

Sam Thiagalingam

List of Publications by Year in descending order

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77
papers

8,018
citations

147726

31
h-index

233338

45
g-index

84
all docs

84
docs citations

84
times ranked

8616
citing authors

#	ARTICLE	IF	CITATIONS
1	Pathogenic histone modifications in schizophrenia are targets for therapy. , 2021, , 309-319.		1
2	Cataloging recent advances in epigenetic alterations in major mental disorders and autism. Epigenomics, 2021, 13, 1231-1245.	1.0	5
3	Targeting RICTOR Sensitizes SMAD4-Negative Colon Cancer to Irinotecan. Molecular Cancer Research, 2020, 18, 414-423.	1.5	12
4	Epigenetic memory in development and disease: Unraveling the mechanism. Biochimica Et Biophysica Acta: Reviews on Cancer, 2020, 1873, 188349.	3.3	25
5	MicroRNA-4417 is a tumor suppressor and prognostic biomarker for triple-negative breast cancer. Cancer Biology and Therapy, 2019, 20, 1113-1120.	1.5	19
6	Activin A Signaling Regulates IL13RÎ±2 Expression to Promote Breast Cancer Metastasis. Frontiers in Oncology, 2019, 9, 32.	1.3	33
7	Aberrant transcriptomes and DNA methylomes define pathways that drive pathogenesis and loss of brain laterality/asymmetry in schizophrenia and bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 138-149.	1.1	31
8	Methamphetamine-induced psychosis is associated with DNA hypomethylation and increased expression of <i>AKT1</i> and key dopaminergic genes. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 1180-1189.	1.1	18
9	SDPR functions as a metastasis suppressor in breast cancer by promoting apoptosis. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 638-643.	3.3	66
10	Tumor Cell-Derived Periostin Regulates Cytokines That Maintain Breast Cancer Stem Cells. Molecular Cancer Research, 2016, 14, 103-113.	1.5	46
11	Antipsychotic drugs attenuate aberrant DNA methylation of <i>DTNBP1</i> (dysbindin) promoter in saliva and postmortem brain of patients with schizophrenia and Psychotic bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 687-696.	1.1	64
12	TGFÎ² and BMP signaling in cancer. , 2015, , 204-221.		1
13	An update on the epigenetics of psychotic diseases and autism. Epigenomics, 2015, 7, 427-449.	1.0	57
14	Targeting IL13RÎ±2 activates STAT6-TP63 pathway to suppress breast cancer lung metastasis. Breast Cancer Research, 2015, 17, 98.	2.2	76
15	Dietary and environmental influences on the genomic and epigenomic codes in cancer. , 2015, , 154-168.		1
16	DNA hypermethylation of serotonin transporter gene promoter in drug naïve patients with schizophrenia. Schizophrenia Research, 2014, 152, 373-380.	1.1	93
17	Pathogenic Histone Modifications in Schizophrenia are Targets for Therapy. , 2014, , 241-251.		5
18	Integrin Signaling in Mammary Epithelial Cells and Breast Cancer. ISRN Oncology, 2012, 2012, 1-9.	2.1	31

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19	Epigenetic dysregulation of HTR2A in the brain of patients with schizophrenia and bipolar disorder. Schizophrenia Research, 2011, 129, 183-190.	1.1	170
20	DNA hypomethylation of MB-COMT promoter in the DNA derived from saliva in schizophrenia and bipolar disorder. Journal of Psychiatric Research, 2011, 45, 1432-1438.	1.5	155
21	Hypomethylation of the serotonin receptor type 2A Gene (HTR2A) at T102C polymorphic site in DNA derived from the saliva of patients with schizophrenia and bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 536-545.	1.1	104
22	Can the schizophrenia epigenome provide clues for the molecular basis of pathogenesis?. Epigenomics, 2011, 3, 679-683.	1.0	17
23	Smad4 Inactivation Promotes Malignancy and Drug Resistance of Colon Cancer. Cancer Research, 2011, 71, 998-1008.	0.4	170
24	Smad Signaling Is Required to Maintain Epigenetic Silencing during Breast Cancer Progression. Cancer Research, 2010, 70, 968-978.	0.4	162
25	hBub1 deficiency triggers a novel p53 mediated early apoptotic checkpoint pathway in mitotic spindle damaged cells. Cancer Biology and Therapy, 2009, 8, 627-635.	1.5	11
26	hBub1 negatively regulates p53 mediated early cell death upon mitotic checkpoint activation. Cancer Biology and Therapy, 2009, 8, 636-644.	1.5	11
27	Epigenetic and pharmacoeconomic studies of major psychoses and potentials for therapeutics. Pharmacogenomics, 2008, 9, 1809-1823.	0.6	44
28	Epigenetic Alterations of the Dopaminergic System in Major Psychiatric Disorders. Methods in Molecular Biology, 2008, 448, 187-212.	0.4	62
29	Epigenetic Modulation of Reelin Function in Schizophrenia and Bipolar Disorder. , 2008, , 365-384.		4
30	The Cancer Epigenome. , 2008, , 97-113.		1
31	DNA Methylation Profiles as Prognostic Markers for Cancer. , 2008, , 333-346.		0
32	Aberrant activation of β -catenin promotes genomic instability and oncogenic effects during tumor progression. Cancer Biology and Therapy, 2007, 6, 1638-1643.	1.5	33
33	A Cascade of Modules of a Network Defines Cancer Progression. Cancer Research, 2006, 66, 7379-7385.	0.4	27
34	Hypomethylation of MB-COMT promoter is a major risk factor for schizophrenia and bipolar disorder. Human Molecular Genetics, 2006, 15, 3132-3145.	1.4	433
35	Hypermethylation of the reelin (RELN) promoter in the brain of schizophrenic patients: A preliminary report. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2005, 134B, 60-66.	1.1	347
36	Loss of Heterozygosity Patterns Provide Fingerprints for Genetic Heterogeneity in Multistep Cancer Progression of Tobacco Smoke-Induced Non-Small Cell Lung Cancer. Cancer Research, 2005, 65, 1664-1669.	0.4	59

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37	Differential DNA Hypermethylation of Critical Genes Mediates the Stage-Specific Tobacco Smoke-Induced Neoplastic Progression of Lung Cancer. <i>Clinical Cancer Research</i> , 2005, 11, 2466-2470.	3.2	140
38	Genetics and Epigenetics in Major Psychiatric Disorders. <i>Molecular Diagnosis and Therapy</i> , 2005, 5, 149-160.	3.3	134
39	Elucidation of Epigenetic Inactivation of SMAD8 in Cancer Using Targeted Expressed Gene Display. <i>Cancer Research</i> , 2004, 64, 1639-1646.	0.4	36
40	Histone Deacetylases: Unique Players in Shaping the Epigenetic Histone Code. <i>Annals of the New York Academy of Sciences</i> , 2003, 983, 84-100.	1.8	635
41	Loss of heterozygosity as a predictor to map tumor suppressor genes in cancer: molecular basis of its occurrence. <i>Current Opinion in Oncology</i> , 2002, 14, 65-72.	1.1	89
42	Molecular Detection of Smad2/Smad4 Alterations in Colorectal Tumors. , 2001, 50, 149-165.		3
43	Mechanisms underlying losses of heterozygosity in human colorectal cancers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 2698-2702.	3.3	194
44	Is a p53-Regulated Inhibitor of G2/M Progression. <i>Molecular Cell</i> , 1997, 1, 3-11.	4.5	1,153
45	Homeosis and polyposis: A tale from the mouse. <i>BioEssays</i> , 1997, 19, 551-555.	1.2	7
46	Evaluation of candidate tumour suppressor genes on chromosome 18 in colorectal cancers. <i>Nature Genetics</i> , 1996, 13, 343-346.	9.4	580
47	Mad-related genes in the human. <i>Nature Genetics</i> , 1996, 13, 347-349.	9.4	359
48	PAK1, a gene that can regulate p53 activity in yeast.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1995, 92, 6062-6066.	3.3	26
49	p53 tagged sites from human genomic DNA. <i>Human Molecular Genetics</i> , 1994, 3, 1537-1542.	1.4	174
50	Sequence-specific transcriptional activation is essential for growth suppression by p53.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1994, 91, 1998-2002.	3.3	368
51	Oncoprotein MDM2 conceals the activation domain of tumour suppressor p53. <i>Nature</i> , 1993, 362, 857-860.	13.7	1,407
52	ATPase activity of the UvrA and UvrAB protein complexes of the <i>Escherichia coli</i> UvrABC endonuclease. <i>Nucleic Acids Research</i> , 1989, 17, 4145-4159.	6.5	78
53	Events at DNA replication origins and genome stability. , 0, , 35-55.		0
54	Regulation and dysregulation of protein synthesis in cancer cells. , 0, , 70-92.		1

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55	Role of network biology and network medicine in early detection of cancer. , 0, , 457-463.		0
56	Application of bioinformatics to analyze the expression of tissue-specific and housekeeping genes in cancer. , 0, , 20-34.		0
57	Tumor microenvironment: blood vascular system in cancer metastasis. , 0, , 309-322.		0
58	PI3K pathway in cancer. , 0, , 193-203.		0
59	The Wnt signaling network in cancer. , 0, , 222-255.		0
60	Genomic instability and carcinogenesis. , 0, , 93-112.		0
61	Molecular links between inflammation and cancer. , 0, , 273-281.		3
62	MicroRNA epigenetic systems and cancer. , 0, , 134-153.		1
63	Cancer metastasis. , 0, , 282-294.		1
64	Cancer metabolism. , 0, , 295-308.		1
65	The role of growth factor-induced changes in cell fate in prostate cancer progression. , 0, , 361-376.		1
66	Systems biology of cancer progression. , 0, , 1-6.		0
67	Lessons from cancer genome sequencing. , 0, , 7-19.		0
68	Systems biology approaches bring new insights in the understanding of global gene regulatory mechanisms and their deregulation in cancer. , 0, , 56-69.		0
69	Epigenomic code. , 0, , 113-133.		0
70	RAS signaling networks. , 0, , 183-192.		0
71	Apoptotic pathways and cancer. , 0, , 256-272.		0
72	Biology of human stomach cancer. , 0, , 386-408.		0

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73	Deregulated signaling networks in lung cancer. , 0, , 421-442.		0
74	Modular signaling in hematopoietic malignancies. , 0, , 443-456.		0
75	Systems biology in cancer biomarkers for early detection, diagnosis, and prognosis. , 0, , 464-472.		0
76	Prognosis of cancer. , 0, , 473-498.		0
77	Cancer pharmacogenomics: challenges, promises, and its application to cancer drug discovery. , 0, , 499-517.		0