Shankar Subramaniam

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6252194/publications.pdf

Version: 2024-02-01

87 5,786 papers citations

33 72
h-index g-index

93 93 docs citations

93 times ranked 11088 citing authors

#	Article	IF	Citations
1	Metabolomics Workbench: An international repository for metabolomics data and metadata, metabolite standards, protocols, tutorials and training, and analysis tools. Nucleic Acids Research, 2016, 44, D463-D470.	14.5	568
2	Update on LIPID MAPS classification, nomenclature, and shorthand notation for MS-derived lipid structures. Journal of Lipid Research, 2020, 61, 1539-1555.	4.2	372
3	p62, Upregulated during Preneoplasia, Induces Hepatocellular Carcinogenesis by Maintaining Survival of Stressed HCC-Initiating Cells. Cancer Cell, 2016, 29, 935-948.	16.8	353
4	Systems Analysis of Immunity to Influenza Vaccination across Multiple Years and in Diverse Populations Reveals Shared Molecular Signatures. Immunity, 2015, 43, 1186-1198.	14.3	286
5	Biomarkers of NAFLD progression: a lipidomics approach to an epidemic. Journal of Lipid Research, 2015, 56, 722-736.	4.2	264
6	Skeletal muscle: A review of molecular structure and function, in health and disease. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2020, 12, e1462.	6.6	262
7	A Mouse Macrophage Lipidome. Journal of Biological Chemistry, 2010, 285, 39976-39985.	3.4	260
8	Adjuvanting a subunit COVID-19 vaccine to induce protective immunity. Nature, 2021, 594, 253-258.	27.8	253
9	Metabolic Phenotypes of Response to Vaccination in Humans. Cell, 2017, 169, 862-877.e17.	28.9	234
10	AMPK promotes mitochondrial biogenesis and function by phosphorylating the epigenetic factors DNMT1, RBBP7, and HAT1. Science Signaling, 2017, 10, .	3.6	170
11	Role of microRNA-23b in flow-regulation of Rb phosphorylation and endothelial cell growth. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 3234-3239.	7.1	160
12	Discovering and linking public omics data sets using the Omics Discovery Index. Nature Biotechnology, 2017, 35, 406-409.	17.5	159
13	Bioinformatics and Systems Biology of the Lipidome. Chemical Reviews, 2011, 111, 6452-6490.	47.7	154
14	Epigenetic Regulation: A New Frontier for Biomedical Engineers. Annual Review of Biomedical Engineering, 2017, 19, 195-219.	12.3	135
15	The single-cell epigenomic and transcriptional landscape of immunity to influenza vaccination. Cell, 2021, 184, 3915-3935.e21.	28.9	133
16	Enhancer-associated long non-coding RNA LEENE regulates endothelial nitric oxide synthase and endothelial function. Nature Communications, 2018, 9, 292.	12.8	129
17	T cell-inducing vaccine durably prevents mucosal SHIV infection even with lower neutralizing antibody titers. Nature Medicine, 2020, 26, 932-940.	30.7	124
18	Systems vaccinology: Enabling rational vaccine design with systems biological approaches. Vaccine, 2015, 33, 5294-5301.	3.8	108

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19	Systems biology analysis of longitudinal functional response of endothelial cells to shear stress. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 10990-10995.	7.1	95
20	A comprehensive classification system for lipids. European Journal of Lipid Science and Technology, 2005, 107, 337-364.	1.5	94
21	Kr $ ilde{A}^1\!\!/\!\!$ ppel-Like Factor 4 Regulation of Cholesterol-25-Hydroxylase and Liver X Receptor Mitigates Atherosclerosis Susceptibility. Circulation, 2017, 136, 1315-1330.	1.6	87
22	LIPID MAPS: Serving the next generation of lipid researchers with tools, resources, data, and training. Science Signaling, 2019, 12, .	3.6	87
23	Technical Variations in Low-Input RNA-seq Methodologies. Scientific Reports, 2014, 4, 3678.	3.3	75
24	Serum microRNAs explain discordance of non-alcoholic fatty liver disease in monozygotic and dizygotic twins: a prospective study. Gut, 2016, 65, 1546-1554.	12.1	75
25	miRNAs that Induce Human Cardiomyocyte Proliferation Converge on the Hippo Pathway. Cell Reports, 2018, 23, 2168-2174.	6.4	73
26	The Role of ARF6 in Biliary Atresia. PLoS ONE, 2015, 10, e0138381.	2.5	66
27	LIPID MAPS-Nature Lipidomics Gateway: An Online Resource for Students and Educators Interested in Lipids. Journal of Chemical Education, 2012, 89, 291-292.	2.3	55
28	LINCO0341 exerts an anti-inflammatory effect on endothelial cells by repressing VCAM1. Physiological Genomics, 2017, 49, 339-345.	2.3	53
29	A Complement–IL-4 Regulatory Circuit Controls Liver Regeneration. Journal of Immunology, 2012, 188, 641-648.	0.8	52
30	RefMet: a reference nomenclature for metabolomics. Nature Methods, 2020, 17, 1173-1174.	19.0	52
31	Right and left-sided colon cancers - specificity of molecular mechanisms in tumorigenesis and progression. BMC Cancer, 2020, 20, 317.	2.6	51
32	Three distinct cell populations express extracellular matrix proteins and increase in number during skeletal muscle fibrosis. American Journal of Physiology - Cell Physiology, 2017, 312, C131-C143.	4.6	49
33	Dedifferentiation and neuronal repression define familial Alzheimer's disease. Science Advances, 2020, 6, .	10.3	44
34	Longitudinal shear stress response in human endothelial cells to atheroprone and atheroprotective conditions. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	43
35	Topological and functional comparison of community detection algorithms in biological networks. BMC Bioinformatics, 2019, 20, 212.	2.6	41
36	GNPS Dashboard: collaborative exploration of mass spectrometry data in the web browser. Nature Methods, 2022, 19, 134-136.	19.0	35

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37	Systems analysis of transcriptional data provides insights into muscle's biological response to botulinum toxin. Muscle and Nerve, 2014, 50, 744-758.	2.2	33
38	MicroRNA-23b Regulates Cyclin-Dependent Kinase–Activating Kinase Complex Through Cyclin H Repression to Modulate Endothelial Transcription and Growth Under Flow. Arteriosclerosis, Thrombosis, and Vascular Biology, 2014, 34, 1437-1445.	2.4	33
39	Scope and limitations of yeast as a model organism for studying human tissue-specific pathways. BMC Systems Biology, 2015, 9, 96.	3.0	33
40	Gene-expression measurement: variance-modeling considerations for robust data analysis. Nature Immunology, 2012, 13, 199-203.	14.5	26
41	Modeling of Eicosanoid Fluxes Reveals Functional Coupling between Cyclooxygenases and Terminal Synthases. Biophysical Journal, 2014, 106, 966-975.	0.5	25
42	Dysregulated mechanisms underlying Duchenne muscular dystrophy from co-expression network preservation analysis. BMC Research Notes, 2015, 8, 182.	1.4	24
43	Immune Response in Severe and Non-Severe Coronavirus Disease 2019 (COVID-19) Infection: A Mechanistic Landscape. Frontiers in Immunology, 2021, 12, 738073.	4.8	24
44	Identification of AMP-activated protein kinase targets by a consensus sequence search of the proteome. BMC Systems Biology, 2015, 9, 13.	3.0	23
45	Systems Analysis of the Complement-Induced Priming Phase of Liver Regeneration. Journal of Immunology, 2016, 197, 2500-2508.	0.8	22
46	Loss of EGFR-ASAP1 signaling in metastatic and unresectable hepatoblastoma. Scientific Reports, 2016, 6, 38347.	3.3	20
47	Co-expression Network Approach Reveals Functional Similarities among Diseases Affecting Human Skeletal Muscle. Frontiers in Physiology, 2017, 8, 980.	2.8	19
48	Identification of therapeutic targets for glioblastoma by network analysis. Oncogene, 2016, 35, 608-620.	5.9	18
49	Enhanced B Cell Alloantigen Presentation and Its Epigenetic Dysregulation in Liver Transplant Rejection. American Journal of Transplantation, 2016, 16, 497-508.	4.7	17
50	The maternal blood lipidome is indicative of the pathogenesis of severe preeclampsia. Journal of Lipid Research, 2021, 62, 100118.	4.2	17
51	Identification of Crosstalk between Phosphoprotein Signaling Pathways in RAW 264.7 Macrophage Cells. PLoS Computational Biology, 2010, 6, e1000654.	3.2	15
52	Biliary-Atresia-Associated Mannosidase-1-Alpha-2 Gene Regulates Biliary and Ciliary Morphogenesis and Laterality. Frontiers in Physiology, 2020, 11, 538701.	2.8	13
53	Healthcare Innovations to Address the Challenges of the COVID-19 Pandemic. IEEE Journal of Biomedical and Health Informatics, 2022, 26, 3294-3302.	6.3	13
54	Plasmin Cascade Mediates Thrombotic Events in SARS-CoV-2 Infection via Complement and Platelet-Activating Systems. IEEE Open Journal of Engineering in Medicine and Biology, 2020, 1, 220-227.	2.3	12

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55	Information theoretic approach to complex biological network reconstruction: application to cytokine release in RAW 264.7 macrophages. BMC Systems Biology, 2014, 8, 77.	3.0	11
56	Computational Modeling of Competitive Metabolism between i‰3- and i‰6-Polyunsaturated Fatty Acids in Inflammatory Macrophages. Journal of Physical Chemistry B, 2016, 120, 8346-8353.	2.6	11
57	Steps Toward Minimal Reporting Standards for Lipidomics Mass Spectrometry in Biomedical Research Publications. Circulation Genomic and Precision Medicine, 2020, 13, e003019.	3.6	11
58	Statistical insights into major human muscular diseases. Human Molecular Genetics, 2014, 23, 3772-3778.	2.9	10
59	Domain architecture of aCaenorhabditis elegansAKAP suggests a novel AKAP function. FEBS Letters, 2000, 486, 107-111.	2.8	9
60	Endotype reversal as a novel strategy for screening drugs targeting familial Alzheimer's disease. Alzheimer's and Dementia, 2022, 18, 2117-2130.	0.8	9
61	Functional classification of skeletal muscle networks. II. Applications to pathophysiology. Journal of Applied Physiology, 2012, 113, 1902-1920.	2.5	8
62	Wnt Pathway Activation Increases Hypoxia Tolerance during Development. PLoS ONE, 2014, 9, e103292.	2.5	8
63	A Cybernetic Approach to Modeling Lipid Metabolism in Mammalian Cells. Processes, 2018, 6, 126.	2.8	7
64	Systems Biology and Medicine: A New Take on an Old Paradigm. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2009, 1, 1-3.	6.6	5
65	Replacing the Transfusion of 1–2 Units of Blood with Plasma Expanders that Increase Oxygen Delivery Capacity: Evidence from Experimental Studies. Journal of Functional Biomaterials, 2014, 5, 232-245.	4.4	5
66	Attractor Ranked Radial Basis Function Network: A Nonparametric Forecasting Approach for Chaotic Dynamic Systems. Scientific Reports, 2020, 10, 3780.	3.3	5
67	Cover Image, Volume 12, Issue 1. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2020, 12, e1473.	6.6	4
68	Influence of serological factors and BMI on the blood pressure/hematocrit association in healthy young men and women. Vascular Health and Risk Management, 2014, 10, 271.	2.3	3
69	Co-Expression Network Approach to Studying the Effects of Botulinum Neurotoxin-A. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2018, 15, 2009-2016.	3.0	3
70	Time varying causal network reconstruction of a mouse cell cycle. BMC Bioinformatics, 2019, 20, 294.	2.6	3
71	Systems Analysis of Biliary Atresia Through Integration of High-Throughput Biological Data. Frontiers in Physiology, 2020, 11, 966.	2.8	3
72	Cloudâ€based archived metabolomics data: A resource for inâ€source fragmentation/annotation, metaâ€analysis and systems biology. Analytical Science Advances, 2020, 1, 70-80.	2.8	3

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73	Histone Signatures Predict Therapeutic Efficacy in Breast Cancer. IEEE Open Journal of Engineering in Medicine and Biology, 2020, 1, 74-82.	2.3	3
74	Temporal mechanisms of myogenic specification in human induced pluripotent stem cells. Science Advances, $2021, 7, .$	10.3	3
75	Heme Oxygenase-1 at the Nexus of Endothelial Cell Fate Decision Under Oxidative Stress. Frontiers in Cell and Developmental Biology, 2021, 9, 702974.	3.7	3
76	Modular and mechanistic changes across stages of colorectal cancer. BMC Cancer, 2022, 22, 436.	2.6	3
77	Systems medicineâ€"viewed through the real and computing lenses. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2010, 2, 383-384.	6.6	2
78	Systems biology—old wine in a new bottle or is the bottle changing the wine?. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2010, 2, 1-2.	6.6	2
79	Drug discovery and the maze of biological complexity: an editorial essay. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2014, 6, 225-226.	6.6	2
80	Mechanisms of Impaired Lung Development and Ciliation in Mannosidase-1-Alpha-2 (Man1a2) Mutants. Frontiers in Physiology, 2021, 12, 658518.	2.8	2
81	Bioinformatics of Cellular Signalling. Novartis Foundation Symposium, 2002, 247, 104-118.	1.1	1
82	Systems Biology and Medicine—Metaâ€Issues and Frameworks. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2010, 2, i-ii.	6.6	0
83	A Clustering Approach to Identify Intergenic Non-coding RNA in Mouse Macrophages. , 2010, , .		0
84	Evolution of Bioengineering at UCSD: Opening New Vistas. IEEE Pulse, 2012, 3, 49-55.	0.3	0
85	Signed Differential Coâ€Expression Network Analysis Suggests Differential Regulation of SP/KLF Family of Transcription Factors in Dilated Cardiomyopathy. FASEB Journal, 2018, 32, .	0.5	0
86	The Mechanistic Metamorphosis. WIREs Mechanisms of Disease, 2021, 13, e1517.	3.3	0
87	Methadone disrupts early synaptic development in human cortical organoids. FASEB Journal, 2022, 36, .	0.5	О