

Michael Hallett

List of Publications by Year in descending order

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Version: 2024-02-01

97
papers

9,369
citations

53660

45
h-index

39575

94
g-index

102
all docs

102
docs citations

102
times ranked

15399
citing authors

#	ARTICLE	IF	CITATIONS
1	The time-varying effect of radiotherapy after breast-conserving surgery for DCIS. <i>Breast Cancer Research and Treatment</i> , 2019, 178, 221-230.	1.1	3
2	Reproducible Data Analysis Pipelines for Precision Medicine. , 2019, , .		2
3	Spatially distinct tumor immune microenvironments stratify triple-negative breast cancers. <i>Journal of Clinical Investigation</i> , 2019, 129, 1785-1800.	3.9	266
4	The Complex Subtype-Dependent Role of Connexin 43 (GJA1) in Breast Cancer. <i>International Journal of Molecular Sciences</i> , 2018, 19, 693.	1.8	30
5	Discovery of Stromal Regulatory Networks that Suppress Ras-Sensitized Epithelial Cell Proliferation. <i>Developmental Cell</i> , 2017, 41, 392-407.e6.	3.1	25
6	Building Applications for Interactive Data Exploration in Systems Biology. , 2017, , .		1
7	Identification of Interacting Stromal Axes in Triple-Negative Breast Cancer. <i>Cancer Research</i> , 2017, 77, 4673-4683.	0.4	25
8	Interactions between the tumor and the blood systemic response of breast cancer patients. <i>PLoS Computational Biology</i> , 2017, 13, e1005680.	1.5	22
9	Structural and Functional Characterization of a <i>Caenorhabditis elegans</i> Genetic Interaction Network within Pathways. <i>PLoS Computational Biology</i> , 2016, 12, e1004738.	1.5	3
10	Molecular Features of Subtype-Specific Progression from Ductal Carcinoma In Situ to Invasive Breast Cancer. <i>Cell Reports</i> , 2016, 16, 1166-1179.	2.9	85
11	The ErbB2 ^{Ex16} splice variant is a major oncogenic driver in breast cancer that promotes a pro-metastatic tumor microenvironment. <i>Oncogene</i> , 2016, 35, 6053-6064.	2.6	54
12	Chordin-Like 1 Suppresses Bone Morphogenetic Protein 4-Induced Breast Cancer Cell Migration and Invasion. <i>Molecular and Cellular Biology</i> , 2016, 36, 1509-1525.	1.1	53
13	Rab11-FIP1C Is a Critical Negative Regulator in ErbB2-Mediated Mammary Tumor Progression. <i>Cancer Research</i> , 2016, 76, 2662-2674.	0.4	31
14	STAT3 Establishes an Immunosuppressive Microenvironment during the Early Stages of Breast Carcinogenesis to Promote Tumor Growth and Metastasis. <i>Cancer Research</i> , 2016, 76, 1416-1428.	0.4	87
15	SOCIO-ETHICAL ISSUES IN PERSONALIZED MEDICINE: A SYSTEMATIC REVIEW OF ENGLISH LANGUAGE HEALTH TECHNOLOGY ASSESSMENTS OF GENE EXPRESSION PROFILING TESTS FOR BREAST CANCER PROGNOSIS. <i>International Journal of Technology Assessment in Health Care</i> , 2015, 31, 36-50.	0.2	5
16	A 12-gene signature to distinguish colon cancer patients with better clinical outcome following treatment with 5-fluorouracil or FOLFIRI. <i>Journal of Pathology: Clinical Research</i> , 2015, 1, 160-172.	1.3	8
17	Role of DNA Methylation in the Nucleus Accumbens in Incubation of Cocaine Craving. <i>Journal of Neuroscience</i> , 2015, 35, 8042-8058.	1.7	137
18	A functional in vitro model of heterotypic interactions reveals a role for interferon-positive carcinoma associated fibroblasts in breast cancer. <i>BMC Cancer</i> , 2015, 15, 130.	1.1	16

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19	Maternal depression is associated with DNA methylation changes in cord blood T lymphocytes and adult hippocampi. <i>Translational Psychiatry</i> , 2015, 5, e545-e545.	2.4	106
20	Disruption of histone methylation in developing sperm impairs offspring health transgenerationally. <i>Science</i> , 2015, 350, aab2006.	6.0	426
21	DNA Methylation Signature of Childhood Chronic Physical Aggression in T Cells of Both Men and Women. <i>PLoS ONE</i> , 2014, 9, e86822.	1.1	81
22	The Prognostic Ease and Difficulty of Invasive Breast Carcinoma. <i>Cell Reports</i> , 2014, 9, 129-142.	2.9	64
23	Autocrine Activation of the Wnt/ β -Catenin Pathway by CUX1 and GLIS1 in Breast Cancers. <i>Biology Open</i> , 2014, 3, 937-946.	0.6	41
24	Naive CD4 T-cell activation identifies MS patients having rapid transition to progressive MS. <i>Neurology</i> , 2014, 82, 681-690.	1.5	22
25	Dynamic Reprogramming of Signaling Upon Met Inhibition Reveals a Mechanism of Drug Resistance in Gastric Cancer. <i>Science Signaling</i> , 2014, 7, ra38.	1.6	40
26	Genome-Wide Study of Hypomethylated and Induced Genes in Patients with Liver Cancer Unravels Novel Anticancer Targets. <i>Clinical Cancer Research</i> , 2014, 20, 3118-3132.	3.2	85
27	p66ShcA Promotes Breast Cancer Plasticity by Inducing an Epithelial-to-Mesenchymal Transition. <i>Molecular and Cellular Biology</i> , 2014, 34, 3689-3701.	1.1	19
28	Synergistic effects of combined DNA methyltransferase inhibition and MBD2 depletion on breast cancer cells; MBD2 depletion blocks 5-aza-2'-deoxycytidine-triggered invasiveness. <i>Carcinogenesis</i> , 2014, 35, 2436-2446.	1.3	16
29	Association of Childhood Chronic Physical Aggression with a DNA Methylation Signature in Adult Human T Cells. <i>PLoS ONE</i> , 2014, 9, e89839.	1.1	76
30	Transcription onset of genes critical in liver carcinogenesis is epigenetically regulated by methylated DNA-binding protein MBD2. <i>Carcinogenesis</i> , 2013, 34, 2738-2749.	1.3	32
31	Effects of Antenatal Synthetic Glucocorticoid on Glucocorticoid Receptor Binding, DNA Methylation, and Genome-Wide mRNA Levels in the Fetal Male Hippocampus. <i>Endocrinology</i> , 2013, 154, 4170-4181.	1.4	62
32	Glucocorticoid Programming of the Fetal Male Hippocampal Epigenome. <i>Endocrinology</i> , 2013, 154, 1168-1180.	1.4	83
33	β -Catenin Signaling Is a Critical Event in ErbB2-Mediated Mammary Tumor Progression. <i>Cancer Research</i> , 2013, 73, 4474-4487.	0.4	65
34	The ShcA PTB Domain Functions as a Biological Sensor of Phosphotyrosine Signaling during Breast Cancer Progression. <i>Cancer Research</i> , 2013, 73, 4521-4532.	0.4	13
35	Ets2 in Tumor Fibroblasts Promotes Angiogenesis in Breast Cancer. <i>PLoS ONE</i> , 2013, 8, e71533.	1.1	33
36	Met synergizes with p53 loss to induce mammary tumors that possess features of claudin-low breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E1301-E1310.	3.3	61

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37	Epigenomic socioeconomic studies more similar than different. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E1246.	3.3	8
38	Posttranscriptional Regulation of <i>PER1</i> Underlies the Oncogenic Function of IRE1 α . Cancer Research, 2013, 73, 4732-4743.	0.4	115
39	Discovery and Validation of DNA Hypomethylation Biomarkers for Liver Cancer Using HRM-Specific Probes. PLoS ONE, 2013, 8, e68439.	1.1	12
40	Differential DNA Methylation Regions in Cytokine and Transcription Factor Genomic Loci Associate with Childhood Physical Aggression. PLoS ONE, 2013, 8, e71691.	1.1	60
41	The Signature of Maternal Rearing in the Methylome in Rhesus Macaque Prefrontal Cortex and T Cells. Journal of Neuroscience, 2012, 32, 15626-15642.	1.7	340
42	Associations with early-life socio-economic position in adult DNA methylation. International Journal of Epidemiology, 2012, 41, 62-74.	0.9	380
43	ABCC5 supports osteoclast formation and promotes breast cancer metastasis to bone. Breast Cancer Research, 2012, 14, R149.	2.2	40
44	More on Frege and Hilbert. The Western Ontario Series in Philosophy of Science, 2012, , 135-162.	0.2	2
45	Broad Epigenetic Signature of Maternal Care in the Brain of Adult Rats. PLoS ONE, 2011, 6, e14739.	1.1	406
46	Simultaneous Identification of Duplications and Lateral Gene Transfers. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2011, 8, 517-535.	1.9	155
47	Intracellular Eukaryotic Parasites Have a Distinct Unfolded Protein Response. PLoS ONE, 2011, 6, e19118.	1.1	45
48	Claudin-2 is selectively enriched in and promotes the formation of breast cancer liver metastases through engagement of integrin complexes. Oncogene, 2011, 30, 1318-1328.	2.6	130
49	Definition of the Landscape of Promoter DNA Hypomethylation in Liver Cancer. Cancer Research, 2011, 71, 5891-5903.	0.4	187
50	Stromal retinoic acid receptor β promotes mammary gland tumorigenesis. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 774-779.	3.3	35
51	Cut homeobox 1 causes chromosomal instability by promoting bipolar division after cytokinesis failure. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 1949-1954.	3.3	43
52	Reverse Genetics in <i>Candida albicans</i> Predicts ARF Cycling Is Essential for Drug Resistance and Virulence. PLoS Pathogens, 2010, 6, e1000753.	2.1	51
53	Searching for Signaling Balance through the Identification of Genetic Interactors of the Rab Guanine-Nucleotide Dissociation Inhibitor <i>gdi-1</i> . PLoS ONE, 2010, 5, e10624.	1.1	11
54	ADAM10 Releases a Soluble Form of the GPNMB/Osteoactivin Extracellular Domain with Angiogenic Properties. PLoS ONE, 2010, 5, e12093.	1.1	149

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55	Receptor Tyrosine Kinase Signaling Favors a Protumorigenic State in Breast Cancer Cells by Inhibiting the Adaptive Immune Response. <i>Cancer Research</i> , 2010, 70, 7776-7787.	0.4	25
56	Glycoprotein Nonmetastatic B Is an Independent Prognostic Indicator of Recurrence and a Novel Therapeutic Target in Breast Cancer. <i>Clinical Cancer Research</i> , 2010, 16, 2147-2156.	3.2	172
57	Molecular diversity in ductal carcinoma <i>in situ</i> (DCIS) and early invasive breast cancer. <i>Molecular Oncology</i> , 2010, 4, 357-368.	2.1	107
58	Crkl Transgene Induces Atypical Mammary Gland Development and Tumorigenesis. <i>American Journal of Pathology</i> , 2010, 176, 446-460.	1.9	23
59	Frege and Hilbert. , 2010, , 413-464.		9
60	A Biochemical Genomics Screen for Substrates of Ste20p Kinase Enables the In Silico Prediction of Novel Substrates. <i>PLoS ONE</i> , 2009, 4, e8279.	1.1	2
61	Identification of a Stat3-Dependent Transcription Regulatory Network Involved in Metastatic Progression. <i>Cancer Research</i> , 2009, 69, 6823-6830.	0.4	96
62	PTEN Deficiency in a Luminal ErbB-2 Mouse Model Results in Dramatic Acceleration of Mammary Tumorigenesis and Metastasis. <i>Journal of Biological Chemistry</i> , 2009, 284, 19018-19026.	1.6	66
63	Genome-Wide Identification of Direct Target Genes Implicates Estrogen-Related Receptor $\hat{\pm}$ as a Determinant of Breast Cancer Heterogeneity. <i>Cancer Research</i> , 2009, 69, 6149-6157.	0.4	146
64	Chemogenomic profiling predicts antifungal synergies. <i>Molecular Systems Biology</i> , 2009, 5, 338.	3.2	71
65	Pten in stromal fibroblasts suppresses mammary epithelial tumours. <i>Nature</i> , 2009, 461, 1084-1091.	13.7	475
66	Stromal signature identifies basal breast cancers. <i>Nature Medicine</i> , 2009, 15, 238-238.	15.2	1
67	Met induces mammary tumors with diverse histologies and is associated with poor outcome and human basal breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 12903-12908.	3.3	188
68	Stromal gene expression predicts clinical outcome in breast cancer. <i>Nature Medicine</i> , 2008, 14, 518-527.	15.2	1,497
69	DNA demethylation induced by the methyl-CpG-binding domain protein MBD3. <i>Gene</i> , 2008, 420, 99-106.	1.0	41
70	Phosphatase and Tensin Homologue Deleted on Chromosome 10 Deficiency Accelerates Tumor Induction in a Mouse Model of ErbB-2 Mammary Tumorigenesis. <i>Cancer Research</i> , 2008, 68, 2122-2131.	0.4	45
71	Reflections on the Purity of Method in Hilbert's <i>Grundlagen der Geometrie</i> . , 2008, , 198-255.		25
72	Peptides you can count on. <i>Nature Biotechnology</i> , 2007, 25, 61-62.	9.4	20

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73	Osteoactivin Promotes Breast Cancer Metastasis to Bone. <i>Molecular Cancer Research</i> , 2007, 5, 1001-1014.	1.5	146
74	Distinct ErbB-2â€“Coupled Signaling Pathways Promote Mammary Tumors with Unique Pathologic and Transcriptional Profiles. <i>Cancer Research</i> , 2007, 67, 7579-7588.	0.4	23
75	Tools for visually exploring biological networks. <i>Bioinformatics</i> , 2007, 23, 2651-2659.	1.8	220
76	A Faster FPT Algorithm for the Maximum Agreement Forest Problem. <i>Theory of Computing Systems</i> , 2007, 41, 539-550.	0.7	18
77	Gene expression signatures of morphologically normal breast tissue identify basal-like tumors. <i>Breast Cancer Research</i> , 2006, 8, R58.	2.2	122
78	Global Survey of Organ and Organelle Protein Expression in Mouse: Combined Proteomic and Transcriptomic Profiling. <i>Cell</i> , 2006, 125, 173-186.	13.5	429
79	Quantitative Proteomics Analysis of the Secretory Pathway. <i>Cell</i> , 2006, 127, 1265-1281.	13.5	425
80	Strain-dependent pulmonary gene expression profiles of a cystic fibrosis mouse model. <i>Physiological Genomics</i> , 2006, 25, 336-345.	1.0	28
81	Mapping cis-acting regulatory variation in recombinant congenic strains. <i>Physiological Genomics</i> , 2006, 25, 294-302.	1.0	20
82	Dynamical properties of model gene networks and implications for the inverse problem. <i>BioSystems</i> , 2006, 84, 115-123.	0.9	9
83	A Fixed-Parameter Approach to 2-Layer Planarization. <i>Algorithmica</i> , 2006, 45, 159-182.	1.0	30
84	Refining Protein Subcellular Localization. <i>PLoS Computational Biology</i> , 2005, 1, e66.	1.5	90
85	Identifying Regulatory Subnetworks for a Set of Genes. <i>Molecular and Cellular Proteomics</i> , 2005, 4, 683-692.	2.5	61
86	Murine candidate bleomycin induced pulmonary fibrosis susceptibility genes identified by gene expression and sequence analysis of linkage regions. <i>Journal of Medical Genetics</i> , 2005, 42, 464-473.	1.5	40
87	Towards Quality Control for DNA Microarrays. <i>Journal of Computational Biology</i> , 2004, 11, 945-970.	0.8	3
88	Predicting Subcellular Localization via Protein Motif Co-Occurrence. <i>Genome Research</i> , 2004, 14, 1957-1966.	2.4	92
89	ANCESTRAL MAXIMUM LIKELIHOOD OF EVOLUTIONARY TREES IS HARD. <i>Journal of Bioinformatics and Computational Biology</i> , 2004, 02, 257-271.	0.3	17
90	Inferring models of gene expression dynamics. <i>Journal of Theoretical Biology</i> , 2004, 230, 289-299.	0.8	55

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91	Analogues & duals of the MAST problem for sequences & trees. <i>Journal of Algorithms</i> , 2003, 49, 192-216.	0.9	44
92	DePIE: Designing Primers for Protein Interaction Experiments. <i>Nucleic Acids Research</i> , 2003, 31, 3755-3757.	6.5	6
93	Cell fingerprinting: An approach to classifying cells according to mass profiles of digests of protein extracts. <i>Proteomics</i> , 2001, 1, 683-690.	1.3	17
94	Identifying the most significant pairwise correlations of residues in different positions of helices. , 2001, , .		0
95	Analogues and Duals of the MAST Problem for Sequences and Trees. <i>Lecture Notes in Computer Science</i> , 1998, , 103-114.	1.0	12
96	Towards a Theory of Mathematical Research Programmes (II). <i>British Journal for the Philosophy of Science</i> , 1979, 30, 135-159.	1.4	25
97	Towards a Theory of Mathematical Research Programmes (I). <i>British Journal for the Philosophy of Science</i> , 1979, 30, 1-25.	1.4	59