

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
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83
all docs

83
docs citations

83
times ranked

6039
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	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
3	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
4	Quantification of Immune Variables from Liquid Biopsy in Breast Cancer Patients Links $\gamma\delta$ T Cell Alterations with Lymph Node Invasion. <i>Cancers</i> , 2021, 13, 441.	1.7	6
5	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
6	Optimization and Standardization of Human Saliva Collection for MALDI-TOF MS. <i>Diagnostics</i> , 2021, 11, 1304.	1.3	6
7	Mouth Washing Impaired SARS-CoV-2 Detection in Saliva. <i>Diagnostics</i> , 2021, 11, 1509.	1.3	5
8	Salivette, a relevant saliva sampling device for SARS-CoV-2 detection. <i>Journal of Oral Microbiology</i> , 2021, 13, 1920226.	1.2	26
9	HVEM has a broader expression than PD-L1 and constitutes a negative prognostic marker and potential treatment target for melanoma. <i>OncImmunology</i> , 2019, 8, e1665976.	2.1	35
10	LIF Drives Neural Remodeling in Pancreatic Cancer and Offers a New Candidate Biomarker. <i>Cancer Research</i> , 2018, 78, 909-921.	0.4	83
11	Evolutionary and expression analyses reveal a pattern of ancient duplications and functional specializations in the diversification of the Downstream of Kinase (DOK) genes. <i>Developmental and Comparative Immunology</i> , 2018, 84, 193-198.	1.0	2
12	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
13	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
14	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
15	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
16	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
17	$\gamma\delta$ memory-like NK cells contribute to the control of HIV viremia during primary infection: OptprimANRS 147. <i>Clinical and Translational Immunology</i> , 2017, 6, e150.	1.7	42
18	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

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19	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
20	HVEM: A novel cosignaling molecule of major interest in melanoma.. <i>Journal of Clinical Oncology</i> , 2017, 35, e14591-e14591.	0.8	3
21	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
22	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
23	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
24	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
25	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
26	Abstract A61: CAF-derived ANXA6+-exosomes support pancreatic cancer aggressiveness and serve as a circulating biomarker. , 2016, , .		0
27	Increased NK Cell Maturation in Patients with Acute Myeloid Leukemia. <i>Frontiers in Immunology</i> , 2015, 6, 564.	2.2	24
28	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
29	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
30	Follicular Lymphoma B Cells Generate Functional Regulatory T Cells Via ICOS/ICOSL Pathway and Are Inhibited By Intratumoral Tregs. <i>Blood</i> , 2015, 126, 5018-5018.	0.6	0
31	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
32	Cerebrospinal Fluid Biomarker Candidates Associated with Human WNV Neuroinvasive Disease. <i>PLoS ONE</i> , 2014, 9, e93637.	1.1	13
33	Djeen (Database for Joomla!™s Extensible Engine): a research information management system for flexible multi-technology project administration. <i>BMC Research Notes</i> , 2013, 6, 223.	0.6	0
34	Tubulin isoforms identified in the brain by MALDI in-source decay. <i>Journal of Proteomics</i> , 2013, 79, 172-179.	1.2	9
35	Achieving automated scorpion venom mass fingerprinting (VMF) in the nanogram range. <i>Toxicon</i> , 2013, 69, 211-218.	0.8	17
36	Altered Protein Networks and Cellular Pathways in Severe West Nile Disease in Mice. <i>PLoS ONE</i> , 2013, 8, e68318.	1.1	20

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37	Anopheles salivary gland proteomes from major malaria vectors. BMC Genomics, 2012, 13, 614.	1.2	23
38	Multiparametric cytometry for exploration of complex cellular dynamics. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2012, 81A, 332-342.	1.1	9
39	Plasmodium falciparum infection-induced changes in erythrocyte membrane proteins. Parasitology Research, 2012, 110, 545-556.	0.6	21
40	Plasmodium falciparum proteome changes in response to doxycycline treatment. Malaria Journal, 2010, 9, 141.	0.8	62
41	Molecular differences between the divergent responses of ovalbumin-specific CD4 T cells to alum-precipitated ovalbumin compared to ovalbumin expressed by Salmonella. Molecular Immunology, 2008, 45, 3558-3566.	1.0	39
42	Entropy Measures Quantify Global Splicing Disorders in Cancer. PLoS Computational Biology, 2008, 4, e1000011.	1.5	53
43	TranscriptomeBrowser: A Powerful and Flexible Toolbox to Explore Productively the Transcriptional Landscape of the Gene Expression Omnibus Database. PLoS ONE, 2008, 3, e4001.	1.1	27
44	Postoperative serum proteomic profiles may predict metastatic relapse in high-risk primary breast cancer patients receiving adjuvant chemotherapy. Oncogene, 2006, 25, 981-989.	2.6	112
45	MicroArray Facility: a laboratory information management system with extended support for Nylon based technologies. BMC Genomics, 2006, 7, 240.	1.2	4
46	The disparate nature of "intergenic" polyadenylation sites. Rna, 2006, 12, 1794-1801.	1.6	25
47	Gene profiling reveals specific oncogenic mechanisms and signaling pathways in oncocytic and papillary thyroid carcinoma. Oncogene, 2005, 24, 4155-4161.	2.6	40
48	Transcriptional Profiling Reveals Coordinated Up-Regulation of Oxidative Metabolism Genes in Thyroid Oncocytic Tumors. Journal of Clinical Endocrinology and Metabolism, 2004, 89, 994-1005.	1.8	80
49	Feature extraction and signal processing for nylon DNA microarrays. BMC Genomics, 2004, 5, 38.	1.2	37
50	Genomic organization and the tissue distribution of alternatively spliced isoforms of the mouse Spatial gene. BMC Genomics, 2004, 5, 41.	1.2	7
51	Small lymphocytic lymphoma, marginal zone B-cell lymphoma, and mantle cell lymphoma exhibit distinct gene-expression profiles allowing molecular diagnosis. Blood, 2004, 103, 2727-2737.	0.6	127
52	Gene expression profiling of breast carcinomas using Nylon DNA arrays. Comptes Rendus - Biologies, 2003, 326, 1031-1039.	0.1	3
53	The Strategy of T Cell Antigen-presenting Cell Encounter in Antigen-draining Lymph Nodes Revealed by Imaging of Initial T Cell Activation. Journal of Experimental Medicine, 2003, 198, 715-724.	4.2	205
54	Gene expression profiling of multiple myeloma reveals molecular portraits in relation to the pathogenesis of the disease. Blood, 2003, 101, 4998-5006.	0.6	124

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55	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
56	Gene expression profiles of poor-prognosis primary breast cancer correlate with survival. <i>Human Molecular Genetics</i> , 2002, 11, 863-872.	1.4	117
57	Inducible Antibacterial Defense System in <i>C. elegans</i> . <i>Current Biology</i> , 2002, 12, 1209-1214.	1.8	417
58	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
59	Gene expression profiling defines molecular subtypes of classical Hodgkin's disease. <i>Oncogene</i> , 2002, 21, 3095-3102.	2.6	91
60	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
61	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
62	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
63	Gene expression profiling of primary breast carcinomas using arrays of candidate genes. <i>Human Molecular Genetics</i> , 2000, 9, 2981-2991.	1.4	168
64	Transcriptional Expression Analysis of T-Cell Activation by Multiplex Messenger Assay (MMA). , 2000, 134, 337-353.		1
65	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
66	'LABNOTE', a laboratory notebook system designed for academic genomics groups. <i>Nucleic Acids Research</i> , 1999, 27, 601-607.	6.5	11
67	Expression scanning of an array of growth control genes in human tumor cell lines. <i>Oncogene</i> , 1999, 18, 3905-3912.	2.6	57
68	An ESTs description of the new Vanin gene family conserved from fly to human. <i>Immunogenetics</i> , 1999, 49, 964-972.	1.2	45
69	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
70	Expression profiling: DNA arrays in many guises. <i>BioEssays</i> , 1999, 21, 781-790.	1.2	102
71	Atomic resolution on small three-dimensional metal clusters by STM. <i>Surface Science</i> , 1997, 391, 19-26.	0.8	71
72	Differential mRNA Expression in Untreated and TNF- α Elicited Murine Dendritic Cells Precursors. <i>Advances in Experimental Medicine and Biology</i> , 1997, 417, 467-473.	0.8	2

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73	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
74	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
75	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
76	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
77	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
78	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
79	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