Ibon Iloro

List of Publications by Year in descending order

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304368 377514 1,265 46 22 34 citations h-index g-index papers 47 47 47 1926 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Comprehensive Proteomic Analysis of Human Endometrial Fluid Aspirate. Journal of Proteome Research, 2009, 8, 4622-4632.	1.8	107
2	DNA-induced Secondary Structure of the Carboxyl-terminal Domain of Histone H1. Journal of Biological Chemistry, 2005, 280, 32141-32147.	1.6	86
3	Human tear proteomics and peptidomics in ophthalmology: Toward the translation of proteomic biomarkers into clinical practice. Journal of Proteomics, 2017, 150, 359-367.	1.2	82
4	Structural and Functional Roles of HIV-1 gp41 Pretransmembrane Sequence Segmentation. Biophysical Journal, 2003, 85, 3769-3780.	0.2	79
5	Thermal and pH-Induced Conformational Changes of a \hat{I}^2 -Sheet Protein Monitored by Infrared Spectroscopy. Biochemistry, 1999, 38, 1525-1530.	1.2	70
6	High-throughput proteomic characterization of plasma rich in growth factors (PRGF-Endoret)-derived fibrin clot interactome. Journal of Tissue Engineering and Regenerative Medicine, 2015, 9, E1-E12.	1.3	66
7	Effect of sphingomyelin and cholesterol on the interaction of St II with lipidic interfaces. Toxicon, 2007, 49, 68-81.	0.8	58
8	Human Basal Tear Peptidome Characterization by CID, HCD, and ETD Followed by in Silico and in Vitro Analyses for Antimicrobial Peptide Identification. Journal of Proteome Research, 2015, 14, 2649-2658.	1.8	48
9	Unveiling the olfactory proteostatic disarrangement in Parkinson's disease by proteome-wide profiling. Neurobiology of Aging, 2019, 73, 123-134.	1.5	48
10	Proteome analysis of human serum proteins adsorbed onto different titanium surfaces used in dental implants. Biofouling, 2017, 33, 98-111.	0.8	45
11	Capsid protein identification and analysis of mature Triatoma virus (TrV) virions and naturally occurring empty particles. Virology, 2011, 409, 91-101.	1.1	42
12	MALDI-Imaging Mass Spectrometry: a step forward in the anatomopathological characterization of stenotic aortic valve tissue. Scientific Reports, 2016, 6, 27106.	1.6	39
13	Differential proteomic analysis of endometrial fluid suggests increased inflammation and impaired glucose metabolism in non-implantative IVF cycles and pinpoints PYGB as a putative implantation marker. Human Reproduction, 2018, 33, 1898-1906.	0.4	38
14	Quantitative structure–activity relationship based screening of bioactive peptides identified in ripened cheese. International Dairy Journal, 2013, 33, 184-190.	1.5	37
15	Do Sequence Repeats Play an Equivalent Role in the Choline-binding Module of Pneumococcal LytA Amidase?. Journal of Biological Chemistry, 2000, 275, 26842-26855.	1.6	33
16	Calcium-Dependent Conformational Rearrangements and Protein Stability in Chicken Annexin A5. Biophysical Journal, 2002, 83, 2280-2291.	0.2	28
17	Role of the Transmembrane Domain in the Stability of TrwB, an Integral Protein Involved in Bacterial Conjugation. Journal of Biological Chemistry, 2004, 279, 10955-10961.	1.6	28
18	Evaluation of inter-day and inter-individual variability of tear peptide/protein profiles by MALDI-TOF MS analyses. Molecular Vision, 2012, 18, 1572-82.	1.1	28

#	Article	lF	Citations
19	Methionine Adenosyltransferase \hat{l}_{\pm} -Helix Structure Unfolds at Lower Temperatures than \hat{l}^{2} -Sheet: A 2D-IR Study. Biophysical Journal, 2004, 86, 3951-3958.	0.2	27
20	LUZP1, a novel regulator of primary cilia and the actin cytoskeleton, is a contributing factor in Townes-Brocks Syndrome. ELife, 2020, 9 , .	2.8	27
21	Analysis of Root Plasma Membrane Aquaporins from <i>Brassica oleracea</i> : Post-Translational Modifications, <i>de novo</i> Sequencing and Detection of Isoforms by High Resolution Mass Spectrometry. Journal of Proteome Research, 2010, 9, 3479-3494.	1.8	25
22	Multicentric study of the effect of pre-analytical variables in the quality of plasma samples stored in biobanks using different complementary proteomic methods. Journal of Proteomics, 2017, 150, 109-120.	1.2	25
23	In-depth proteomics and natural peptidomics analyses reveal antibacterial peptides in human endometrial fluid. Journal of Proteomics, 2020, 216, 103652.	1.2	22
24	Proteomic analysis of processing by-products from canned and fresh tuna: Identification of potentially functional food proteins. Food Chemistry, 2012, 134, 1211-1219.	4.2	19
25	Characterization of Ejl, the cell-wall amidase coded by the pneumococcal bacteriophage Ej-1. Protein Science, 2009, 11, 1788-1799.	3.1	18
26	A twoâ€'dimensional IR spectroscopic (2Dâ€'IR) simulation of protein conformational changes. Spectroscopy, 2004, 18, 49-58.	0.8	14
27	Characterization of serum proteins attached to distinct sol–gel hybrid surfaces. Journal of Biomedical Materials Research - Part B Applied Biomaterials, 2018, 106, 1477-1485.	1.6	14
28	A Bacterial TrwC Relaxase Domain Contains a Thermally Stable \hat{l}_{\pm} -Helical Core. Journal of Bacteriology, 2003, 185, 4226-4232.	1.0	13
29	Human tear peptide/protein profiling study of ocular surface diseases by SPE-MALDI-TOF mass spectrometry analyses. EuPA Open Proteomics, 2014, 3, 206-215.	2.5	13
30	Langartech: A Custom-Made MALDI Matrix Sprayer for MALDI Imaging Mass Spectrometry. Journal of the Association for Laboratory Automation, 2016, 21, 260-267.	2.8	13
31	Simulation of FT-IR spectra and 2D-COS analysis for the H/D exchange of two related ligands. Journal of Molecular Structure, 2006, 799, 153-157.	1.8	11
32	Affinity for the Interface Underpins Potency of Antibodies Operating In Membrane Environments. Cell Reports, 2020, 32, 108037.	2.9	10
33	The Kinetics of the Hydrogen/Deuterium Exchange of Epidermal Growth Factor Receptor Ligands. Biophysical Journal, 2008, 94, 4041-4055.	0.2	9
34	Non-invasive detection of drug toxicity in rats by solid-phase extraction and MALDI-TOF analysis of urine samples. Analytical and Bioanalytical Chemistry, 2013, 405, 2311-2320.	1.9	9
35	Smelling the Dark Proteome: Functional Characterization of PITH Domain-Containing Protein 1 (C1orf128) in Olfactory Metabolism. Journal of Proteome Research, 2020, 19, 4826-4843.	1.8	8
36	Borrelia burgdorferi infection induces long-term memory-like responses in macrophages with tissue-wide consequences in the heart. PLoS Biology, 2021, 19, e3001062.	2.6	7

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37	Mass Spectrometric Identification of Endogenous Peptides. Methods in Molecular Biology, 2018, 1719, 59-70.	0.4	5
38	Silica-gelatin hybrid sol-gel coatings: A proteomic study with biocompatibility implications. Journal of Tissue Engineering and Regenerative Medicine, 2018, 12, 1769-1779.	1.3	5
39	Identification of bioactive peptides in raw sheep milk ripened cheese by mass spectrometry combined with quantitative structure activity relationship (QSAR) Analysis Journal of Biotechnology, 2010, 150, 60-60.	1.9	3
40	Two-Dimensional Infrared Correlation Spectroscopy. , 2006, , 73-88.		3
41	Focal accumulation of aromaticity at the CDRH3 loop mitigates 4E10 polyreactivity without altering its HIV neutralization profile. IScience, 2021, 24, 102987.	1.9	1
42	Methods for Human Olfactory Bulb Tissue Studies Using Peptide/Protein MALDI-TOF Imaging Mass Spectrometry (MALDI-IMS). Neuromethods, 2017, , 91-106.	0.2	1
43	Molecular Histology Analysis of Cryopreserved Tissue Using Peptide/Protein MALDI-TOF Imaging Mass Spectrometry (MALDI-IMS). Methods in Molecular Biology, 2022, 2420, 177-190.	0.4	1
44	VP4 Protein Appears as a Product of RNA Encapsidation in Triatoma Virus (TRV). Biophysical Journal, 2011, 100, 404a.	0.2	0
45	Identification and Characterization of Immunodominant Proteins from Tick Tissue Extracts Inducing a Protective Immune Response against Ixodes ricinus in Cattle. Vaccines, 2021, 9, 636.	2.1	0
46	A 2D-Infrared Study of Human Lipoproteins. , 2005, , 135-143.		0