## JesÃ<sup>o</sup>s Mateos MartÃ-n

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3248992/publications.pdf

Version: 2024-02-01

52 papers 1,660 citations

257450 24 h-index 289244 40 g-index

53 all docs 53 docs citations

53 times ranked 2662 citing authors

#	Article	IF	CITATIONS
1	Mitochondrial Dysregulation of Osteoarthritic Human Articular Chondrocytes Analyzed by Proteomics. Molecular and Cellular Proteomics, 2009, 8, 172-189.	3.8	177
2	Differential protein profiling of synovial fluid from rheumatoid arthritis and osteoarthritis patients using LC–MALDI TOF/TOF. Journal of Proteomics, 2012, 75, 2869-2878.	2.4	106
3	Molecular Mechanisms Involved in the Response to Desiccation Stress and Persistence in Acinetobacter baumannii. Journal of Proteome Research, 2014, 13, 460-476.	3.7	90
4	Identification of a Panel of Novel Serum Osteoarthritis Biomarkers. Journal of Proteome Research, 2011, 10, 5095-5101.	3.7	86
5	The FhaB/FhaC two-partner secretion system is involved in adhesion of <i>Acinetobacter baumannii </i> AbH12O-A2 strain. Virulence, 2017, 8, 959-974.	4.4	72
6	Effect of age on pro-inflammatory miRNAs contained in mesenchymal stem cell-derived extracellular vesicles. Scientific Reports, 2017, 7, 43923.	3.3	69
7	Quantitative Proteomic Profiling of Human Articular Cartilage Degradation in Osteoarthritis. Journal of Proteome Research, 2014, 13, 6096-6106.	3.7	66
8	Influence of age on rat bone-marrow mesenchymal stem cells potential. Scientific Reports, 2015, 5, 16765.	3.3	59
9	Pharmacoproteomic study of the effects of chondroitin and glucosamine sulfate on human articular chondrocytes. Arthritis Research and Therapy, 2010, 12, R138.	3.5	52
10	Lamin A deregulation in human mesenchymal stem cells promotes an impairment in their chondrogenic potential and imbalance in their response to oxidative stress. Stem Cell Research, 2013, 11, 1137-1148.	0.7	50
11	Extracellular Proteome of a Highly Invasive Multidrug-resistant Clinical Strain of <i>Acinetobacter baumannii</i> . Journal of Proteome Research, 2012, 11, 5678-5694.	3.7	48
12	Analysis of the Chondrogenic Potential and Secretome of Mesenchymal Stem Cells Derived from Human Umbilical Cord Stroma. Stem Cells and Development, 2011, 20, 1199-1212.	2.1	47
13	Secretome analysis of chondroitin sulfate-treated chondrocytes reveals anti-angiogenic, anti-inflammatory and anti-catabolic properties. Arthritis Research and Therapy, 2012, 14, R202.	3.5	44
14	Quantitative proteomic analysis of hostâ€"pathogen interactions: a study of Acinetobacter baumannii responses to host airways. BMC Genomics, 2015, 16, 422.	2.8	42
15	Metabolic Labeling of Chondrocytes for the Quantitative Analysis of the Interleukin-1-beta-mediated Modulation of Their Intracellular and Extracellular Proteomes. Journal of Proteome Research, 2011, 10, 3701-3711.	3.7	40
16	Proteome profiling of L3 and L4 Anisakis simplex development stages by TMT-based quantitative proteomics. Journal of Proteomics, 2019, 201, 1-11.	2.4	38
17	Proteomic Analysis of Connexin 43 Reveals Novel Interactors Related to Osteoarthritis. Molecular and Cellular Proteomics, 2015, 14, 1831-1845.	3.8	35
18	Pharmacoproteomic Study of Three Different Chondroitin Sulfate Compounds on Intracellular and Extracellular Human Chondrocyte Proteomes. Molecular and Cellular Proteomics, 2012, 11, M111.013417.	3.8	34

#	Article	IF	Citations
19	Metabolic Labeling of Human Bone Marrow Mesenchymal Stem Cells for the Quantitative Analysis of their Chondrogenic Differentiation. Journal of Proteome Research, 2012, 11, 5350-5361.	3.7	30
20	A pharmacoproteomic study confirms the synergistic effect of chondroitin sulfate and glucosamine. Scientific Reports, 2014, 4, 5069.	3.3	30
21	Serum proteomics of active tuberculosis patients and contacts reveals unique processes activated during Mycobacterium tuberculosis infection. Scientific Reports, 2020, 10, 3844.	3.3	29
22	iTRAQ-based analysis of progerin expression reveals mitochondrial dysfunction, reactive oxygen species accumulation and altered proteostasis. Stem Cell Research and Therapy, 2015, 6, 119.	5.5	28
23	CD105+-mesenchymal stem cells migrate into osteoarthritis joint: An animal model. PLoS ONE, 2017, 12, e0188072.	2.5	28
24	Proteome Analysis During Chondrocyte Differentiation in a New Chondrogenesis Model Using Human Umbilical Cord Stroma Mesenchymal Stem Cells. Molecular and Cellular Proteomics, 2012, 11, M111.010496.	3.8	26
25	Secretome analysis of human articular chondrocytes unravels catabolic effects of nicotine on the joint. Proteomics - Clinical Applications, 2016, 10, 671-680.	1.6	26
26	Molecular characterization of B-cell epitopes for the major fish allergen, parvalbumin, by shotgun proteomics, protein-based bioinformatics and IgE-reactive approaches. Journal of Proteomics, 2019, 200, 123-133.	2.4	26
27	The Coevolution of Insect Muscle TpnT and TpnI Gene Isoforms. Molecular Biology and Evolution, 2005, 22, 2231-2242.	8.9	25
28	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.	2.4	25
29	Diversification and Independent Evolution of Troponin C Genes in Insects. Journal of Molecular Evolution, 2005, 60, 31-44.	1.8	23
30	Hypoxia Conditions Differentially Modulate Human Normal and Osteoarthritic Chondrocyte Proteomes. Journal of Proteome Research, 2010, 9, 3035-3045.	3.7	22
31	The structural role of high molecular weight tropomyosins in dipteran indirect flight muscle and the effect of phosphorylation. Journal of Muscle Research and Cell Motility, 2006, 27, 189-201.	2.0	21
32	High-resolution quantitative proteomics applied to the study of the specific protein signature in the sputum and saliva of active tuberculosis patients and their infected and uninfected contacts. Journal of Proteomics, 2019, 195, 41-52.	2.4	20
33	Strategies to optimize two-dimensional gel electrophoresis analysis of the human joint proteome. Talanta, 2010, 80, 1552-1560.	5.5	18
34	Next-Generation Sequencing and Quantitative Proteomics of Hutchinson-Gilford progeria syndrome-derived cells point to a role of nucleotide metabolism in premature aging. PLoS ONE, 2018, 13, e0205878.	2.5	16
35	Comparative Proteomics Analysis of Anisakis simplex s.s.—Evaluation of the Response of Invasive Larvae to Ivermectin. Genes, 2020, 11, 710.	2.4	15
36	Proteomic analysis and biochemical alterations in marine mussel gills after exposure to the organophosphate flame retardant TDCPP. Aquatic Toxicology, 2021, 230, 105688.	4.0	15

#	Article	IF	CITATIONS
37	Dimethylarginine dimethylaminohydrolase 2, a newly identified mitochondrial protein modulating nitric oxide synthesis in normal human chondrocytes. Arthritis and Rheumatism, 2012, 64, 204-212.	6.7	12
38	Predictive modeling of therapeutic response to chondroitin sulfate/glucosamine hydrochloride in knee osteoarthritis. Therapeutic Advances in Chronic Disease, 2019, 10, 204062231987001.	2.5	11
39	3, 3′, 5â€triiodoâ€Lâ€thyronine Increases In Vitro Chondrogenesis of Mesenchymal Stem Cells From Human Umbilical Cord Stroma Through SRC2. Journal of Cellular Biochemistry, 2016, 117, 2097-2108.	2.6	9
40	Proteomic Applications in the Study of Human Mesenchymal Stem Cells. Proteomes, 2014, 2, 53-71.	3.5	7
41	LC-MALDI-TOF/TOF for Shotgun Proteomics. Methods in Molecular Biology, 2014, 1156, 27-38.	0.9	7
42	The role of gravity in the evolutionary emergence of multicellular complexity: Microgravity effects on arthropod development and aging. Advances in Space Research, 1999, 23, 2075-2082.	2.6	5
43	Cryoconservation of Peptide Extracts from Trypsin Digestion of Proteins for Proteomic Analysis in a Hospital Biobank Facility. Journal of Proteome Research, 2014, 13, 1930-1937.	3.7	5
44	Mesenchymal Stem Cell-Derived Extracellular Isolation and Their Protein Cargo Characterization. Methods in Molecular Biology, 2021, 2259, 3-12.	0.9	5
45	Drosophila melanogaster and the Future of â€~Evo-Devo' Biology in Space. Challenges and Problems in the Path of an Eventual Colonization Project Outside the Earth. Advances in Space Biology and Medicine, 2003, 9, 41-81.	0.5	4
46	Shotgun for L3 and L4 Development Stages. Methods in Molecular Biology, 2021, 2259, 59-75.	0.9	3
47	A Complex Proteomic Response of the Parasitic Nematode Anisakis simplex s.s. to Escherichia coli Lipopolysaccharide. Molecular and Cellular Proteomics, 2021, 20, 100166.	3.8	3
48	Identification of autoantibodies in serum from osteoarthritis patients using microarrays. Osteoarthritis and Cartilage, 2014, 22, S425.	1.3	2
49	The Spanish biology/disease initiative within the human proteome project: Application to rheumatic diseases. Journal of Proteomics, 2015, 127, 406-413.	2.4	2
50	Tandem Mass Tagging (TMT) Reveals Tissue-Specific Proteome of L4 Larvae of Anisakis simplex s. s.: Enzymes of Energy and/or Carbohydrate Metabolism as Potential Drug Targets in Anisakiasis. International Journal of Molecular Sciences, 2022, 23, 4336.	4.1	2
51	Data Treatment in Food Proteomics. , 2021, , 324-338.		1
52	Identification of an acetyl esterase in the supernatant of the environmental strain Bacillus sp. HR21-6. Biochimie, 2022, 198, 48-59.	2.6	0