Manoj K Bhasin

List of Publications by Year
in descending order

140 papers	8,162 citations	44069 48 h-index	51608 86 g-index
149 all docs	149 docs citations	149 times ranked	12652 citing authors

#	Article	IF	Citations
1	PGC-1Î± promotes recovery after acute kidney injury during systemic inflammation in mice. Journal of Clinical Investigation, 2011, 121, 4003-4014.	8.2	404
2	PGC1Î± drives NAD biosynthesis linking oxidative metabolism to renal protection. Nature, 2016, 531, 528-532.	27.8	395
3	Prediction of CTL epitopes using QM, SVM and ANN techniques. Vaccine, 2004, 22, 3195-3204.	3.8	296
4	ESLpred: SVM-based method for subcellular localization of eukaryotic proteins using dipeptide composition and PSI-BLAST. Nucleic Acids Research, 2004, 32, W414-W419.	14.5	280
5	Classification of Nuclear Receptors Based on Amino Acid Composition and Dipeptide Composition. Journal of Biological Chemistry, 2004, 279, 23262-23266.	3.4	244
6	Notch1 Contributes to Mouse T-Cell Leukemia by Directly Inducing the Expression of c -myc. Molecular and Cellular Biology, 2006, 26, 8022-8031.	2.3	241
7	Relaxation Response Induces Temporal Transcriptome Changes in Energy Metabolism, Insulin Secretion and Inflammatory Pathways. PLoS ONE, 2013, 8, e62817.	2.5	223
8	A metabolic prosurvival role for PML in breast cancer. Journal of Clinical Investigation, 2012, 122, 3088-3100.	8.2	220
9	PSLpred: prediction of subcellular localization of bacterial proteins. Bioinformatics, 2005, 21, 2522-2524.	4.1	204
10	Support Vector Machine-based Method for Subcellular Localization of Human Proteins Using Amino Acid Compositions, Their Order, and Similarity Search. Journal of Biological Chemistry, 2005, 280, 14427-14432.	3.4	202
11	Genomic Counter-Stress Changes Induced by the Relaxation Response. PLoS ONE, 2008, 3, e2576.	2.5	198
12	MHCBN: a comprehensive database of MHC binding and non-binding peptides. Bioinformatics, 2003, 19, 665-666.	4.1	191
13	Identification of the Transcription Factor Single-Minded Homologue 2 as a Potential Biomarker and Immunotherapy Target in Prostate Cancer. Clinical Cancer Research, 2009, 15, 5794-5802.	7.0	184
14	Bcipep: A database of B-cell epitopes. BMC Genomics, 2005, 6, 79.	2.8	179
15	Analysis and prediction of affinity of TAP binding peptides using cascade SVM. Protein Science, 2004, 13, 596-607.	7.6	167
16	MSC-Regulated MicroRNAs Converge on the Transcription Factor FOXP2 and Promote Breast Cancer Metastasis. Cell Stem Cell, 2014, 15, 762-774.	11.1	155
17	GPCRpred: an SVM-based method for prediction of families and subfamilies of G-protein coupled receptors. Nucleic Acids Research, 2004, 32, W383-W389.	14.5	154
18	Curative and Î² cell regenerative effects of Î±1-antitrypsin treatment in autoimmune diabetic NOD mice. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 16242-16247.	7.1	154

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19	SVM based method for predicting HLA-DRB1*0401 binding peptides in an antigen sequence. Bioinformatics, 2004, 20, 421-423.	4.1	153
20	A hybrid approach for predicting promiscuous MHC class I restricted T cell epitopes. Journal of Biosciences, 2007, 32, 31-42.	1.1	117
21	The receptor tyrosine kinase Axl is an essential regulator of prostate cancer proliferation and tumor growth and represents a new therapeutic target. Oncogene, 2013, 32, 689-698.	5.9	112
22	Single cell transcriptomic landscape of diabetic foot ulcers. Nature Communications, 2022, 13, 181.	12.8	111
23	Prediction of methylated CpGs in DNA sequences using a support vector machine. FEBS Letters, 2005, 579, 4302-4308.	2.8	103
24	Pcleavage: an SVM based method for prediction of constitutive proteasome and immunoproteasome cleavage sites in antigenic sequences. Nucleic Acids Research, 2005, 33, W202-W207.	14.5	95
25	Preoperative stimulation of resolution and inflammation blockade eradicates micrometastases. Journal of Clinical Investigation, 2019, 129, 2964-2979.	8.2	94
26	Evidence for a role of the histone deacetylase SIRT6 in DNA damage response of multiple myeloma cells. Blood, 2016, 127, 1138-1150.	1.4	89
27	The KDM3Aâ€“KLF2â€“IRF4 axis maintains myeloma cell survival. Nature Communications, 2016, 7, 10258.	12.8	87
28	Reduced PDEF Expression Increases Invasion and Expression of Mesenchymal Genes in Prostate Cancer Cells. Cancer Research, 2007, 67, 4219-4226.	0.9	86
29	Recognition and Classification of Histones Using Support Vector Machine. Journal of Computational Biology, 2006, 13, 102-112.	1.6	79
30	Antiinflammatory Effects of the ETS Factor ERG in Endothelial Cells Are Mediated Through Transcriptional Repression of the Interleukin-8 Gene. Circulation Research, 2009, 104, 1049-1057.	4.5	77
31	MHCBN 4.0: A database of MHC/TAP binding peptides and T-cell epitopes. BMC Research Notes, 2009, 2, 61.	1.4	77
32	Optimizing a Proteomics Platform for Urine Biomarker Discovery. Molecular and Cellular Proteomics, 2010, 9, 2195-2204.	3.8	74
33	CARM1 is required for proper control of proliferation and differentiation of pulmonary epithelial cells. Development (Cambridge), 2010, 137, 2147-2156.	2.5	73
34	RhoJ is an endothelial cell-restricted Rho GTPase that mediates vascular morphogenesis and is regulated by the transcription factor ERG. Blood, 2011, 118, 1145-1153.	1.4	70
35	Upregulation of inflammatory gene transcripts in periosteum of chronic migraineurs: Implications for extracranial origin of headache. Annals of Neurology, 2016, 79, 1000-1013.	5.3	68
36	Integrated Skin Transcriptomics and Serum Multiplex Assays Reveal Novel Mechanisms of Wound Healing in Diabetic Foot Ulcers. Diabetes, 2020, 69, 2157-2169.	0.6	68

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37	Anti-S1P Antibody as a Novel Therapeutic Strategy for VEGFR TKI-Resistant Renal Cancer. Clinical Cancer Research, 2015, 21, 1925-1934.	7.0	67
38	BhairPred: prediction of Â-hairpins in a protein from multiple alignment information using ANN and SVM techniques. Nucleic Acids Research, 2005, 33, W154-W159.	14.5	65
39	Resistance of Renal Cell Carcinoma to Sorafenib Is Mediated by Potentially Reversible Gene Expression. PLoS ONE, 2011, 6, e19144.	2.5	64
40	Genomic and Clinical Effects Associated with a Relaxation Response Mind-Body Intervention in Patients with Irritable Bowel Syndrome and Inflammatory Bowel Disease. PLoS ONE, 2015, 10, e0123861.	2.5	62
41	GPCRsclass: a web tool for the classification of amine type of G-protein-coupled receptors. Nucleic Acids Research, 2005, 33, W143-W147.	14.5	59
42	Topical Application of a Mast Cell Stabilizer Improves Impaired Diabetic Wound Healing. Journal of Investigative Dermatology, 2020, 140, 901-911.e11.	0.7	58
43	Computational Repositioning and Preclinical Validation of Pentamidine for Renal Cell Cancer. Molecular Cancer Therapeutics, 2014, 13, 1929-1941.	4.1	57
44	Alpha 1-antitrypsin reduces inflammation and enhances mouse pancreatic islet transplant survival. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 15443-15448.	7.1	56
45	Multicenter Analysis of Immune Biomarkers and Heart Transplant Outcomes: Results of the Clinical Trials in Organ Transplantation-05 Study. American Journal of Transplantation, 2016, 16, 121-136.	4.7	56
46	Neoadjuvant-Intensive Androgen Deprivation Therapy Selects for Prostate Tumor Foci with Diverse Subclonal Oncogenic Alterations. Cancer Research, 2018, 78, 4716-4730.	0.9	56
47	Bench-to-bedside review: Future novel diagnostics for sepsis - a systems biology approach. Critical Care, 2013, 17, 231.	5.8	55
48	Application of Machine Learning Techniques in Predicting MHC Binders. Methods in Molecular Biology, 2007, 409, 201-215.	0.9	55
49	ERG is required for the differentiation of embryonic stem cells along the endothelial lineage. BMC Developmental Biology, 2009, 9, 72.	2.1	54
50	Bioinformatic identification and characterization of human endothelial cell-restricted genes. BMC Genomics, 2010, 11, 342.	2.8	54
51	Combinatorial Effect of Non-Steroidal Anti-inflammatory Drugs and NF-ÎºB Inhibitors in Ovarian Cancer Therapy. PLoS ONE, 2011, 6, e24285.	2.5	50
52	Detecting Microbial Dysbiosis Associated with Pediatric Crohn Disease Despite the High Variability of the Gut Microbiota. Cell Reports, 2016, 14, 945-955.	6.4	49
53	Meta-analysis of transcriptome data identifies a novel 5-gene pancreatic adenocarcinoma classifier. Oncotarget, 2016, 7, 23263-23281.	1.8	49
54	Identification of key regulators of pancreatic cancer progression through multidimensional systems-level analysis. Genome Medicine, 2016, 8, 38.	8.2	48

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55	Gene expression profile of mouse prostate tumors reveals dysregulations in major biological processes and identifies potential murine targets for preclinical development of human prostate cancer therapy. Prostate, 2008, 68, 1517-1530.	2.3	47
56	Pentraxin-3 is a PI3K signaling target that promotes stem cellâ€“like traits in basal-like breast cancers. Science Signaling, 2017, 10, .	3.6	43
57	Pericytes Elicit Resistance to Vemurafenib and Sorafenib Therapy in Thyroid Carcinoma via the TSP-1/TGFÎ²1 Axis. Clinical Cancer Research, 2018, 24, 6078-6097.	7.0	43
58	Requirement of the Epithelium-specific Ets Transcription Factor Spdef for Mucous Gland Cell Function in the Gastric Antrum. Journal of Biological Chemistry, 2010, 285, 35047-35055.	3.4	42
59	Downregulation of Dipeptidyl Peptidase 4 Accelerates Progression to Castration-Resistant Prostate Cancer. Cancer Research, 2018, 78, 6354-6362.	0.9	42
60	TFEB-driven lysosomal biogenesis is pivotal for PGC1Î±-dependent renal stress resistance. JCI Insight, 2019, 4, .	5.0	40
61	Vemurafenib-resistance via de novo RBM genes mutations and chromosome 5 aberrations is overcome by combined therapy with palbociclib in thyroid carcinoma with BRAFV600E. Oncotarget, 2017, 8, 84743-84760.	1.8	40
62	TAPPred Prediction of TAP-Binding Peptides in Antigens. Methods in Molecular Biology, 2007, 409, 381-386.	0.9	39
63	Prediction of Promiscuous and High-Affinity Mutated MHC Binders. Hybridoma, 2003, 22, 229-234.	0.4	38
64	Novel non-invasive biomarkers that distinguish between benign prostate hyperplasia and prostate cancer. BMC Cancer, 2015, 15, 259.	2.6	37
65	Determination of system level alterations in host transcriptome due to Zika virus (ZIKV) Infection in retinal pigment epithelium. Scientific Reports, 2018, 8, 11209.	3.3	37
66	The Role of TNF-Î± in Mice with Type 1- and 2- Diabetes. PLoS ONE, 2012, 7, e33254.	2.5	35
67	Multicenter evaluation of a standardized protocol for noninvasive gene expression profiling. American Journal of Transplantation, 2013, 13, 1891-1897.	4.7	35
68	Chemotherapyâ€generated cell debris stimulates colon carcinoma tumor growthviaosteopontin. FASEB Journal, 2019, 33, 114-125.	0.5	35
69	Mitochondrial Dysfunction in Atrial Tissue of Patients Developing Postoperative Atrial Fibrillation. Annals of Thoracic Surgery, 2017, 104, 1547-1555.	1.3	33
70	A20 Modulates Lipid Metabolism and Energy Production to Promote Liver Regeneration. PLoS ONE, 2011, 6, e17715.	2.5	33
71	Modulation of lymphocyte-mediated tissue repair by rational design of heterocyclic aryl hydrocarbon receptor agonists. Science Advances, 2020, 6, eaay8230.	10.3	31
72	Temporal retinal transcriptome and systems biology analysis identifies key pathways and hub genes in Staphylococcus aureus endophthalmitis. Scientific Reports, 2016, 6, 21502.	3.3	30

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73	A Phase II Trial of Abiraterone Combined with Dutasteride for Men with Metastatic Castration-Resistant Prostate Cancer. Clinical Cancer Research, 2017, 23, 935-945.	7.0	30
74	Smad2/3â€pathway ligand trap luspatercept enhances erythroid differentiation in murine Î²â€thalassaemia by increasing GATAâ€1 availability. Journal of Cellular and Molecular Medicine, 2020, 24, 6162-6177.	3.6	30
75	Pressure-overload hypertrophy of the developing heart reveals activation of divergent gene and protein pathways in the left and right ventricular myocardium. American Journal of Physiology - Heart and Circulatory Physiology, 2013, 304, H697-H708.	3.2	29
76	ACE2 abrogates tumor resistance to VEGFR inhibitors suggesting angiotensin-(1-7) as a therapy for clear cell renal cell carcinoma. Science Translational Medicine, 2021, 13, .	12.4	29
77	p16Ink4a or p19Arf loss contributes to Tal1-induced leukemogenesis in mice. Oncogene, 2006, 25, 3023-3031.	5.9	25
78	Quantitative proteomic analysis in HCV-induced HCC reveals sets of proteins with potential significance for racial disparity. Journal of Translational Medicine, 2013, 11, 239.	4.4	25
79	High dose intermittent sorafenib shows improved efficacy over conventional continuous dose in renal cell carcinoma. Journal of Translational Medicine, 2011, 9, 220.	4.4	24
80	HIV-1 Infection Impairs Regulatory T-Cell Suppressive Capacity on a Per-Cell Basis. Journal of Infectious Diseases, 2014, 210, 899-903.	4.0	24
81	Functional and Transcriptomic Characterization of Peritoneal Immune-Modulation by Addition of Alanyl-Glutamine to Dialysis Fluid. Scientific Reports, 2017, 7, 6229.	3.3	24
82	Temporal Network Based Analysis of Cell Specific Vein Graft Transcriptome Defines Key Pathways and Hub Genes in Implantation Injury. PLoS ONE, 2012, 7, e39123.	2.5	23
83	SARS-CoV-2 morbidity and mortality in racial/ethnic minority populations: A window into the stress related inflammatory basis of health disparities?. Brain, Behavior, & Immunity - Health, 2020, 9, 100158.	2.5	22
84	Gene expression analysis of embryonic stem cells expressing VE-cadherin (CD144) during endothelial differentiation. BMC Genomics, 2008, 9, 240.	2.8	21
85	Preliminary Biomarkers for Identification of Human Ascending Thoracic Aortic Aneurysm. Journal of the American Heart Association, 2013, 2, e000138.	3.7	21
86	Inhibition of ALK1 signaling with dalantercept combined with VEGFR TKI leads to tumor stasis in renal cell carcinoma. Oncotarget, 0, 7, 41857-41869.	1.8	21
87	A20 Regulates Atherogenic Interferon (IFN)-Î³ Signaling in Vascular Cells by Modulating Basal IFNÎ² Levels. Journal of Biological Chemistry, 2014, 289, 30912-30924.	3.4	20
88	Induced Human Decidual NK-Like Cells Improve Utero-Placental Perfusion in Mice. PLoS ONE, 2016, 11, e0164353.	2.5	20
89	Transcriptomic and proteomic analysis of global ischemia and cardioprotection in the rabbit heart. Physiological Genomics, 2009, 38, 125-137.	2.3	19
90	Prevention of Nonimmunologic Loss of Transplanted Islets in Monkeys. American Journal of Transplantation, 2014, 14, 1543-1551.	4.7	19

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91	Leukemia vaccine overcomes limitations of checkpoint blockade by evoking clonal T cell responses in a murine acute myeloid leukemia model. Haematologica, 2021, 106, 1330-1342.	3.5	19
92	Binding of the WASP/N-WASP-Interacting Protein WIP to Actin Regulates Focal Adhesion Assembly and Adhesion. Molecular and Cellular Biology, 2014, 34, 2600-2610.	2.3	18
93	Microarray and proteomic analysis of the cardioprotective effects of cold blood cardioplegia in the mature and aged male and female. Physiological Genomics, 2012, 44, 1027-1041.	2.3	17
94	Significant lethality following liver resection in A20 heterozygous knockout mice uncovers a key role for A20 in liver regeneration. Cell Death and Differentiation, 2015, 22, 2068-2077.	11.2	17
95	The effects of transfection reagent polyethyleneimine (PEI) and non-targeting control siRNAs on global gene expression in human aortic smooth muscle cells. BMC Genomics, 2016, 17, 20.	2.8	17
96	Transcriptional Patterns in Peritoneal Tissue of Encapsulating Peritoneal Sclerosis, a Complication of Chronic Peritoneal Dialysis. PLoS ONE, 2013, 8, e56389.	2.5	17
97	Specific Transcriptome Changes Associated with Blood Pressure Reduction in Hypertensive Patients After Relaxation Response Training. Journal of Alternative and Complementary Medicine, 2018, 24, 486-504.	2.1	16
98	A Transcriptomics-Based Meta-Analysis Combined With Machine Learning Identifies a Secretory Biomarker Panel for Diagnosis of Pancreatic Adenocarcinoma. Frontiers in Genetics, 2020, 11, 572284.	2.3	15
99	Functional genomics in the study of mind-body therapies. Ochsner Journal, 2014, 14, 681-95.	1.1	15
100	Autocrine Canonical Wnt Signaling Primes Noncanonical Signaling through ROR1 in Metastatic Castration-Resistant Prostate Cancer. Cancer Research, 2022, 82, 1518-1533.	0.9	15
101	Analysis of Host Gene Expression Changes Reveals Distinct Roles for the Cytoplasmic Domain of the Epstein-Barr Virus Receptor/CD21 in B-Cell Maturation, Activation, and Initiation of Virus Infection. Journal of Virology, 2014, 88, 5559-5577.	3.4	14
102	Study of Cathepsin B inhibition in VEGFR TKI treated human renal cell carcinoma xenografts. Oncogenesis, 2019, 8, 15.	4.9	14
103	Analysis of Multiple Sarcoma Expression Datasets: Implications for Classification, Oncogenic Pathway Activation and Chemotherapy Resistance. PLoS ONE, 2010, 5, e9747.	2.5	14
104	Alterations of the gene expression profile in renal cell carcinoma after treatment with the histone deacetylase-inhibitor valproic acid and interferon-alpha. World Journal of Urology, 2011, 29, 779-786.	2.2	13
105	SB225002 Induces Cell Death and Cell Cycle Arrest in Acute Lymphoblastic Leukemia Cells through the Activation of GLIPR1. PLoS ONE, 2015, 10, e0134783.	2.5	13
106	Computational Methods in Genome Research. Applied Mycology and Biotechnology, 2006, , 179-207.	0.3	10
107	Cardiopulmonary Bypass Decreases Activation of the Signal Transducer and Activator of Transcription 3 (STAT3) Pathway in Diabetic Human Myocardium. Annals of Thoracic Surgery, 2015, 100, 1636-1645.	1.3	9
108	Rap-536 (Murine ACE-536/Luspatercept) Inhibits Smad2/3 Signaling and Promotes Erythroid Differentiation By Restoring GATA-1 Function in Murine b-Thalassemia. Blood, 2015, 126, 751-751.	1.4	8

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109	Systems Biology Approach to Identify Novel Genomic Determinants for Pancreatic Cancer Pathogenesis. Scientific Reports, 2019, 9, 123.	3.3	7
110	A <i>Grammastola spatulata</i> mechanotoxin-4 (GsMTx4)-sensitive cation channel mediates increased cation permeability in human hereditary spherocytosis of multiple genetic etiologies. Haematologica, 2021, 106, 2759-2762.	3.5	5
111	Single Cell Transcriptomics Revealed AML and Non-AML Cell Clusters Relevant to Relapse and Remission in Pediatric AML. Blood, 2020, 136, 24-25.	1.4	5
112	Cellular and immunometabolic mechanisms of inflammation in depression: Preliminary findings from single cell RNA sequencing and a tribute to Bruce McEwen. Neurobiology of Stress, 2022, 19, 100462.	4.0	4
113	Leukocyte Transcriptional Response in Sepsis. Shock, 2019, 52, 166-173.	2.1	3
114	Embryonic periventricular endothelial cells demonstrate a unique pro-neurodevelopment and anti-inflammatory gene signature. Scientific Reports, 2020, 10, 20393.	3.3	3
115	Single Cell RNA-Seq Analyses of Healthy Lower Extremity Skin and Diabetic Foot Ulcers Uncover Distinct Immune Landscape of Diabetic Wound Healing. Diabetes, 2018, 67, 647-P.	0.6	3
116	Pediatric Single Cell Cancer Atlas: An Integrative Web-Based Resource for Single Cell Transcriptome Data from Pediatric Leukemias. Blood, 2021, 138, 3488-3488.	1.4	3
117	Comprehensive Genomic Profiling of High-Risk Pediatric Cancer Patients Has a Measurable Impact on Clinical Care. JCO Precision Oncology, 2022, 6, e2100451.	3.0	3
118	Searching and Mapping of T-Cell Epitopes, MHC Binders, and TAP Binders. Methods in Molecular Biology, 2007, 409, 95-112.	0.9	2
119	Integrated Cytof, Scrna-Seq and Cite-Seq Analysis of Bone Marrow Immune Microenvironment in the Mmrf Commpass Study. Blood, 2020, 136, 28-29.	1.4	2
120	Functional and Clinical Impact of Splicing Factor Dysregulation in Multiple Myeloma. Blood, 2015, 126, 726-726.	1.4	1
121	Characterization of T-Cell Exhaustion in Rapid Progressing Multiple Myeloma Using Cross Center Scrna-Seq Study. Blood, 2021, 138, 401-401.	1.4	1
122	Analysis of Single Cell Transcriptomics in Paired Pediatric T-ALL Samples Collected at Diagnosis and Following End of Induction Therapy Reveals an MRD-Associated Stem Cell Signature. Blood, 2021, 138, 1311-1311.	1.4	1
123	Survival Genie: A Web Portal for Single-Cell Data, Gene-Ratio, and Cell Composition-Based Survival Analyses. Blood, 2021, 138, 276-276.	1.4	1
124	Differential gene expression in vein graft endothelial and smooth muscle cells isolated by laser capture microdissection. Journal of the American College of Surgeons, 2009, 209, S137.	0.5	0
125	Identification of Inflammatory Bowel Disease (IBD)-Specific Diagnostic Biomarkers Using Transcriptional Profiling of Peripheral Blood Mononuclear Cells (PBMC) From Treatment NaÃ¯VE Children With Either IBD or Non-IBD Gastrointestinal Inflammatory Disorders. Gastroenterology, 2011, 140, S-509.	1.3	0
126	Genomic Determinants of a Relaxation Response Resiliency Program in Inflammatory Bowel Disease and Irritable Bowel Syndrome. Gastroenterology, 2011, 140, S-524.	1.3	0

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127	Sa1162 Transcriptional Profiling of Peripheral Blood Mononuclear Cells Identifies Genes and Gene Pathways Associated With Pediatric IBD and That Differentiate Crohn's Disease and Ulcerative Colitis. Gastroenterology, 2014, 146, S-216.	1.3	0
128	A Distinct Treg Transcriptome Signature in HIV-1 Elite Controllers Might Contribute to Improved Disease Outcome. AIDS Research and Human Retroviruses, 2014, 30, A180-A180.	1.1	0
129	Next-generation sequencing demonstrates genomic signature of resistance patterns following phosphatidylinositol 3-kinase (PI3K) inhibition. Gynecologic Oncology, 2015, 136, 406.	1.4	0
130	[31-OR]. Pregnancy Hypertension, 2015, 5, 15.	1.4	0
131	Notch1 Provides a Proliferative Signal in Mouse T Cell Leukemia by Directly Targeting c-Myc.. Blood, 2006, 108, 1434-1434.	1.4	0
132	Selective isolation of vein graft endothelial and smooth muscle cells using laser capture microdissection for microarray gene analysis. FASEB Journal, 2009, 23, LB322.	0.5	0
133	CARM1 is required for proper control of proliferation and differentiation of pulmonary epithelial cells. Journal of Cell Science, 2010, 123, e1-e1.	2.0	0
134	Abstract 3188: Discovery and validation of non-invasive biomarkers for benign prostate hyperplasia and prostate cancer. , 2011, , .		0
135	Abstract 1150: Biomarkers discovery and racial disparity in hepatitis C-associated hepatocellular carcinoma.. , 2013, , .		0
136	Abstract LB-173: Early detection of pancreatic cancer using a new 5-gene classifier. , 2014, , .		0
137	The KDM3A-KLF2-IRF4 Axis Maintains Myeloma Cell Survival. Blood, 2015, 126, 3633-3633.	1.4	0
138	Single-Cell RNA-Seq Analysis of CD138-Depleted Bone Marrow Samples Reveals Genetic Alterations and Disease Progression Correlate with Tumor and Bone Marrow Immune Microenvironment in the Mmrf Commpass Study. Blood, 2021, 138, 2691-2691.	1.4	0
139	Mitochondrial Electron Transport Chain Inhibition Promotes Resistance to Proteasome Inhibitors in Multiple Myeloma. Blood, 2021, 138, 1611-1611.	1.4	0
140	Single Cell RNA Sequencing Driven Characterization of Rare B/Myeloid and T/Myeloid Mixed Phenotype Acute Leukemia. Blood, 2021, 138, 3455-3455.	1.4	0