Jesus Vazquez

List of Publications by Year in descending order

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255 papers 17,629 citations

63 h-index 19190 118 g-index

266 all docs 266
docs citations

times ranked

266

27255 citing authors

#	Article	IF	CITATIONS
1	Sumoylated hnRNPA2B1 controls the sorting of miRNAs into exosomes through binding to specific motifs. Nature Communications, 2013, 4, 2980.	12.8	1,522
2	Vesiclepedia: A Compendium for Extracellular Vesicles with Continuous Community Annotation. PLoS Biology, 2012, 10, e1001450.	5.6	1,064
3	Multiple evidence strands suggest that there may be as few as 19 000 human protein-coding genes. Human Molecular Genetics, 2014, 23, 5866-5878.	2.9	463
4	Genomic insights into the Ixodes scapularis tick vector of Lyme disease. Nature Communications, 2016, 7, 10507.	12.8	450
5	The Intracellular Interactome of Tetraspanin-enriched Microdomains Reveals Their Function as Sorting Machineries toward Exosomes. Journal of Biological Chemistry, 2013, 288, 11649-11661.	3.4	377
6	Glutathionylation of the p50 Subunit of NF-κB:  a Mechanism for Redox-Induced Inhibition of DNA Binding. Biochemistry, 2001, 40, 14134-14142.	2.5	366
7	A Network of Macrophages Supports Mitochondrial Homeostasis in the Heart. Cell, 2020, 183, 94-109.e23.	28.9	360
8	Mitochondrial and nuclear DNA matching shapes metabolism and healthy ageing. Nature, 2016, 535, 561-565.	27.8	333
9	SQANTI: extensive characterization of long-read transcript sequences for quality control in full-length transcriptome identification and quantification. Genome Research, 2018, 28, 396-411.	5.5	299
10	S-nitrosylation of Hsp90 promotes the inhibition of its ATPase and endothelial nitric oxide synthase regulatory activities. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 8525-8530.	7.1	294
11	A polymorphism in the regulatory region of APOE associated with risk for Alzheimer's dementia. Nature Genetics, 1998, 18, 69-71.	21.4	291
12	Mammalian lipid droplets are innate immune hubs integrating cell metabolism and host defense. Science, 2020, 370, .	12.6	245
13	The CoQH2/CoQ Ratio Serves as a Sensor of Respiratory Chain Efficiency. Cell Reports, 2016, 15, 197-209.	6.4	215
14	Allelic polymorphisms in the transcriptional regulatory region of apolipoprotein E gene. FEBS Letters, 1998, 421, 105-108.	2.8	213
15	ITAM-Based Interaction of ERM Proteins with Syk Mediates Signaling by the Leukocyte Adhesion Receptor PSGL-1. Immunity, 2002, 17, 401-412.	14.3	200
16	ATP-Dependent Lon Protease Controls Tumor Bioenergetics by Reprogramming Mitochondrial Activity. Cell Reports, 2014, 8, 542-556.	6.4	186
17	Use of toxins to study potassium channels. Journal of Bioenergetics and Biomembranes, 1991, 23, 615-646.	2.3	184
18	Programmed \hat{a} e disarming \hat{a} e of the neutrophil proteome reduces the magnitude of inflammation. Nature Immunology, 2020, 21, 135-144.	14.5	180

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19	MTOC translocation modulates IS formation and controls sustained T cell signaling. Journal of Cell Biology, 2008, 182, 951-962.	5.2	165
20	General Statistical Framework for Quantitative Proteomics by Stable Isotope Labeling. Journal of Proteome Research, 2014, 13, 1234-1247.	3.7	165
21	A Refined Method To Calculate False Discovery Rates for Peptide Identification Using Decoy Databases. Journal of Proteome Research, 2009, 8, 1792-1796.	3.7	161
22	Mechanism of super-assembly of respiratory complexes III and IV. Nature, 2016, 539, 579-582.	27.8	157
23	Ultra Fast Trypsin Digestion of Proteins by High Intensity Focused Ultrasound. Journal of Proteome Research, 2005, 4, 1569-1574.	3.7	155
24	Properties of Average Score Distributions of SEQUEST. Molecular and Cellular Proteomics, 2008, 7, 1135-1145.	3.8	142
25	Systems Biology of Tissue-Specific Response to Anaplasma phagocytophilum Reveals Differentiated Apoptosis in the Tick Vector Ixodes scapularis. PLoS Genetics, 2015, 11, e1005120.	3 . 5	139
26	Risk for Alzheimer's disease correlates with transcriptional activity of the APOE gene. Human Molecular Genetics, 1998, 7, 1887-1892.	2.9	135
27	APPRIS 2017: principal isoforms for multiple gene sets. Nucleic Acids Research, 2018, 46, D213-D217.	14.5	134
28	Interplay between hepatic mitochondria-associated membranes, lipid metabolism and caveolin-1 in mice. Scientific Reports, 2016, 6, 27351.	3.3	131
29	Self-Renewing Human Bone Marrow Mesenspheres Promote Hematopoietic Stem Cell Expansion. Cell Reports, 2013, 3, 1714-1724.	6.4	128
30	Optic Atrophy 1 Is Epistatic to the Core MICOS Component MIC60 in Mitochondrial Cristae Shape Control. Cell Reports, 2016, 17, 3024-3034.	6.4	127
31	Connexin43 in cardiomyocyte mitochondria contributes to mitochondrial potassium uptake. Cardiovascular Research, 2009, 83, 747-756.	3.8	124
32	Bone Marrow Mesenchymal Stem Cells Support Acute Myeloid Leukemia Bioenergetics and Enhance Antioxidant Defense and Escape from Chemotherapy. Cell Metabolism, 2020, 32, 829-843.e9.	16.2	122
33	The cristae modulator Optic atrophy 1 requires mitochondrial ATP synthase oligomers to safeguard mitochondrial function. Nature Communications, 2018, 9, 3399.	12.8	111
34	Most Highly Expressed Protein-Coding Genes Have a Single Dominant Isoform. Journal of Proteome Research, 2015, 14, 1880-1887.	3.7	106
35	Peptide Rearrangement during Quadrupole Ion Trap Fragmentation:Â Added Complexity to MS/MS Spectra. Analytical Chemistry, 2003, 75, 1524-1535.	6.5	101
36	Statistical Model for Large-Scale Peptide Identification in Databases from Tandem Mass Spectra Using SEQUEST. Analytical Chemistry, 2004, 76, 6853-6860.	6.5	101

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37	Analyzing the First Drafts of the Human Proteome. Journal of Proteome Research, 2014, 13, 3854-3855.	3.7	101
38	Anaplasma phagocytophilum Inhibits Apoptosis and Promotes Cytoskeleton Rearrangement for Infection of Tick Cells. Infection and Immunity, 2013, 81, 2415-2425.	2.2	99
39	CD69 controls the uptake of L-tryptophan through LAT1-CD98 and AhR-dependent secretion of IL-22 in psoriasis. Nature Immunology, 2016, 17, 985-996.	14.5	98
40	Characterization and partial sequencing of species-specific sarcoplasmic polypeptides from commercial hake species by mass spectrometry following two-dimensional electrophoresis. Electrophoresis, 2001, 22, 1545-1552.	2.4	96
41	A Robust Method for Quantitative High-throughput Analysis of Proteomes by 18O Labeling. Molecular and Cellular Proteomics, 2011, 10, M110.003335.	3.8	95
42	Dissecting the proteome dynamics of the early heat stress response leading to plant survival or death in Arabidopsis. Plant, Cell and Environment, 2016, 39, 1264-1278.	5.7	94
43	Proteomics as a Tool for the Investigation of Seafood and Other Marine Products. Journal of Proteome Research, 2003, 2, 127-135.	3.7	92
44	Caveolin-1 Modulates Mechanotransduction Responses to Substrate Stiffness through Actin-Dependent Control of YAP. Cell Reports, 2018, 25, 1622-1635.e6.	6.4	91
45	Identification of commercial hake and grenadier species by proteomic analysis of the parvalbumin fraction. Proteomics, 2006, 6, 5278-5287.	2.2	90
46	A Novel Systems-Biology Algorithm for the Analysis of Coordinated Protein Responses Using Quantitative Proteomics. Molecular and Cellular Proteomics, 2016, 15, 1740-1760.	3.8	86
47	Defective sarcoplasmic reticulum–mitochondria calcium exchange in aged mouse myocardium. Cell Death and Disease, 2014, 5, e1573-e1573.	6.3	85
48	Fast Monitoring of Species-Specific Peptide Biomarkers Using High-Intensity-Focused-Ultrasound-Assisted Tryptic Digestion and Selected MS/MS Ion Monitoring. Analytical Chemistry, 2011, 83, 5688-5695.	6.5	81
49	Alternatively Spliced Homologous Exons Have Ancient Origins and Are Highly Expressed at the Protein Level. PLoS Computational Biology, 2015, 11, e1004325.	3.2	80
50	The Leukocyte Activation Receptor CD69 Controls T Cell Differentiation through Its Interaction with Galectin-1. Molecular and Cellular Biology, 2014, 34, 2479-2487.	2.3	79
51	Caveolinâ€1 deficiency induces a <scp>MEK</scp> â€ <scp>ERK</scp> 1/2â€Snailâ€1â€dependent epithelial–mesenchymal transition and fibrosis during peritoneal dialysis. EMBO Molecular Medicine, 2015, 7, 102-123.	6.9	79
52	Improved Method for Differential Expression Proteomics Using Trypsin-catalyzed 18O Labeling with a Correction for Labeling Efficiency. Molecular and Cellular Proteomics, 2007, 6, 1274-1286.	3.8	78
53	Differential Association of HLA-B*2705 and B*2709 to Ankylosing Spondylitis Correlates with Limited Peptide Subsets but Not with Altered Cell Surface Stability. Journal of Biological Chemistry, 2002, 277, 28749-28756.	3.4	77
54	Statistical Model to Analyze Quantitative Proteomics Data Obtained by 180/160 Labeling and Linear Ion Trap Mass Spectrometry. Molecular and Cellular Proteomics, 2009, 8, 1130-1149.	3.8	76

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55	ISC15 governs mitochondrial function in macrophages following vaccinia virus infection. PLoS Pathogens, 2017, 13, e1006651.	4.7	75
56	The glycine receptor: pharmacological studies and mathematical modeling of the allosteric interaction between the glycine- and strychnine-binding sites. Molecular Pharmacology, 1986, 30, 590-7.	2.3	75
57	<i>De </i> Novo Mass Spectrometry Sequencing and Characterization of Species-Specific Peptides from Nucleoside Diphosphate Kinase B for the Classification of Commercial Fish Species Belonging to the Family Merlucciidae. Journal of Proteome Research, 2007, 6, 3070-3080.	3.7	74
58	$p38\hat{l}^3$ is essential for cell cycle progression and liver tumorigenesis. Nature, 2019, 568, 557-560.	27.8	72
59	ALDH4A1 is an atherosclerosis auto-antigen targeted by protective antibodies. Nature, 2021, 589, 287-292.	27.8	72
60	Application of proteomics for fast identification of species-specific peptides from marine species. Proteomics, 2002, 2, 1658-1665.	2.2	70
61	Transcriptional activation by AP-2alpha is modulated by the oncogene DEK. Nucleic Acids Research, 2003, 31, 1571-1575.	14.5	70
62	Highâ€sensitivity analysis of specific peptides in complex samples by selected MS/MS ion monitoring and linear ion trap mass spectrometry: Application to biological studies. Journal of Mass Spectrometry, 2007, 42, 1391-1403.	1.6	68
63	p38 \hat{l}^3 and \hat{l}' promote heart hypertrophy by targeting the mTOR-inhibitory protein DEPTOR for degradation. Nature Communications, 2016, 7, 10477.	12.8	68
64	Functional role of respiratory supercomplexes in mice: SCAF1 relevance and segmentation of the Q _{pool} . Science Advances, 2020, 6, eaba7509.	10.3	68
65	The Peptide Repertoires of HLA-B27 Subtypes Differentially Associated to Spondyloarthropathy (B*2704) Tj ETQq1 2002, 277, 16744-16749.		14 rgBT /0 66
66	A proteomic approach to the study of the marine mussels Mytilus edulis and M . galloprovincialis. Marine Biology, 2002, 141, 217-223.	1.5	66
67	Flow Cytometry Has a Significant Impact on the Cellular Metabolome. Journal of Proteome Research, 2019, 18, 169-181.	3.7	66
68	A Novel Strategy for Global Analysis of the Dynamic Thiol Redox Proteome. Molecular and Cellular Proteomics, 2012, 11, 800-813.	3.8	65
69	Revisiting Peptide Identification by High-Accuracy Mass Spectrometry: Problems Associated with the Use of Narrow Mass Precursor Windows. Journal of Proteome Research, 2015, 14, 700-710.	3.7	65
70	CD81 regulates cell migration through its association with Rac GTPase. Molecular Biology of the Cell, 2013, 24, 261-273.	2.1	64
71	Transcription factor AP-2 activity is modulated by protein kinase A-mediated phosphorylation. FEBS Letters, 1999, 444, 27-31.	2.8	62
72	Blockade of NFAT Activation by the Second Calcineurin Binding Site. Journal of Biological Chemistry, 2006, 281, 6227-6235.	3.4	62

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73	Loose ends: almost one in five human genes still have unresolved coding status. Nucleic Acids Research, 2018, 46, 7070-7084.	14.5	62
74	Severe Cardiac Dysfunction and Death Caused by Arrhythmogenic Right Ventricular Cardiomyopathy Type 5 Are Improved by Inhibition of Glycogen Synthase Kinase-3Î ² . Circulation, 2019, 140, 1188-1204.	1.6	62
75	Ryanodine Receptor Glycation Favors Mitochondrial Damage in the Senescent Heart. Circulation, 2019, 139, 949-964.	1.6	62
76	Cyclosporine A-induced nitration of tyrosine 34 MnSOD in endothelial cells: role of mitochondrial superoxide. Cardiovascular Research, 2010, 87, 356-365.	3.8	61
77	Proteome-wide alterations on adipose tissue from obese patients as age-, diabetes- and gender-specific hallmarks. Scientific Reports, 2016, 6, 25756.	3.3	61
78	Transcription Factor AP-2 Regulates Human Apolipoprotein E Gene Expression in Astrocytoma Cells. Journal of Neuroscience, 1996, 16, 7550-7556.	3.6	60
79	Extensive <i>De Novo</i> Sequencing of New Parvalbumin Isoforms Using a Novel Combination of Bottom-Up Proteomics, Accurate Molecular Mass Measurement by FTICRâ^'MS, and Selected MS/MS Ion Monitoring. Journal of Proteome Research, 2010, 9, 4393-4406.	3.7	60
80	Arabidopsis SWC4 Binds DNA and Recruits the SWR1 Complex to Modulate Histone H2A.Z Deposition at Key Regulatory Genes. Molecular Plant, 2018, 11, 815-832.	8.3	60
81	Characterization of high affinity binding sites for charybdotoxin in synaptic plasma membranes from rat brain. Evidence for a direct association with an inactivating, voltage-dependent, potassium channel. Journal of Biological Chemistry, 1990, 265, 15564-71.	3.4	60
82	c-Jun N-terminal Kinase (JNK) Positively Regulates NFATc2 Transactivation through Phosphorylation within the N-terminal Regulatory Domain. Journal of Biological Chemistry, 2005, 280, 20867-20878.	3.4	59
83	SanXoT: a modular and versatile package for the quantitative analysis of high-throughput proteomics experiments. Bioinformatics, 2019, 35, 1594-1596.	4.1	59
84	High-sensitivity analysis and sequencing of peptides and proteins by quadrupole ion trap mass spectrometry. Journal of Mass Spectrometry, 1999, 34, 17-27.	1.6	58
85	Ischemic preconditioning protects cardiomyocyte mitochondria through mechanisms independent of cytosol. Journal of Molecular and Cellular Cardiology, 2014, 68, 79-88.	1.9	58
86	The immunomodulatory activity of extracellular vesicles derived from endometrial mesenchymal stem cells on CD4+ T cells is partially mediated by TGFbeta. Journal of Tissue Engineering and Regenerative Medicine, 2018, 12, 2088-2098.	2.7	58
87	ECM deposition is driven by caveolin-1–dependent regulation of exosomal biogenesis and cargo sorting. Journal of Cell Biology, 2020, 219, .	5.2	58
88	Interaction of bilirubin with the synaptosomal plasma membrane. Journal of Biological Chemistry, 1988, 263, 1255-65.	3.4	58
89	Quantitative proteomics using $160/180$ labeling and linear ion trap mass spectrometry. Proteomics, 2006, 6, S4-S11.	2.2	57
90	Sample treatment for protein identification by mass spectrometry-based techniques. TrAC - Trends in Analytical Chemistry, 2006, 25, 996-1005.	11.4	57

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91	Expression of Heat Shock and Other Stress Response Proteins in Ticks and Cultured Tick Cells in Response to <i>Anaplasma</i> Spp. Infection and Heat Shock. International Journal of Proteomics, 2010, 2010, 1-11.	2.0	55
92	An analysis of tissue-specific alternative splicing at the protein level. PLoS Computational Biology, 2020, 16, e1008287.	3.2	55
93	Purification and characterization of paraoxon hydrolase from rat liver. Biochemical Journal, 1997, 321, 595-601.	3.7	54
94	F-actin-binding protein drebrin regulates CXCR4 recruitment to the immune synapse. Journal of Cell Science, 2010, 123, 1160-1170.	2.0	54
95	Cutting Edge: Association of the Motor Protein Nonmuscle Myosin Heavy Chain-IIA with the C Terminus of the Chemokine Receptor CXCR4 in T Lymphocytes. Journal of Immunology, 2002, 169, 5410-5414.	0.8	53
96	Deficiency of MMP17/MT4-MMP Proteolytic Activity Predisposes to Aortic Aneurysm in Mice. Circulation Research, 2015, 117, e13-26.	4.5	53
97	Aurora A drives early signalling and vesicle dynamics during T-cell activation. Nature Communications, 2016, 7, 11389.	12.8	53
98	Characterization of high affinity binding sites for charybdotoxin in sarcolemmal membranes from bovine aortic smooth muscle. Evidence for a direct association with the high conductance calcium-activated potassium channel. Journal of Biological Chemistry, 1989, 264, 20902-9.	3.4	53
99	The same natural ligand is involved in allorecognition of multiple HLA-B27 subtypes by a single T cell clone: role of peptide and the MHC molecule in alloreactivity. Journal of Immunology, 1998, 161, 5481-90.	0.8	53
100	Differential effects of the tricyclic antidepressant amoxapine on glycine uptake mediated by the recombinant GLYT1 and GLYT2 glycine transporters. British Journal of Pharmacology, 2000, 129, 200-206.	5 . 4	52
101	Growth, mortality, pathological conditions and protein expression of Mytilus edulis and M. galloprovincialis crosses cultured in the Rıla de Arousa (NW of Spain). Aquaculture, 2002, 213, 233-251.	3.5	52
102	White matter injury restoration after stem cell administration in subcortical ischemic stroke. Stem Cell Research and Therapy, 2015, 6, 121.	5 . 5	52
103	miR-28 regulates the germinal center reaction and blocks tumor growth in preclinical models of non-Hodgkin lymphoma. Blood, 2017, 129, 2408-2419.	1.4	52
104	Two-dimensional gel electrophoresis of Mytilus galloprovincialis differences in protein expression between intertidal and cultured mussels. Marine Ecology - Progress Series, 2001, 224, 149-156.	1.9	51
105	New protein–protein interactions of mitochondrial connexin 43 in mouse heart. Journal of Cellular and Molecular Medicine, 2016, 20, 794-803.	3.6	49
106	Proteomic and transcriptomic analyses of differential stress/inflammatory responses in mandibular lymph nodes and oropharyngeal tonsils of European wild boars naturally infected withMycobacterium bovis. Proteomics, 2007, 7, 220-231.	2.2	48
107	Targeting L-type amino acid transporter 1 in innate and adaptive T cells efficiently controls skin inflammation. Journal of Allergy and Clinical Immunology, 2020, 145, 199-214.e11.	2.9	47
108	Altered FoF1 ATP synthase and susceptibility to mitochondrial permeability transition pore during ischaemia and reperfusion in aging cardiomyocytes. Thrombosis and Haemostasis, 2015, 113, 441-451.	3.4	46

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109	Activation of Serine One-Carbon Metabolism by Calcineurin A \hat{l}^21 Reduces Myocardial Hypertrophy and Improves Ventricular Function. Journal of the American College of Cardiology, 2018, 71, 654-667.	2.8	45
110	Thermodynamics of Agonist and Antagonist Interaction with the Strychnine-Sensitive Glycine Receptor. Journal of Neurochemistry, 1989, 52, 1775-1780.	3.9	44
111	EWI-2 Association with α-Actinin Regulates T Cell Immune Synapses and HIV Viral Infection. Journal of Immunology, 2012, 189, 689-700.	0.8	44
112	Extracellular vesicles derived from endometrial human mesenchymal stem cells enhance embryo yield and quality in an aged murine modelâ€. Biology of Reproduction, 2019, 100, 1180-1192.	2.7	44
113	Scaf1 promotes respiratory supercomplexes and metabolic efficiency in zebrafish. EMBO Reports, 2020, 21, e50287.	4.5	42
114	Purification of the sodium- and chloride-coupled glycine transporter from central nervous system. Journal of Biological Chemistry, 1991, 266, 24809-14.	3.4	42
115	Beneficial effects of omega-3 fatty acids in the proteome of high-density lipoprotein proteome. Lipids in Health and Disease, 2012, 11, 116.	3.0	41
116	ApoA-I/HDL-C levels are inversely associated with abdominal aortic aneurysm progression. Thrombosis and Haemostasis, 2015, 113, 1335-1346.	3.4	41
117	Loss of SRSF3 in Cardiomyocytes Leads to Decapping of Contraction-Related mRNAs and Severe Systolic Dysfunction. Circulation Research, 2019, 125, 170-183.	4.5	41
118	Inhibition of insulin release by synthetic peptides shows that the H3 region at the C-terminal domain of syntaxin-1 is crucial for Ca2+- but not for guanosine 5′-[γ-thio]triphosphate-induced secretion. Biochemical Journal, 1996, 320, 201-205.	3.7	40
119	Limited Diversity of Peptides Related to an Alloreactive T Cell Epitope in the HLA-B27-Bound Peptide Repertoire Results from Restrictions at Multiple Steps Along the Processing-Loading Pathway. Journal of Immunology, 2000, 164, 329-337.	0.8	40
120	Differential proteomic and oxidative profiles unveil dysfunctional protein import to adipocyte mitochondria in obesity-associated aging and diabetes. Redox Biology, 2017, 11, 415-428.	9.0	40
121	APOA1 oxidation is associated to dysfunctional high-density lipoproteins in human abdominal aortic aneurysm. EBioMedicine, 2019, 43, 43-53.	6.1	40
122	Evidence of saxitoxin derivatives as causative agents in the 1997 mass mortality of monk seals in the Cape Blanc Peninsula. Natural Toxins, 1999, 7, 311-315.	1.0	39
123	CXCL6 is an important paracrine factor in the pro-angiogenic human cardiac progenitor-like cell secretome. Scientific Reports, 2017, 7, 12490.	3.3	39
124	Comprehensive Quantification of the Modified Proteome Reveals Oxidative Heart Damage in Mitochondrial Heteroplasmy. Cell Reports, 2018, 23, 3685-3697.e4.	6.4	39
125	Arabidopsis YAF9 histone readers modulate flowering time through NuA4â€complexâ€dependent H4 and H2A.Z histone acetylation at <i>FLC</i> chromatin. New Phytologist, 2019, 222, 1893-1908.	7.3	39
126	Caveolin1 and YAP drive mechanically induced mesothelial to mesenchymal transition and fibrosis. Cell Death and Disease, 2020, 11, 647.	6.3	39

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127	The Cys-67 Residue of HLA-B27 Influences Cell Surface Stability, Peptide Specificity, and T-cell Antigen Presentation. Journal of Biological Chemistry, 2001, 276, 48740-48747.	3.4	38
128	HDAC6 controls innate immune and autophagy responses to TLR-mediated signalling by the intracellular bacteria Listeria monocytogenes. PLoS Pathogens, 2017, 13, e1006799.	4.7	38
129	Unraveling the Molecular Signature of Extracellular Vesicles From Endometrial-Derived Mesenchymal Stem Cells: Potential Modulatory Effects and Therapeutic Applications. Frontiers in Bioengineering and Biotechnology, 2019, 7, 431.	4.1	38
130	Executioner Caspase-3 and 7 Deficiency Reduces Myocyte Number in the Developing Mouse Heart. PLoS ONE, 2015, 10, e0131411.	2.5	38
131	Specific interaction of heterogeneous nuclear ribonucleoprotein A1 with the -219T allelic form modulates APOE promoter activity. Nucleic Acids Research, 2003, 31, 3063-3070.	14.5	37
132	Loss of the proteostasis factor AIRAPL causes myeloid transformation by deregulating IGF-1 signaling. Nature Medicine, 2016, 22, 91-96.	30.7	37
133	Proteomic footprint of myocardial ischemia/reperfusion injury: Longitudinal study of the at-risk and remote regions in the pig model. Scientific Reports, 2017, 7, 12343.	3.3	37
134	The metalloprotease ADAM8 is associated with and regulates the function of the adhesion receptor PSGL†through ERM proteins. European Journal of Immunology, 2011, 41, 3436-3442.	2.9	36
135	Oxidized Low-Density Lipoprotein Receptor in Lymphocytes Prevents Atherosclerosis and Predicts Subclinical Disease. Circulation, 2019, 139, 243-255.	1.6	36
136	Na+/K+-ATPase Is a New Interacting Partner for the Neuronal Glycine Transporter GlyT2 That Downregulates Its Expression In Vitro and In Vivo. Journal of Neuroscience, 2013, 33, 14269-14281.	3.6	35
137	NOX4-dependent Hydrogen peroxide promotes shear stress-induced SHP2 sulfenylation and eNOS activation. Free Radical Biology and Medicine, 2015, 89, 419-430.	2.9	35
138	A Single In-Vial Dual Extraction Strategy for the Simultaneous Lipidomics and Proteomics Analysis of HDL and LDL Fractions. Journal of Proteome Research, 2016, 15, 1762-1775.	3.7	35
139	Arabidopsis DNA polymerase ϵ recruits components of Polycomb repressor complex to mediate epigenetic gene silencing. Nucleic Acids Research, 2016, 44, 5597-5614.	14.5	34
140	MMP-25 Metalloprotease Regulates Innate Immune Response through NF-κB Signaling. Journal of Immunology, 2016, 197, 296-302.	0.8	34
141	The intracellular bacterium Anaplasma phagocytophilum selectively manipulates the levels of vertebrate host proteins in the tick vector lxodes scapularis. Parasites and Vectors, 2016, 9, 467.	2.5	33
142	The apparent variability of silkworm (Bombyx mori) silk and its relationship with degumming. European Polymer Journal, 2016, 78, 129-140.	5.4	33
143	Urinary exosomes reveal protein signatures in hypertensive patients with albuminuria. Oncotarget, 2017, 8, 44217-44231.	1.8	33
144	Cardiomyocyte hypertrophy induced by Endonuclease G deficiency requires reactive oxygen radicals accumulation and is inhibitable by the micropeptide humanin. Redox Biology, 2018, 16, 146-156.	9.0	32

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145	Complement C5 Protein as a Marker of Subclinical Atherosclerosis. Journal of the American College of Cardiology, 2020, 75, 1926-1941.	2.8	32
146	Mechanical control of nuclear import by Importin-7 is regulated by its dominant cargo YAP. Nature Communications, 2022, 13, 1174.	12.8	32
147	Identification and Characterization of Anaplasma phagocytophilum Proteins Involved in Infection of the Tick Vector, Ixodes scapularis. PLoS ONE, 2015, 10, e0137237.	2.5	31
148	Successful aging: insights from proteome analyses of healthy centenarians. Aging, 2020, 12, 3502-3515.	3.1	31
149	Identification of phosphorylation sites in proteins by nanospray quadrupole ion trap mass spectrometry., 2000, 35, 556-565.		30
150	The human HDL proteome displays high inter-individual variability and is altered dynamically in response to angioplasty-induced atheroma plaque rupture. Journal of Proteomics, 2014, 106, 61-73.	2.4	30
151	Quantitative HDL Proteomics Identifies Peroxiredoxin-6 as a Biomarker of Human Abdominal Aortic Aneurysm. Scientific Reports, 2016, 6, 38477.	3.3	29
152	HDAC6 regulates the dynamics of lytic granules in cytotoxic T lymphocytes. Journal of Cell Science, 2016, 129, 1305-1311.	2.0	29
153	APPRIS: selecting functionally important isoforms. Nucleic Acids Research, 2022, 50, D54-D59.	14.5	29
154	Aortic disease in Marfan syndrome is caused by overactivation of sGC-PRKG signaling by NO. Nature Communications, 2021, 12, 2628.	12.8	28
155	Functional reconstitution of the glycine receptor. Biochemistry, 1989, 28, 6405-6409.	2.5	27
156	Apolipoprotein E forms stable complexes with recombinant Alzheimer's disease β-amyloid precursor protein. Biochemical Journal, 1997, 325, 169-175.	3.7	27
157	P24, a glycogen synthase kinase 3 (GSK 3) inhibitor. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2002, 1586, 113-122.	3.8	27
158	Trypanosoma cruzi mitochondrial malate dehydrogenase triggers polyclonal B-cell activation. Clinical and Experimental Immunology, 2002, 127, 27-36.	2.6	27
159	Characterization of natural peptide ligands from HLA-DP2: new insights into HLA-DP peptide-binding motifs. Immunogenetics, 2005, 56, 754-759.	2.4	27
160	Production, partial characterization and mass spectrometric studies of the extracellular laccase activity from Fusarium proliferatum. Applied Microbiology and Biotechnology, 2006, 70, 212-221.	3.6	27
161	Application of highly sensitive saturation labeling to the analysis of differential protein expression in infected ticks from limited samples. Proteome Science, 2010, 8, 43.	1.7	27
162	Characterization of the solubilized charybdotoxin receptor from bovine aortic smooth muscle. Biochemistry, 1991, 30, 11157-11164.	2.5	26

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163	Phosphatidylcholine oated Iron Oxide Nanomicelles for In Vivo Prolonged Circulation Time with an Antibiofouling Protein Corona. Chemistry - A European Journal, 2014, 20, 16662-16671.	3.3	26
164	The potential clinical impact of the release of two drafts of the human proteome. Expert Review of Proteomics, 2015, 12, 579-593.	3.0	26
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