7	67
59	The metastable decomposition of a peptide containing oxidized methionine(s) in matrix-assisted laser desorption/ionization time-of-flight mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 1999, 13, 1075-1076.	0.7	12
60	Automated interpretation of high-energy collision-induced dissociation spectra of singly protonated peptides by 'seqms', a software aid for de novo sequencing by tandem mass spectrometry. , 1998, 12, 1867-1878.		43
61	Effect of the position of a basic amino acid on C-terminal rearrangement of protonated peptides upon collision-induced dissociation. , 1996, 31, 150-158.		55
62	Concanavalin A- and Wheat Germ Agglutinin-Conjugated Lectins as a Tool for the Identification of Multiple N-Glycosylation Sites in Heterologous Protein Expressed in Yeast. <i>Analytical Biochemistry</i> , 1995, 231, 342-348.	1.1	18
63	A method for determination of N-glycosylation sites in glycoproteins by collision-induced dissociation analysis in fast atom bombardment mass spectrometry: Identification of the positions of carbohydrate-linked asparagine in recombinant α -amylase by treatment with peptide-N-glycosidase F in ^{18}O -labeled water. <i>Analytical Biochemistry</i> , 1992, 205, 151-158.	1.1	116