Raphael Sandaltzopoulos

List of Publications by Year in descending order

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46 papers

4,236 citations

257450 24 h-index 233421 45 g-index

47 all docs 47 docs citations

47 times ranked

6458 citing authors

#	Article	IF	CITATIONS
1	TEM1-targeting PEGylated PLGA shikonin nanoformulation for immunomodulation and eradication of ovarian cancer BioImpacts, 2022, 12, 65-86.	1.5	1
2	Higher-order connections between stereotyped subsets: implications for improved patient classification in CLL. Blood, 2021, 137, 1365-1376.	1.4	72
3	Higher-order immunoglobulin repertoire restrictions in CLL: the illustrative case of stereotyped subsets 2 and 169. Blood, 2021, 137, 1895-1904.	1.4	21
4	Plant Yield Efficiency by Homeostasis as Selection Tool at Ultra-Low Density. A Comparative Study with Common Stability Measures in Maize. Agronomy, 2020, 10, 1203.	3.0	4
5	Improved plant yield efficiency alleviates the erratic optimum density in maize. Agronomy Journal, 2020, 112, 1690-1701.	1.8	19
6	TRIP - T cell receptor/immunoglobulin profiler. BMC Bioinformatics, 2020, 21, 422.	2.6	11
7	VH CDR3-Focused Somatic Hypermutation in CLL IGHV-IGHD-IGHJ Gene Rearrangements with 100% IGHV Germline Identity. Blood, 2019, 134, 4277-4277.	1.4	3
8	Higher Order Restrictions of the Immunoglobulin Repertoire in CLL: The Illustrative Case of Stereotyped Subsets #2 and #169. Blood, 2019, 134, 5453-5453.	1.4	1
9	The homeodomain transcription factor MEIS1 triggers chemokine expression and is involved in CD8+ Tâ€lymphocyte infiltration in early stage ovarian cancer. Molecular Carcinogenesis, 2018, 57, 1251-1263.	2.7	19
10	Roles of NF-κB Signaling in the Regulation of miRNAs Impacting on Inflammation in Cancer. Biomedicines, 2018, 6, 40.	3.2	75
11	Remarkable Functional Constraints on the Antigen Receptors of CLL Stereotyped Subset #2: High-Throughput Immunogenetic Evidence. Blood, 2018, 132, 1839-1839.	1.4	5
12	Senescence-associated microRNAs target cell cycle regulatory genes in normal human lung fibroblasts. Experimental Gerontology, 2017, 96, 110-122.	2.8	50
13	A step-by-step microRNA guide to cancer development and metastasis. Cellular Oncology (Dordrecht), 2017, 40, 303-339.	4.4	129
14	Editorial: The microRNA 221/222 Cluster: Inaugurating a New Era in Cardiovascular Disease and Cancer?. Current Vascular Pharmacology, 2016, 15, 47-50.	1.7	1
15	Editorial: Novel microRNAs as Putative Therapeutic Targets in Cardiovascular Diseases. Current Vascular Pharmacology, 2015, 13, 564-565.	1.7	6
16	Overexpression of <i>GPC6</i> and <i>TMEM132D</i> in Early Stage Ovarian Cancer Correlates with CD8+ T-Lymphocyte Infiltration and Increased Patient Survival. BioMed Research International, 2015, 2015, 1-9.	1.9	20
17	Beads-on-a-String on a Bead: Reconstitution and Analysis of Chromatin on a Solid Support. Methods in Molecular Biology, 2015, 1288, 1-14.	0.9	0
18	Melt-curve-multiplex-haplotype-specific-PCR, a valuable tool for biological studies: Application in congeneric species discrimination assay. Biochemical Systematics and Ecology, 2014, 56, 271-277.	1.3	2

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19	Overexpression of SMARCE1 is associated with CD8+ T-cell infiltration in early stage ovarian cancer. International Journal of Biochemistry and Cell Biology, 2014, 53, 389-398.	2.8	15
20	Shikonin-loaded antibody-armed nanoparticles for targeted therapy of ovarian cancer. International Journal of Nanomedicine, 2014, 9, 1855.	6.7	48
21	Chimeric Antigen Receptor T Cells with Dissociated Signaling Domains Exhibit Focused Antitumor Activity with Reduced Potential for Toxicity <i>In Vivo</i> . Cancer Immunology Research, 2013, 1, 43-53.	3.4	284
22	Novel Recombinant Human B7-H4 Antibodies Overcome Tumoral Immune Escape to Potentiate T-Cell Antitumor Responses. Cancer Research, 2013, 73, 4820-4829.	0.9	94
23	Redirected Antitumor Activity of Primary Human Lymphocytes Transduced With a Fully Human Anti-mesothelin Chimeric Receptor. Molecular Therapy, 2012, 20, 633-643.	8.2	161
24	Primary Human Ovarian Epithelial Cancer Cells Broadly Express HER2 at Immunologically-Detectable Levels. PLoS ONE, 2012, 7, e49829.	2.5	60
25	Biochemical and molecular analysis of the interaction between ERK2 MAP kinase and hypoxia inducible factor- $1\hat{l}_{\pm}$. International Journal of Biochemistry and Cell Biology, 2011, 43, 1582-1590.	2.8	14
26	Novel surface targets and serum biomarkers from the ovarian cancer vasculature. Cancer Biology and Therapy, 2011, 12, 169-180.	3.4	33
27	Mannose Receptor (MR) Engagement by Mesothelin GPI Anchor Polarizes Tumor-Associated Macrophages and Is Blocked by Anti-MR Human Recombinant Antibody. PLoS ONE, 2011, 6, e28386.	2.5	34
28	TAF4b and Jun/Activating Protein-1 Collaborate to Regulate the Expression of Integrin $\hat{l}\pm 6$ and Cancer Cell Migration Properties. Molecular Cancer Research, 2010, 8, 554-568.	3.4	13
29	Distinct Region-Specific α-Synuclein Oligomers in A53T Transgenic Mice: Implications for Neurodegeneration. Journal of Neuroscience, 2010, 30, 3409-3418.	3.6	89
30	Metal-induced carcinogenesis, oxidative stress and hypoxia signalling. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2009, 674, 31-35.	1.7	107
31	Analysis of Reconstituted Chromatin Using a Solid-Phase Approach. Methods in Molecular Biology, 2009, 523, 11-25.	0.9	1
32	Reliable and rapid discrimination of congeneric species by mtDNA SNP analysis by multiplex PCR: application on three Trachurus and two Mullus fish species as model cases. Hydrobiologia, 2008, 614, 401-404.	2.0	6
33	Reactive oxygen species and HIF-1 signalling in cancer. Cancer Letters, 2008, 266, 12-20.	7.2	186
34	miR-210 links hypoxia with cell cycle regulation and is deleted in human epithelial ovarian cancer. Cancer Biology and Therapy, 2008, 7, 255-264.	3.4	324
35	Genomic and epigenetic alterations deregulate microRNA expression in human epithelial ovarian cancer. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 7004-7009.	7.1	491
36	Tumor Vascular Proteins As Biomarkers in Ovarian Cancer. Journal of Clinical Oncology, 2007, 25, 852-861.	1.6	172

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37	miRNA genetic alterations in human cancers. Expert Opinion on Biological Therapy, 2007, 7, 1375-1386.	3.1	50
38	Rapid identification of Salmo trutta lineages by multiplex PCR utilizing primers tailored to discriminate single nucleotide polymorphisms (SNPs) of the mitochondrial control region. Conservation Genetics, 2007, 8, 1025-1028.	1.5	9
39	Analysis of Activator-Dependent Transcription Reinitiation In Vitro. Methods in Enzymology, 2003, 370, 487-501.	1.0	3
40	Dual Functions of Largest NURF Subunit NURF301 in Nucleosome Sliding and Transcription Factor Interactions. Molecular Cell, 2001, 8, 531-543.	9.7	229
41	ATP-Dependent Histone Octamer Sliding Mediated by the Chromatin Remodeling Complex NURF. Cell, 1999, 97, 833-842.	28.9	302
42	Purification of Drosophila nucleosome remodeling factor. Methods in Enzymology, 1999, 304, 757-765.	1.0	4
43	Heat Shock Factor Increases the Reinitiation Rate from Potentiated Chromatin Templates. Molecular and Cellular Biology, 1998, 18, 361-367.	2.3	56
44	The bifunctional protein DCoH modulates interactions of the homeodomain transcription factor HNF1 with nucleic acids 1 1Edited by M. Yaniv. Journal of Molecular Biology, 1997, 265, 20-29.	4.2	39
45	Self-organization of microtubules into bipolar spindles around artificial chromosomes in Xenopus egg extracts. Nature, 1996, 382, 420-425.	27.8	921
46	Solid phase DNase I footprinting: quick and versatile. Nucleic Acids Research, 1994, 22, 1511-1512.	14.5	52