

Manoj K Bhasin

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/82480/publications.pdf>

Version: 2024-02-01

140
papers

8,162
citations

43973

48
h-index

51492

86
g-index

149
all docs

149
docs citations

149
times ranked

12652
citing authors

#	ARTICLE	IF	CITATIONS
1	PGC-1 β promotes recovery after acute kidney injury during systemic inflammation in mice. <i>Journal of Clinical Investigation</i> , 2011, 121, 4003-4014.	3.9	404
2	PGC1 β drives NAD biosynthesis linking oxidative metabolism to renal protection. <i>Nature</i> , 2016, 531, 528-532.	13.7	395
3	Prediction of CTL epitopes using QM, SVM and ANN techniques. <i>Vaccine</i> , 2004, 22, 3195-3204.	1.7	296
4	ESLPred: SVM-based method for subcellular localization of eukaryotic proteins using dipeptide composition and PSI-BLAST. <i>Nucleic Acids Research</i> , 2004, 32, W414-W419.	6.5	280
5	Classification of Nuclear Receptors Based on Amino Acid Composition and Dipeptide Composition. <i>Journal of Biological Chemistry</i> , 2004, 279, 23262-23266.	1.6	244
6	Notch1 Contributes to Mouse T-Cell Leukemia by Directly Inducing the Expression of c-myc. <i>Molecular and Cellular Biology</i> , 2006, 26, 8022-8031.	1.1	241
7	Relaxation Response Induces Temporal Transcriptome Changes in Energy Metabolism, Insulin Secretion and Inflammatory Pathways. <i>PLoS ONE</i> , 2013, 8, e62817.	1.1	223
8	A metabolic prosurvival role for PML in breast cancer. <i>Journal of Clinical Investigation</i> , 2012, 122, 3088-3100.	3.9	220
9	PSLPred: prediction of subcellular localization of bacterial proteins. <i>Bioinformatics</i> , 2005, 21, 2522-2524.	1.8	204
10	Support Vector Machine-based Method for Subcellular Localization of Human Proteins Using Amino Acid Compositions, Their Order, and Similarity Search. <i>Journal of Biological Chemistry</i> , 2005, 280, 14427-14432.	1.6	202
11	Genomic Counter-Stress Changes Induced by the Relaxation Response. <i>PLoS ONE</i> , 2008, 3, e2576.	1.1	198
12	MHCBN: a comprehensive database of MHC binding and non-binding peptides. <i>Bioinformatics</i> , 2003, 19, 665-666.	1.8	191
13	Identification of the Transcription Factor Single-Minded Homologue 2 as a Potential Biomarker and Immunotherapy Target in Prostate Cancer. <i>Clinical Cancer Research</i> , 2009, 15, 5794-5802.	3.2	184
14	Bcipep: A database of B-cell epitopes. <i>BMC Genomics</i> , 2005, 6, 79.	1.2	179
15	Analysis and prediction of affinity of TAP binding peptides using cascade SVM. <i>Protein Science</i> , 2004, 13, 596-607.	3.1	167
16	MSC-Regulated MicroRNAs Converge on the Transcription Factor FOXP2 and Promote Breast Cancer Metastasis. <i>Cell Stem Cell</i> , 2014, 15, 762-774.	5.2	155
17	GPCRpred: an SVM-based method for prediction of families and subfamilies of G-protein coupled receptors. <i>Nucleic Acids Research</i> , 2004, 32, W383-W389.	6.5	154
18	Curative and β cell regenerative effects of β 1-antitrypsin treatment in autoimmune diabetic NOD mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 16242-16247.	3.3	154

#	ARTICLE	IF	CITATIONS
19	SVM based method for predicting HLA-DRB1*0401 binding peptides in an antigen sequence. <i>Bioinformatics</i> , 2004, 20, 421-423.	1.8	153
20	A hybrid approach for predicting promiscuous MHC class I restricted T cell epitopes. <i>Journal of Biosciences</i> , 2007, 32, 31-42.	0.5	117
21	The receptor tyrosine kinase Axl is an essential regulator of prostate cancer proliferation and tumor growth and represents a new therapeutic target. <i>Oncogene</i> , 2013, 32, 689-698.	2.6	112
22	Single cell transcriptomic landscape of diabetic foot ulcers. <i>Nature Communications</i> , 2022, 13, 181.	5.8	111
23	Prediction of methylated CpGs in DNA sequences using a support vector machine. <i>FEBS Letters</i> , 2005, 579, 4302-4308.	1.3	103
24	Pcleavage: an SVM based method for prediction of constitutive proteasome and immunoproteasome cleavage sites in antigenic sequences. <i>Nucleic Acids Research</i> , 2005, 33, W202-W207.	6.5	95
25	Preoperative stimulation of resolution and inflammation blockade eradicates micrometastases. <i>Journal of Clinical Investigation</i> , 2019, 129, 2964-2979.	3.9	94
26	Evidence for a role of the histone deacetylase SIRT6 in DNA damage response of multiple myeloma cells. <i>Blood</i> , 2016, 127, 1138-1150.	0.6	89
27	The KDM3A-KLF2-IRF4 axis maintains myeloma cell survival. <i>Nature Communications</i> , 2016, 7, 10258.	5.8	87
28	Reduced PDEF Expression Increases Invasion and Expression of Mesenchymal Genes in Prostate Cancer Cells. <i>Cancer Research</i> , 2007, 67, 4219-4226.	0.4	86
29	Recognition and Classification of Histones Using Support Vector Machine. <i>Journal of Computational Biology</i> , 2006, 13, 102-112.	0.8	79
30	Antiinflammatory Effects of the ETS Factor ERG in Endothelial Cells Are Mediated Through Transcriptional Repression of the Interleukin-8 Gene. <i>Circulation Research</i> , 2009, 104, 1049-1057.	2.0	77
31	MHCBN 4.0: A database of MHC/TAP binding peptides and T-cell epitopes. <i>BMC Research Notes</i> , 2009, 2, 61.	0.6	77
32	Optimizing a Proteomics Platform for Urine Biomarker Discovery. <i>Molecular and Cellular Proteomics</i> , 2010, 9, 2195-2204.	2.5	74
33	CARM1 is required for proper control of proliferation and differentiation of pulmonary epithelial cells. <i>Development (Cambridge)</i> , 2010, 137, 2147-2156.	1.2	73
34	Rhoj is an endothelial cell-restricted Rho GTPase that mediates vascular morphogenesis and is regulated by the transcription factor ERG. <i>Blood</i> , 2011, 118, 1145-1153.	0.6	70
35	Upregulation of inflammatory gene transcripts in periosteum of chronic migraineurs: Implications for extracranial origin of headache. <i>Annals of Neurology</i> , 2016, 79, 1000-1013.	2.8	68
36	Integrated Skin Transcriptomics and Serum Multiplex Assays Reveal Novel Mechanisms of Wound Healing in Diabetic Foot Ulcers. <i>Diabetes</i> , 2020, 69, 2157-2169.	0.3	68

#	ARTICLE	IF	CITATIONS
37	Anti-S1P Antibody as a Novel Therapeutic Strategy for VEGFR TKI-Resistant Renal Cancer. <i>Clinical Cancer Research</i> , 2015, 21, 1925-1934.	3.2	67
38	BhairPred: prediction of \hat{A} -hairpins in a protein from multiple alignment information using ANN and SVM techniques. <i>Nucleic Acids Research</i> , 2005, 33, W154-W159.	6.5	65
39	Resistance of Renal Cell Carcinoma to Sorafenib Is Mediated by Potentially Reversible Gene Expression. <i>PLoS ONE</i> , 2011, 6, e19144.	1.1	64
40	Genomic and Clinical Effects Associated with a Relaxation Response Mind-Body Intervention in Patients with Irritable Bowel Syndrome and Inflammatory Bowel Disease. <i>PLoS ONE</i> , 2015, 10, e0123861.	1.1	62
41	GPCRclass: a web tool for the classification of amine type of G-protein-coupled receptors. <i>Nucleic Acids Research</i> , 2005, 33, W143-W147.	6.5	59
42	Topical Application of a Mast Cell Stabilizer Improves Impaired Diabetic Wound Healing. <i>Journal of Investigative Dermatology</i> , 2020, 140, 901-911.e11.	0.3	58
43	Computational Repositioning and Preclinical Validation of Pentamidine for Renal Cell Cancer. <i>Molecular Cancer Therapeutics</i> , 2014, 13, 1929-1941.	1.9	57
44	Alpha 1-antitrypsin reduces inflammation and enhances mouse pancreatic islet transplant survival. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 15443-15448.	3.3	56
45	Multicenter Analysis of Immune Biomarkers and Heart Transplant Outcomes: Results of the Clinical Trials in Organ Transplantation-05 Study. <i>American Journal of Transplantation</i> , 2016, 16, 121-136.	2.6	56
46	Neoadjuvant-Intensive Androgen Deprivation Therapy Selects for Prostate Tumor Foci with Diverse Subclonal Oncogenic Alterations. <i>Cancer Research</i> , 2018, 78, 4716-4730.	0.4	56
47	Bench-to-bedside review: Future novel diagnostics for sepsis - a systems biology approach. <i>Critical Care</i> , 2013, 17, 231.	2.5	55
48	Application of Machine Learning Techniques in Predicting MHC Binders. <i>Methods in Molecular Biology</i> , 2007, 409, 201-215.	0.4	55
49	ERG is required for the differentiation of embryonic stem cells along the endothelial lineage. <i>BMC Developmental Biology</i> , 2009, 9, 72.	2.1	54
50	Bioinformatic identification and characterization of human endothelial cell-restricted genes. <i>BMC Genomics</i> , 2010, 11, 342.	1.2	54
51	Combinatorial Effect of Non-Steroidal Anti-inflammatory Drugs and NF- $\hat{\kappa}$ B Inhibitors in Ovarian Cancer Therapy. <i>PLoS ONE</i> , 2011, 6, e24285.	1.1	50
52	Detecting Microbial Dysbiosis Associated with Pediatric Crohn Disease Despite the High Variability of the Gut Microbiota. <i>Cell Reports</i> , 2016, 14, 945-955.	2.9	49
53	Meta-analysis of transcriptome data identifies a novel 5-gene pancreatic adenocarcinoma classifier. <i>Oncotarget</i> , 2016, 7, 23263-23281.	0.8	49
54	Identification of key regulators of pancreatic cancer progression through multidimensional systems-level analysis. <i>Genome Medicine</i> , 2016, 8, 38.	3.6	48

#	ARTICLE	IF	CITATIONS
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. <i>Prostate</i> , 2008, 68, 1517-1530.	1.2	47
56	Pentraxin-3 is a PI3K signaling target that promotes stem cell-like traits in basal-like breast cancers. <i>Science Signaling</i> , 2017, 10, .	1.6	43
57	Pericytes Elicit Resistance to Vemurafenib and Sorafenib Therapy in Thyroid Carcinoma via the TSP-1/TGF β 1 Axis. <i>Clinical Cancer Research</i> , 2018, 24, 6078-6097.	3.2	43
58	Requirement of the Epithelium-specific Ets Transcription Factor Spdef for Mucous Gland Cell Function in the Gastric Antrum. <i>Journal of Biological Chemistry</i> , 2010, 285, 35047-35055.	1.6	42
59	Downregulation of <i>Dipeptidyl Peptidase 4</i> Accelerates Progression to Castration-Resistant Prostate Cancer. <i>Cancer Research</i> , 2018, 78, 6354-6362.	0.4	42
60	TFEB-driven lysosomal biogenesis is pivotal for PGC1 α -dependent renal stress resistance. <i>JCI Insight</i> , 2019, 4, .	2.3	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. <i>Oncotarget</i> , 2017, 8, 84743-84760.	0.8	40
62	TAPPred Prediction of TAP-Binding Peptides in Antigens. <i>Methods in Molecular Biology</i> , 2007, 409, 381-386.	0.4	39
63	Prediction of Promiscuous and High-Affinity Mutated MHC Binders. <i>Hybridoma</i> , 2003, 22, 229-234.	0.6	38
64	Novel non-invasive biomarkers that distinguish between benign prostate hyperplasia and prostate cancer. <i>BMC Cancer</i> , 2015, 15, 259.	1.1	37
65	Determination of system level alterations in host transcriptome due to Zika virus (ZIKV) Infection in retinal pigment epithelium. <i>Scientific Reports</i> , 2018, 8, 11209.	1.6	37
66	The Role of TNF α in Mice with Type 1- and 2- Diabetes. <i>PLoS ONE</i> , 2012, 7, e33254.	1.1	35
67	Multicenter evaluation of a standardized protocol for noninvasive gene expression profiling. <i>American Journal of Transplantation</i> , 2013, 13, 1891-1897.	2.6	35
68	Chemotherapy-generated cell debris stimulates colon carcinoma tumor growth via osteopontin. <i>FASEB Journal</i> , 2019, 33, 114-125.	0.2	35
69	Mitochondrial Dysfunction in Atrial Tissue of Patients Developing Postoperative Atrial Fibrillation. <i>Annals of Thoracic Surgery</i> , 2017, 104, 1547-1555.	0.7	33
70	A20 Modulates Lipid Metabolism and Energy Production to Promote Liver Regeneration. <i>PLoS ONE</i> , 2011, 6, e17715.	1.1	33
71	Modulation of lymphocyte-mediated tissue repair by rational design of heterocyclic aryl hydrocarbon receptor agonists. <i>Science Advances</i> , 2020, 6, eaay8230.	4.7	31
72	Temporal retinal transcriptome and systems biology analysis identifies key pathways and hub genes in <i>Staphylococcus aureus</i> endophthalmitis. <i>Scientific Reports</i> , 2016, 6, 21502.	1.6	30

#	ARTICLE	IF	CITATIONS
73	A Phase II Trial of Abiraterone Combined with Dutasteride for Men with Metastatic Castration-Resistant Prostate Cancer. <i>Clinical Cancer Research</i> , 2017, 23, 935-945.	3.2	30
74	Smad2/3 pathway ligand trap luspatercept enhances erythroid differentiation in murine β -thalassaemia by increasing GATA-1 availability. <i>Journal of Cellular and Molecular Medicine</i> , 2020, 24, 6162-6177.	1.6	30
75	Pressure-overload hypertrophy of the developing heart reveals activation of divergent gene and protein pathways in the left and right ventricular myocardium. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2013, 304, H697-H708.	1.5	29
76	ACE2 abrogates tumor resistance to VEGFR inhibitors suggesting angiotensin-(1-7) as a therapy for clear cell renal cell carcinoma. <i>Science Translational Medicine</i> , 2021, 13, .	5.8	29
77	p16Ink4a or p19Arf loss contributes to Tal1-induced leukemogenesis in mice. <i>Oncogene</i> , 2006, 25, 3023-3031.	2.6	25
78	Quantitative proteomic analysis in HCV-induced HCC reveals sets of proteins with potential significance for racial disparity. <i>Journal of Translational Medicine</i> , 2013, 11, 239.	1.8	25
79	High dose intermittent sorafenib shows improved efficacy over conventional continuous dose in renal cell carcinoma. <i>Journal of Translational Medicine</i> , 2011, 9, 220.	1.8	24
80	HIV-1 Infection Impairs Regulatory T-Cell Suppressive Capacity on a Per-Cell Basis. <i>Journal of Infectious Diseases</i> , 2014, 210, 899-903.	1.9	24
81	Functional and Transcriptomic Characterization of Peritoneal Immune-Modulation by Addition of Alanine-Glutamine to Dialysis Fluid. <i>Scientific Reports</i> , 2017, 7, 6229.	1.6	24
82	Temporal Network Based Analysis of Cell Specific Vein Graft Transcriptome Defines Key Pathways and Hub Genes in Implantation Injury. <i>PLoS ONE</i> , 2012, 7, e39123.	1.1	23
83	SARS-CoV-2 morbidity and mortality in racial/ethnic minority populations: A window into the stress related inflammatory basis of health disparities?. <i>Brain, Behavior, & Immunity - Health</i> , 2020, 9, 100158.	1.3	22
84	Gene expression analysis of embryonic stem cells expressing VE-cadherin (CD144) during endothelial differentiation. <i>BMC Genomics</i> , 2008, 9, 240.	1.2	21
85	Preliminary Biomarkers for Identification of Human Ascending Thoracic Aortic Aneurysm. <i>Journal of the American Heart Association</i> , 2013, 2, e000138.	1.6	21
86	Inhibition of ALK1 signaling with dalantercept combined with VEGFR TKI leads to tumor stasis in renal cell carcinoma. <i>Oncotarget</i> , 0, 7, 41857-41869.	0.8	21
87	A20 Regulates Atherogenic Interferon (IFN)- γ Signaling in Vascular Cells by Modulating Basal IFN γ Levels. <i>Journal of Biological Chemistry</i> , 2014, 289, 30912-30924.	1.6	20
88	Induced Human Decidual NK-Like Cells Improve Utero-Placental Perfusion in Mice. <i>PLoS ONE</i> , 2016, 11, e0164353.	1.1	20
89	Transcriptomic and proteomic analysis of global ischemia and cardioprotection in the rabbit heart. <i>Physiological Genomics</i> , 2009, 38, 125-137.	1.0	19
90	Prevention of Nonimmunologic Loss of Transplanted Islets in Monkeys. <i>American Journal of Transplantation</i> , 2014, 14, 1543-1551.	2.6	19

#	ARTICLE	IF	CITATIONS
91	Leukemia vaccine overcomes limitations of checkpoint blockade by evoking clonal T cell responses in a murine acute myeloid leukemia model. <i>Haematologica</i> , 2021, 106, 1330-1342.	1.7	19
92	Binding of the WASP/N-WASP-Interacting Protein WIP to Actin Regulates Focal Adhesion Assembly and Adhesion. <i>Molecular and Cellular Biology</i> , 2014, 34, 2600-2610.	1.1	18
93	Microarray and proteomic analysis of the cardioprotective effects of cold blood cardioplegia in the mature and aged male and female. <i>Physiological Genomics</i> , 2012, 44, 1027-1041.	1.0	17
94	Significant lethality following liver resection in A20 heterozygous knockout mice uncovers a key role for A20 in liver regeneration. <i>Cell Death and Differentiation</i> , 2015, 22, 2068-2077.	5.0	17
95	The effects of transfection reagent polyethyleneimine (PEI) and non-targeting control siRNAs on global gene expression in human aortic smooth muscle cells. <i>BMC Genomics</i> , 2016, 17, 20.	1.2	17
96	Transcriptional Patterns in Peritoneal Tissue of Encapsulating Peritoneal Sclerosis, a Complication of Chronic Peritoneal Dialysis. <i>PLoS ONE</i> , 2013, 8, e56389.	1.1	17
97	Specific Transcriptome Changes Associated with Blood Pressure Reduction in Hypertensive Patients After Relaxation Response Training. <i>Journal of Alternative and Complementary Medicine</i> , 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. <i>Frontiers in Genetics</i> , 2020, 11, 572284.	1.1	15
99	Functional genomics in the study of mind-body therapies. <i>Ochsner Journal</i> , 2014, 14, 681-95.	0.5	15
100	Autocrine Canonical Wnt Signaling Primes Noncanonical Signaling through ROR1 in Metastatic Castration-Resistant Prostate Cancer. <i>Cancer Research</i> , 2022, 82, 1518-1533.	0.4	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. <i>Journal of Virology</i> , 2014, 88, 5559-5577.	1.5	14
102	Study of Cathepsin B inhibition in VEGFR TKI treated human renal cell carcinoma xenografts. <i>Oncogenesis</i> , 2019, 8, 15.	2.1	14
103	Analysis of Multiple Sarcoma Expression Datasets: Implications for Classification, Oncogenic Pathway Activation and Chemotherapy Resistance. <i>PLoS ONE</i> , 2010, 5, e9747.	1.1	14
104	Alterations of the gene expression profile in renal cell carcinoma after treatment with the histone deacetylase-inhibitor valproic acid and interferon-alpha. <i>World Journal of Urology</i> , 2011, 29, 779-786.	1.2	13
105	SB225002 Induces Cell Death and Cell Cycle Arrest in Acute Lymphoblastic Leukemia Cells through the Activation of GLIPR1. <i>PLoS ONE</i> , 2015, 10, e0134783.	1.1	13
106	Computational Methods in Genome Research. <i>Applied Mycology and Biotechnology</i> , 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. <i>Annals of Thoracic Surgery</i> , 2015, 100, 1636-1645.	0.7	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. <i>Blood</i> , 2015, 126, 751-751.	0.6	8

#	ARTICLE	IF	CITATIONS
109	Systems Biology Approach to Identify Novel Genomic Determinants for Pancreatic Cancer Pathogenesis. <i>Scientific Reports</i> , 2019, 9, 123.	1.6	7
110	A <i></i>Grammastola spatulata</i> mechanotoxin-4 (GsMTx4)-sensitive cation channel mediates increased cation permeability in human hereditary spherocytosis of multiple genetic etiologies. <i>Haematologica</i> , 2021, 106, 2759-2762.	1.7	5
111	Single Cell Transcriptomics Revealed AML and Non-AML Cell Clusters Relevant to Relapse and Remission in Pediatric AML. <i>Blood</i> , 2020, 136, 24-25.	0.6	5
112	Cellular and immunometabolic mechanisms of inflammation in depression: Preliminary findings from single cell RNA sequencing and a tribute to Bruce McEwen. <i>Neurobiology of Stress</i> , 2022, 19, 100462.	1.9	4
113	Leukocyte Transcriptional Response in Sepsis. <i>Shock</i> , 2019, 52, 166-173.	1.0	3
114	Embryonic periventricular endothelial cells demonstrate a unique pro-neurodevelopment and anti-inflammatory gene signature. <i>Scientific Reports</i> , 2020, 10, 20393.	1.6	3
115	Single Cell RNA-Seq Analyses of Healthy Lower Extremity Skin and Diabetic Foot Ulcers Uncover Distinct Immune Landscape of Diabetic Wound Healing. <i>Diabetes</i> , 2018, 67, 647-P.	0.3	3
116	<i>Pediatric Single Cell Cancer Atlas: An Integrative Web-Based Resource for Single Cell Transcriptome Data from Pediatric Leukemias</i>. <i>Blood</i> , 2021, 138, 3488-3488.	0.6	3
117	Comprehensive Genomic Profiling of High-Risk Pediatric Cancer Patients Has a Measurable Impact on Clinical Care. <i>JCO Precision Oncology</i> , 2022, 6, e2100451.	1.5	3
118	Searching and Mapping of T-Cell Epitopes, MHC Binders, and TAP Binders. <i>Methods in Molecular Biology</i> , 2007, 409, 95-112.	0.4	2
119	Integrated Cytof, Scrna-Seq and Cite-Seq Analysis of Bone Marrow Immune Microenvironment in the Mmrf Commpass Study. <i>Blood</i> , 2020, 136, 28-29.	0.6	2
120	Functional and Clinical Impact of Splicing Factor Dysregulation in Multiple Myeloma. <i>Blood</i> , 2015, 126, 726-726.	0.6	1
121	Characterization of T-Cell Exhaustion in Rapid Progressing Multiple Myeloma Using Cross Center Scrna-Seq Study. <i>Blood</i> , 2021, 138, 401-401.	0.6	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. <i>Blood</i> , 2021, 138, 1311-1311.	0.6	1
123	Survival Genie: A Web Portal for Single-Cell Data, Gene-Ratio, and Cell Composition-Based Survival Analyses. <i>Blood</i> , 2021, 138, 276-276.	0.6	1
124	Differential gene expression in vein graft endothelial and smooth muscle cells isolated by laser capture microdissection. <i>Journal of the American College of Surgeons</i> , 2009, 209, S137.	0.2	0
125	Identification of Inflammatory Bowel Disease (IBD)-Specific Diagnostic Biomarkers Using Transcriptional Profiling of Peripheral Blood Mononuclear Cells (PBMC) From Treatment Na ⁺ VE Children With Either IBD or Non-IBD Gastrointestinal Inflammatory Disorders. <i>Gastroenterology</i> , 2011, 140, S-509.	0.6	0
126	Genomic Determinants of a Relaxation Response Resiliency Program in Inflammatory Bowel Disease and Irritable Bowel Syndrome. <i>Gastroenterology</i> , 2011, 140, S-524.	0.6	0

#	ARTICLE	IF	CITATIONS
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. <i>Gastroenterology</i> , 2014, 146, S-216.	0.6	0
128	A Distinct Treg Transcriptome Signature in HIV-1 Elite Controllers Might Contribute to Improved Disease Outcome. <i>AIDS Research and Human Retroviruses</i> , 2014, 30, A180-A180.	0.5	0
129	Next-generation sequencing demonstrates genomic signature of resistance patterns following phosphatidylinositol 3-kinase (PI3K) inhibition. <i>Gynecologic Oncology</i> , 2015, 136, 406.	0.6	0
130	[31-OR]. <i>Pregnancy Hypertension</i> , 2015, 5, 15.	0.6	0
131	Notch1 Provides a Proliferative Signal in Mouse T Cell Leukemia by Directly Targeting c-Myc.. <i>Blood</i> , 2006, 108, 1434-1434.	0.6	0
132	Selective isolation of vein graft endothelial and smooth muscle cells using laser capture microdissection for microarray gene analysis. <i>FASEB Journal</i> , 2009, 23, LB322.	0.2	0
133	CARM1 is required for proper control of proliferation and differentiation of pulmonary epithelial cells. <i>Journal of Cell Science</i> , 2010, 123, e1-e1.	1.2	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. <i>Blood</i> , 2015, 126, 3633-3633.	0.6	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 Compass Study. <i>Blood</i> , 2021, 138, 2691-2691.	0.6	0
139	Mitochondrial Electron Transport Chain Inhibition Promotes Resistance to Proteasome Inhibitors in Multiple Myeloma. <i>Blood</i> , 2021, 138, 1611-1611.	0.6	0
140	<i>Single Cell RNA Sequencing Driven Characterization of Rare B/Myeloid and T/Myeloid Mixed Phenotype Acute Leukemia</i>. <i>Blood</i> , 2021, 138, 3455-3455.	0.6	0