

Samuel Granjeaud

List of Publications by Year in descending order

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79
papers

4,083
citations

117453

34
h-index

118652

62
g-index

83
all docs

83
docs citations

83
times ranked

6039
citing authors

#	ARTICLE	IF	CITATIONS
1	Inducible Antibacterial Defense System in <i>C. elegans</i> . <i>Current Biology</i> , 2002, 12, 1209-1214.	1.8	417
2	The Strategy of T Cell Antigen-presenting Cell Encounter in Antigen-draining Lymph Nodes Revealed by Imaging of Initial T Cell Activation. <i>Journal of Experimental Medicine</i> , 2003, 198, 715-724.	4.2	205
3	Differential Gene Expression in the Murine Thymus Assayed by Quantitative Hybridization of Arrayed cDNA Clones. <i>Genomics</i> , 1995, 29, 207-216.	1.3	185
4	Sensitivity Issues in DNA Array-Based Expression Measurements and Performance of Nylon Microarrays for Small Samples. <i>Human Molecular Genetics</i> , 1999, 8, 1715-1722.	1.4	172
5	Cancer-associated fibroblast-derived annexin A6+ extracellular vesicles support pancreatic cancer aggressiveness. <i>Journal of Clinical Investigation</i> , 2016, 126, 4140-4156.	3.9	169
6	Gene expression profiling of primary breast carcinomas using arrays of candidate genes. <i>Human Molecular Genetics</i> , 2000, 9, 2981-2991.	1.4	168
7	Highly effective NK cells are associated with good prognosis in patients with metastatic prostate cancer. <i>Oncotarget</i> , 2015, 6, 14360-14373.	0.8	164
8	Inherent and Tumor-Driven Immune Tolerance in the Prostate Microenvironment Impairs Natural Killer Cell Antitumor Activity. <i>Cancer Research</i> , 2016, 76, 2153-2165.	0.4	154
9	Small lymphocytic lymphoma, marginal zone B-cell lymphoma, and mantle cell lymphoma exhibit distinct gene-expression profiles allowing molecular diagnosis. <i>Blood</i> , 2004, 103, 2727-2737.	0.6	127
10	Gene expression profiling of multiple myeloma reveals molecular portraits in relation to the pathogenesis of the disease. <i>Blood</i> , 2003, 101, 4998-5006.	0.6	124
11	Gene expression profiles of poor-prognosis primary breast cancer correlate with survival. <i>Human Molecular Genetics</i> , 2002, 11, 863-872.	1.4	117
12	Postoperative serum proteomic profiles may predict metastatic relapse in high-risk primary breast cancer patients receiving adjuvant chemotherapy. <i>Oncogene</i> , 2006, 25, 981-989.	2.6	112
13	Expression profiling: DNA arrays in many guises. <i>BioEssays</i> , 1999, 21, 781-790.	1.2	102
14	Gene expression profiling defines molecular subtypes of classical Hodgkin's disease. <i>Oncogene</i> , 2002, 21, 3095-3102.	2.6	91
15	LIF Drives Neural Remodeling in Pancreatic Cancer and Offers a New Candidate Biomarker. <i>Cancer Research</i> , 2018, 78, 909-921.	0.4	83
16	Transcriptional Profiling Reveals Coordinated Up-Regulation of Oxidative Metabolism Genes in Thyroid Oncocytic Tumors. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2004, 89, 994-1005.	1.8	80
17	Multiplex messenger assay: simultaneous, quantitative measurement of expression of many genes in the context of T cell activation. <i>Nucleic Acids Research</i> , 1996, 24, 1435-1442.	6.5	72
18	Atomic resolution on small three-dimensional metal clusters by STM. <i>Surface Science</i> , 1997, 391, 19-26.	0.8	71

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19	Vanin genes are clustered (human 6q22-24 and mouse 10A2B1) and encode isoforms of pantetheinase ectoenzymes. <i>Immunogenetics</i> , 2001, 53, 296-306.	1.2	68
20	Follicular B Lymphomas Generate Regulatory T Cells via the ICOS/ICOSL Pathway and Are Susceptible to Treatment by Anti-ICOS/ICOSL Therapy. <i>Cancer Research</i> , 2016, 76, 4648-4660.	0.4	65
21	Differential gene expression in CD3e- and RAG1-deficient thymuses: definition of a set of genes potentially involved in thymocyte maturation. <i>Immunogenetics</i> , 1999, 50, 255-270.	1.2	64
22	Plasmodium falciparum proteome changes in response to doxycycline treatment. <i>Malaria Journal</i> , 2010, 9, 141.	0.8	62
23	Expression scanning of an array of growth control genes in human tumor cell lines. <i>Oncogene</i> , 1999, 18, 3905-3912.	2.6	57
24	Reconstitution of Natural Killer Cells in HLA-Matched HSCT after Reduced-Intensity Conditioning: Impact on Clinical Outcome. <i>Biology of Blood and Marrow Transplantation</i> , 2015, 21, 429-439.	2.0	55
25	Entropy Measures Quantify Global Splicing Disorders in Cancer. <i>PLoS Computational Biology</i> , 2008, 4, e1000011.	1.5	53
26	Natural Killer Defective Maturation Is Associated with Adverse Clinical Outcome in Patients with Acute Myeloid Leukemia. <i>Frontiers in Immunology</i> , 2017, 8, 573.	2.2	47
27	An ESTs description of the new Vanin gene family conserved from fly to human. <i>Immunogenetics</i> , 1999, 49, 964-972.	1.2	45
28	NKG2C ⁺ memory-like NK cells contribute to the control of HIV viremia during primary infection: Optprim ⁺ ANRS 147. <i>Clinical and Translational Immunology</i> , 2017, 6, e150.	1.7	42
29	From hybridization image to numerical values: a practical, high throughput quantification system for high density filter hybridizations. <i>Genetic Analysis, Techniques and Applications</i> , 1996, 12, 151-162.	1.5	41
30	Gene profiling reveals specific oncogenic mechanisms and signaling pathways in oncocytic and papillary thyroid carcinoma. <i>Oncogene</i> , 2005, 24, 4155-4161.	2.6	40
31	Molecular differences between the divergent responses of ovalbumin-specific CD4 T cells to alum-precipitated ovalbumin compared to ovalbumin expressed by Salmonella. <i>Molecular Immunology</i> , 2008, 45, 3558-3566.	1.0	39
32	Ultrahigh vacuum and air observations of Pd clusters grown on clean graphite. <i>Journal of Vacuum Science & Technology an Official Journal of the American Vacuum Society B, Microelectronics Processing and Phenomena</i> , 1991, 9, 804.	1.6	37
33	Feature extraction and signal processing for nylon DNA microarrays. <i>BMC Genomics</i> , 2004, 5, 38.	1.2	37
34	Kinetic Analysis of Mouse Brain Proteome Alterations Following Chikungunya Virus Infection before and after Appearance of Clinical Symptoms. <i>PLoS ONE</i> , 2014, 9, e91397.	1.1	35
35	HVEM has a broader expression than PD-L1 and constitutes a negative prognostic marker and potential treatment target for melanoma. <i>Oncolmmunology</i> , 2019, 8, e1665976.	2.1	35
36	NKp30 expression is a prognostic immune biomarker for stratification of patients with intermediate-risk acute myeloid leukemia. <i>Oncotarget</i> , 2017, 8, 49548-49563.	0.8	34

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37	Prognosis of Breast Cancer and Gene Expression Profiling Using DNA Arrays. <i>Annals of the New York Academy of Sciences</i> , 2002, 975, 217-231.	1.8	33
38	A Mature NK Profile at the Time of HIV Primary Infection Is Associated with an Early Response to cART. <i>Frontiers in Immunology</i> , 2017, 8, 54.	2.2	30
39	CXCR5 and ICOS expression identifies a CD8 T-cell subset with TFH features in Hodgkin lymphomas. <i>Blood Advances</i> , 2018, 2, 1889-1900.	2.5	30
40	Two New Neutrophil Subsets Define a Discriminating Sepsis Signature. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2022, 205, 46-59.	2.5	30
41	High-dimensional mass cytometry analysis of NK cell alterations in AML identifies a subgroup with adverse clinical outcome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	29
42	TranscriptomeBrowser: A Powerful and Flexible Toolbox to Explore Productively the Transcriptional Landscape of the Gene Expression Omnibus Database. <i>PLoS ONE</i> , 2008, 3, e4001.	1.1	27
43	Salivette, a relevant saliva sampling device for SARS-CoV-2 detection. <i>Journal of Oral Microbiology</i> , 2021, 13, 1920226.	1.2	26
44	The disparate nature of "intergenic" polyadenylation sites. <i>Rna</i> , 2006, 12, 1794-1801.	1.6	25
45	Immunomodulatory Drugs Exert Anti-Leukemia Effects in Acute Myeloid Leukemia by Direct and Immunostimulatory Activities. <i>Frontiers in Immunology</i> , 2018, 9, 977.	2.2	25
46	Increased NK Cell Maturation in Patients with Acute Myeloid Leukemia. <i>Frontiers in Immunology</i> , 2015, 6, 564.	2.2	24
47	Anopheles salivary gland proteomes from major malaria vectors. <i>BMC Genomics</i> , 2012, 13, 614.	1.2	23
48	JAM-C Identifies Src Family Kinase-Activated Leukemia-Initiating Cells and Predicts Poor Prognosis in Acute Myeloid Leukemia. <i>Cancer Research</i> , 2017, 77, 6627-6640.	0.4	23
49	Overexpression of dominant-negative Ikaros 6 protein is restricted to a subset of B common adult acute lymphoblastic leukemias that express high levels of the CD34 antigen. <i>The Hematology Journal</i> , 2003, 4, 104-109.	2.0	23
50	STM/TEM comparative study of Pd clusters epitaxially grown on highly oriented pyrolytic graphite. <i>Microscopy Microanalysis Microstructures</i> , 1993, 4, 409-418.	0.4	23
51	Plasmodium falciparum infection-induced changes in erythrocyte membrane proteins. <i>Parasitology Research</i> , 2012, 110, 545-556.	0.6	21
52	Altered Protein Networks and Cellular Pathways in Severe West Nile Disease in Mice. <i>PLoS ONE</i> , 2013, 8, e68318.	1.1	20
53	Achieving automated scorpion venom mass fingerprinting (VMF) in the nanogram range. <i>Toxicon</i> , 2013, 69, 211-218.	0.8	17
54	Development of parallel reaction monitoring (PRM)-based quantitative proteomics applied to HER2-Positive breast cancer. <i>Oncotarget</i> , 2018, 9, 33762-33777.	0.8	17

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55	Structure and morphology of nanometer-sized Pd clusters grown at high temperature on natural graphite single crystals. <i>EPJ Applied Physics</i> , 2001, 13, 23-30.	0.3	16
56	Temporal trends in Human T-Lymphotropic virus 1 (HTLV-1) associated myelopathy/tropical spastic paraparesis (HAM/TSP) incidence in Martinique over 25 years (1986-2010). <i>PLoS Neglected Tropical Diseases</i> , 2018, 12, e0006304.	1.3	14
57	Cerebrospinal Fluid Biomarker Candidates Associated with Human WNV Neuroinvasive Disease. <i>PLoS ONE</i> , 2014, 9, e93637.	1.1	13
58	Exploratory Study on Application of MALDI-TOF-MS to Detect SARS-CoV-2 Infection in Human Saliva. <i>Journal of Clinical Medicine</i> , 2022, 11, 295.	1.0	13
59	'LABNOTE', a laboratory notebook system designed for academic genomics groups. <i>Nucleic Acids Research</i> , 1999, 27, 601-607.	6.5	11
60	Multiparametric cytometry for exploration of complex cellular dynamics. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2012, 81A, 332-342.	1.1	9
61	Tubulin isoforms identified in the brain by MALDI in-source decay. <i>Journal of Proteomics</i> , 2013, 79, 172-179.	1.2	9
62	Multi-parametric cytometry from a complex cellular sample: Improvements and limits of manual versus computational-based interactive analyses. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2016, 89, 480-490.	1.1	9
63	Time-limited alterations in cortical activity of a knock-in mouse model of <i>KCNQ2</i> -related developmental and epileptic encephalopathy. <i>Journal of Physiology</i> , 2022, 600, 2429-2460.	1.3	9
64	Genomic organization and the tissue distribution of alternatively spliced isoforms of the mouse <i>Spatial</i> gene. <i>BMC Genomics</i> , 2004, 5, 41.	1.2	7
65	Scanning tunneling investigation of DNA structures involved in gene regulation. <i>Journal of Vacuum Science & Technology an Official Journal of the American Vacuum Society B, Microelectronics Processing and Phenomena</i> , 1994, 12, 1521.	1.6	6
66	Quantification of Immune Variables from Liquid Biopsy in Breast Cancer Patients Links β 2-T Cell Alterations with Lymph Node Invasion. <i>Cancers</i> , 2021, 13, 441.	1.7	6
67	Optimization and Standardization of Human Saliva Collection for MALDI-TOF MS. <i>Diagnostics</i> , 2021, 11, 1304.	1.3	6
68	Mouth Washing Impaired SARS-CoV-2 Detection in Saliva. <i>Diagnostics</i> , 2021, 11, 1509.	1.3	5
69	Scanning tunneling microscopy study of a DNA fragment of known size and sequence. <i>Microscopy Microanalysis Microstructures</i> , 1994, 5, 47-56.	0.4	5
70	Expression profiling in mouse fetal thymus reveals clusters of coordinately expressed genes that mark individual stages of T-cell ontogeny. <i>Immunogenetics</i> , 2002, 54, 469-478.	1.2	4
71	MicroArray Facility: a laboratory information management system with extended support for Nylon based technologies. <i>BMC Genomics</i> , 2006, 7, 240.	1.2	4
72	Gene expression profiling of breast carcinomas using Nylon DNA arrays. <i>Comptes Rendus - Biologies</i> , 2003, 326, 1031-1039.	0.1	3

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73	HVEM: A novel cosignaling molecule of major interest in melanoma.. Journal of Clinical Oncology, 2017, 35, e14591-e14591.	0.8	3
74	Evolutionary and expression analyses reveal a pattern of ancient duplications and functional specializations in the diversification of the Downstream of Kinase (DOK) genes. Developmental and Comparative Immunology, 2018, 84, 193-198.	1.0	2
75	Differential mRNA Expression in Untreated and TNF- α Elicited Murine Dendritic Cells Precursors. Advances in Experimental Medicine and Biology, 1997, 417, 467-473.	0.8	2
76	Transcriptional Expression Analysis of T-Cell Activation by Multiplex Messenger Assay (MMA). , 2000, 134, 337-353.		1
77	Djeen (Database for Joomla!â€™s Extensible Engine): a research information management system for flexible multi-technology project administration. BMC Research Notes, 2013, 6, 223.	0.6	0
78	Follicular Lymphoma B Cells Generate Functional Regulatory T Cells Via ICOS/ICOSL Pathway and Are Inhibited By Intratumoral Tregs. Blood, 2015, 126, 5018-5018.	0.6	0
79	Abstract A61: CAF-derived ANXA6+-exosomes support pancreatic cancer aggressiveness and serve as a circulating biomarker. , 2016, , .		0