

# Kumar Selvarajoo

## List of Publications by Year in descending order

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Version: 2024-02-01

61  
papers

1,033  
citations

393982

19  
h-index

454577

30  
g-index

77  
all docs

77  
docs citations

77  
times ranked

1174  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identifying toggle genes from transcriptome-wide scatter: A new perspective for biological regulation. <i>Genomics</i> , 2022, 114, 215-228.	1.3	6
2	The transformation of our food system using cellular agriculture: What lies ahead and who will lead it?. <i>Trends in Food Science and Technology</i> , 2022, 127, 368-376.	7.8	12
3	The need for integrated systems biology approaches for biotechnological applications. <i>Biotechnology Notes</i> , 2021, 2, 39-43.	0.7	4
4	Searching for unifying laws of general adaptation syndrome. <i>Physics of Life Reviews</i> , 2021, 37, 97-99.	1.5	1
5	GeneCloudOmics: A Data Analytic Cloud Platform for High-Throughput Gene Expression Analysis. <i>Frontiers in Bioinformatics</i> , 2021, 1, .	1.0	4
6	Systems Biology to Understand and Regulate Human Retroviral Proinflammatory Response. <i>Frontiers in Immunology</i> , 2021, 12, 736349.	2.2	5
7	ScatLay: utilizing transcriptome-wide noise for identifying and visualizing differentially expressed genes. <i>Scientific Reports</i> , 2020, 10, 17483.	1.6	6
8	Systems biology approaches integrated with artificial intelligence for optimized metabolic engineering. <i>Metabolic Engineering Communications</i> , 2020, 11, e00149.	1.9	46
9	Attractor Concepts to Evaluate the Transcriptome-wide Dynamics Guiding Anaerobic to Aerobic State Transition in <i>Escherichia coli</i> . <i>Scientific Reports</i> , 2020, 10, 5878.	1.6	12
10	Systems Biology Approaches for Understanding Biofilm Response. <i>ACS Symposium Series</i> , 2020, , 9-29.	0.5	0
11	ABioTrans: A Biostatistical Tool for Transcriptomics Analysis. <i>Frontiers in Genetics</i> , 2019, 10, 499.	1.1	7
12	Defining rules for cancer cell proliferation in TRAIL stimulation. <i>Npj Systems Biology and Applications</i> , 2019, 5, 5.	1.4	7
13	Long-range order and short-range disorder in <i>Saccharomyces cerevisiae</i> biofilm. <i>Engineering Biology</i> , 2019, 3, 12-19.	0.8	7
14	Searching for simple rules in <i>Pseudomonas aeruginosa</i> biofilm formation. <i>BMC Research Notes</i> , 2019, 12, 763.	0.6	3
15	Large-scale free network organisation is likely key for biofilm phase transition. <i>Engineering Biology</i> , 2019, 3, 67-71.	0.8	4
16	Complexity of Biochemical and Genetic Responses Reduced Using Simple Theoretical Models. <i>Methods in Molecular Biology</i> , 2018, 1702, 171-201.	0.4	10
17	Hints from Information Theory for Analyzing Dynamic and High-Dimensional Biological Data. <i>RNA Technologies</i> , 2018, , 313-336.	0.2	1
18	Order Parameter in Bacterial Biofilm Adaptive Response. <i>Frontiers in Microbiology</i> , 2018, 9, 1721.	1.5	8

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19	A systems biology approach to overcome TRAIL resistance in cancer treatment. <i>Progress in Biophysics and Molecular Biology</i> , 2017, 128, 142-154.	1.4	24
20	Can the second law of thermodynamics hold in cell cultures?. <i>Frontiers in Genetics</i> , 2015, 6, 262.	1.1	4
21	Systems Biology Strategy Reveals PKC $\zeta$ is Key for Sensitizing TRAIL-Resistant Human Fibrosarcoma. <i>Frontiers in Immunology</i> , 2015, 5, 659.	2.2	12
22	The reduction of gene expression variability from single cells to populations follows simple statistical laws. <i>Genomics</i> , 2015, 105, 137-144.	1.3	33
23	Tracking global gene expression responses in T cell differentiation. <i>Gene</i> , 2015, 569, 259-266.	1.0	20
24	Measuring merit: Take the risk. <i>Science</i> , 2015, 347, 139-140.	6.0	2
25	Advances in systems immunology and cancer. <i>Frontiers in Physiology</i> , 2014, 5, 249.	1.3	1
26	Beyond MyD88 and TRIF Pathways in Toll-Like Receptor Signaling. <i>Frontiers in Immunology</i> , 2014, 5, 70.	2.2	61
27	Parameter-less approaches for interpreting dynamic cellular response. <i>Journal of Biological Engineering</i> , 2014, 8, 23.	2.0	8
28	Transcriptome-wide Variability in Single Embryonic Development Cells. <i>Scientific Reports</i> , 2014, 4, 7137.	1.6	66
29	Physical Laws Shape Biology. <i>Science</i> , 2013, 339, 646-646.	6.0	11
30	A systems biology approach to suppress TNF-induced proinflammatory gene expressions. <i>Cell Communication and Signaling</i> , 2013, 11, 84.	2.7	28
31	Uncertainty and certainty in cellular dynamics. <i>Frontiers in Genetics</i> , 2013, 4, 68.	1.1	4
32	Non-genetic adaptive dynamics for cellular robustness. <i>Frontiers in Genetics</i> , 2013, 4, 287.	1.1	3
33	Basics of the Mammalian Immune System. <i>Systems Biology</i> , 2013, , 25-33.	0.1	0
34	Stochasticity and Variability: Insights from Single-Cell Dynamics. <i>Systems Biology</i> , 2013, , 105-116.	0.1	0
35	Systems Biology of Population Cell Response. <i>Systems Biology</i> , 2013, , 1-11.	0.1	0
36	Investigating the TLR3 Signaling Dynamics. <i>Systems Biology</i> , 2013, , 65-74.	0.1	0

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37	Inferring Novel Features of the TLR4 Pathways. <i>Systems Biology</i> , 2013, , 35-52.	0.1	0
38	Investigating Single-Cell Stochasticity in TRAIL Signaling. <i>Systems Biology</i> , 2013, , 117-124.	0.1	0
39	Finding Chaos in Biology. <i>Systems Biology</i> , 2013, , 131-140.	0.1	0
40	Finding Self-organization from the Dynamic Gene Expressions of Innate Immune Responses. <i>Frontiers in Physiology</i> , 2012, 3, 192.	1.3	7
41	Is central dogma a global property of cellular information flow?. <i>Frontiers in Physiology</i> , 2012, 3, 439.	1.3	28
42	Investigation of stochasticity in TRAIL signaling cancer model. , 2012, , .		0
43	The Recognition of Chaos in Host-Pathogen Response. <i>Frontiers in Physiology</i> , 2012, 3, 7.	1.3	1
44	Understanding multimodal biological decisions from single cell and population dynamics. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , 2012, 4, 385-399.	6.6	24
45	Emergence of macroscopic simplicity from the Tumor Necrosis Factor-alpha signaling dynamics. <i>Nature Precedings</i> , 2011, , .	0.1	1
46	Macroscopic law of conservation revealed in the population dynamics of Toll-like receptor signaling. <i>Cell Communication and Signaling</i> , 2011, 9, 9.	2.7	24
47	Enhancing apoptosis in TRAIL-resistant cancer cells using fundamental response rules. <i>Scientific Reports</i> , 2011, 1, 144.	1.6	26
48	Signaling Flux Redistribution concept can switch survival to apoptosis in cancer cells. <i>Nature Precedings</i> , 2010, , .	0.1	0
49	Genetic vehicle comprising majority of lowly expressed genes guides cell fate decision. <i>Nature Precedings</i> , 2010, , .	0.1	0
50	Collective Dynamics of Specific Gene Ensembles Crucial for Neutrophil Differentiation: The Existence of Genome Vehicles Revealed. <i>PLoS ONE</i> , 2010, 5, e12116.	1.1	23
51	Emergent Genome-Wide Control in Wildtype and Genetically Mutated Lipopolysaccharides-Stimulated Macrophages. <i>PLoS ONE</i> , 2009, 4, e4905.	1.1	45
52	CAN COMPLEX CELLULAR PROCESSES BE GOVERNED BY SIMPLE LINEAR RULES?. <i>Journal of Bioinformatics and Computational Biology</i> , 2009, 07, 243-268.	0.3	20
53	Local and global responses in complex gene regulation networks. <i>Physica A: Statistical Mechanics and Its Applications</i> , 2009, 388, 1738-1746.	1.2	40
54	In Silico Models for Metabolic Systems Engineering. , 2009, , .		1

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55	Predicting Novel Features of Toll-Like Receptor 3 Signaling in Macrophages. PLoS ONE, 2009, 4, e4661.	1.1	25
56	Signaling Flux Redistribution at Toll-Like Receptor Pathway Junctions. PLoS ONE, 2008, 3, e3430.	1.1	43
57	Toll-like receptor signal transduction. Experimental and Molecular Medicine, 2007, 39, 421-438.	3.2	211
58	Gene expression waves. FEBS Journal, 2007, 274, 2878-2886.	2.2	38
59	Systematic Determination of Biological Network Topology: Nonintegral Connectivity Method (NICM). , 2007, , 449-471.		4
60	Sequential Logic Model Deciphers Dynamic Transcriptional Control of Gene Expressions. PLoS ONE, 2007, 2, e776.	1.1	7
61	Discovering differential activation machinery of the Toll-like receptor 4 signaling pathways in MyD88 knockouts. FEBS Letters, 2006, 580, 1457-1464.	1.3	30