Debashis Sahoo

List of Publications by Year in descending order

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70 papers

6,523 citations

147801 31 h-index 95266 68 g-index

88 all docs 88 docs citations

88 times ranked 11857 citing authors

#	Article	IF	CITATIONS
1	Isolation and characterization of human embryonic stem cell-derived heart field-specific cardiomyocytes unravels new insights into their transcriptional and electrophysiological profiles. Cardiovascular Research, 2022, 118, 828-843.	3.8	14
2	A Microsatellite in the Coding Sequence of HLA-A/B Is a Mutation Hotspot in Colon Cancer With Microsatellite Instability. Gastroenterology, 2022, 162, 960-963.e3.	1.3	0
3	Upregulation of BMI1-suppressor miRNAs (miR-200c, miR-203) during terminal differentiation of colon epithelial cells. Journal of Gastroenterology, 2022, , 1.	5.1	3
4	Artificial intelligence-rationalized balanced PPARÎ \pm Î3 dual agonism resets dysregulated macrophage processes in inflammatory bowel disease. Communications Biology, 2022, 5, 231.	4.4	7
5	An Artificial Intelligence-guided signature reveals the shared host immune response in MIS-C and Kawasaki disease. Nature Communications, 2022, 13, 2687.	12.8	37
6	Hepatocyte Deletion of IGF2 Prevents DNA Damage and Tumor Formation in Hepatocellular Carcinoma. Advanced Science, 2022, 9, .	11.2	9
7	A long isoform of GIV/Girdin contains a PDZ-binding module that regulates localization and G-protein binding. Journal of Biological Chemistry, 2021, 296, 100493.	3.4	8
8	Boolean implication analysis unveils candidate universal relationships in microbiome data. BMC Bioinformatics, 2021, 22, 49.	2.6	1
9	E-cigarettes compromise the gut barrier and trigger inflammation. IScience, 2021, 24, 102035.	4.1	36
10	A zebrafish model of granulin deficiency reveals essential roles in myeloid cell differentiation. Blood Advances, 2021, 5, 796-811.	5 . 2	17
11	Immunosuppression of Macrophages Underlies the Cardioprotective Effects of CST (Catestatin). Hypertension, 2021, 77, 1670-1682.	2.7	31
12	Al-guided discovery of the invariant host response to viral pandemics. EBioMedicine, 2021, 68, 103390.	6.1	37
13	Drug repurposing screens identify chemical entities for the development of COVID-19 interventions. Nature Communications, 2021, 12, 3309.	12.8	81
14	Artificial intelligence guided discovery of a barrier-protective therapy in inflammatory bowel disease. Nature Communications, 2021, 12, 4246.	12.8	37
15	GIV/Girdin, a non-receptor modulator for $\widehat{Gl}\pm i/s$, regulates spatiotemporal signaling during sperm capacitation and is required for male fertility. ELife, 2021, 10, .	6.0	7
16	Adult stem cell-derived complete lung organoid models emulate lung disease in COVID-19. ELife, 2021, 10, .	6.0	64
17	Aged skeletal stem cells generate an inflammatory degenerative niche. Nature, 2021, 597, 256-262.	27.8	143
18	SPT6 loss permits the transdifferentiation of keratinocytes into an intestinal fate that resembles Barrett's metaplasia. IScience, 2021, 24, 103121.	4.1	5

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19	The Gαâ€interacting vesicleâ€associated protein interacts with and promotes cell surface localization of GRP78 during endoplasmic reticulum stress. FEBS Letters, 2020, 594, 1088-1100.	2.8	5
20	Urothelial Proliferation of Unknown Malignant Potential Involving the Bladder: Histopathologic Features and Risk of Progression in De Novo Cases and Cases With Prior Neoplasia. Archives of Pathology and Laboratory Medicine, 2020, 144, 853-862.	2.5	8
21	GIV•Kindlin Interaction Is Required for Kindlin-Mediated Integrin Recognition and Activation. IScience, 2020, 23, 101209.	4.1	11
22	Haematopoietic stem cell-dependent Notch transcription is mediated by p53 through the Histone chaperone Supt16h. Nature Cell Biology, 2020, 22, 1411-1422.	10.3	9
23	GIV/Girdin and Exo70 Collaboratively Regulate the Mammalian Polarized Exocytic Machinery. IScience, 2020, 23, 101246.	4.1	3
24	The DNA Glycosylase NEIL2 Suppresses Fusobacterium-Infection-Induced Inflammation and DNA Damage in Colonic Epithelial Cells. Cells, 2020, 9, 1980.	4.1	28
25	The PVT1 lncRNA is a novel epigenetic enhancer of MYC, and a promising risk-stratification biomarker in colorectal cancer. Molecular Cancer, 2020, 19, 155.	19.2	56
26	Transcriptional profiling of lung macrophages identifies a predictive signature for inflammatory lung disease in preterm infants. Communications Biology, 2020, 3, 259.	4.4	25
27	Upregulation of CD47 Is a Host Checkpoint Response to Pathogen Recognition. MBio, 2020, 11, .	4.1	29
28	Helicobacter pylori infection downregulates the DNA glycosylase NEIL2, resulting in increased genome damage and inflammation in gastric epithelial cells. Journal of Biological Chemistry, 2020, 295, 11082-11098.	3.4	35
29	Enhanced mitochondrial fission suppresses signaling and metastasis in triple-negative breast cancer. Breast Cancer Research, 2020, 22, 60.	5.0	46
30	Computational Approach to Identifying Universal Macrophage Biomarkers. Frontiers in Physiology, 2020, 11, 275.	2.8	26
31	Primary Female Urethral Carcinoma. American Journal of Surgical Pathology, 2020, 44, 1591-1601.	3.7	11
32	The stress polarity signaling (SPS) pathway serves as a marker and a target in the leaky gut barrier: implications in aging and cancer. Life Science Alliance, 2020, 3, e201900481.	2.8	28
33	DDX5 promotes oncogene C3 and FABP1 expressions and drives intestinal inflammation and tumorigenesis. Life Science Alliance, 2020, 3, e202000772.	2.8	21
34	miR-221 Targets QKI to Enhance the Tumorigenic Capacity of Human Colorectal Cancer Stem Cells. Cancer Research, 2019, 79, 5151-5158.	0.9	51
35	Non-coding and Coding Transcriptional Profiles Are Significantly Altered in Pediatric Retinoblastoma Tumors. Frontiers in Oncology, 2019, 9, 221.	2.8	27
36	Unbiased Boolean analysis of public gene expression data for cell cycle gene identification. Molecular Biology of the Cell, 2019, 30, 1770-1779.	2.1	28

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37	Identification of gene expression logical invariants in <i>Arabidopsis</i> . Plant Direct, 2019, 3, e00123.	1.9	7
38	<i>Fusobacterium nucleatum</i> promotes colorectal cancer by inducing Wnt/β atenin modulator Annexin A1. EMBO Reports, 2019, 20, .	4.5	283
39	High fat diet induces sex-specific differential gene expression in Drosophila melanogaster. PLoS ONE, 2019, 14, e0213474.	2.5	27
40	Analysis of cardiomyocyte clonal expansion during mouse heart development and injury. Nature Communications, 2018, 9, 754.	12.8	94
41	Notch1 regulates the initiation of metastasis and self-renewal of Group 3 medulloblastoma. Nature Communications, 2018, 9, 4121.	12.8	36
42	Identification of the Human Skeletal Stem Cell. Cell, 2018, 175, 43-56.e21.	28.9	425
43	Boolean analysis identifies CD38 as a biomarker of aggressive localized prostate cancer. Oncotarget, 2018, 9, 6550-6561.	1.8	16
44	Deep Sequencing of Urinary RNAs for Bladder Cancer Molecular Diagnostics. Clinical Cancer Research, 2017, 23, 3700-3710.	7.0	29
45	The innate immune response in fetal lung mesenchymal cells targets VEGFR2 expression and activity. American Journal of Physiology - Lung Cellular and Molecular Physiology, 2017, 312, L861-L872.	2.9	4
46	A Quiescent Bcl11b High Stem Cell Population Is Required for Maintenance of the Mammary Gland. Cell Stem Cell, 2017, 20, 247-260.e5.	11.1	86
47	Identification of tumorigenic cells and therapeutic targets in pancreatic neuroendocrine tumors. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 4464-4469.	7.1	70
48	Antibody Therapy Targeting CD47 and CD271 Effectively Suppresses Melanoma Metastasis in Patient-Derived Xenografts. Cell Reports, 2016, 16, 1701-1716.	6.4	56
49	CD13 and ROR2 Permit Isolation of Highly Enriched Cardiac Mesoderm from Differentiating Human Embryonic Stem Cells. Stem Cell Reports, 2016, 6, 95-108.	4.8	30
50	CDX2 as a Prognostic Biomarker in Colon Cancer. New England Journal of Medicine, 2016, 374, 2182-2184.	27.0	23
51	CDX2 as a Prognostic Biomarker in Stage II and Stage III Colon Cancer. New England Journal of Medicine, 2016, 374, 211-222.	27.0	388
52	Hoxb5 marks long-term haematopoietic stem cells and reveals a homogenous perivascular niche. Nature, 2016, 530, 223-227.	27.8	275
53	Identification and Specification of the Mouse Skeletal Stem Cell. Cell, 2015, 160, 285-298.	28.9	571
54	CD14-expressing cancer cells establish the inflammatory and proliferative tumor microenvironment in bladder cancer. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 4725-4730.	7.1	83

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55	Endoscopic molecular imaging of human bladder cancer using a CD47 antibody. Science Translational Medicine, 2014, 6, 260ra148.	12.4	124
56	A cell-intrinsic role for TLR2–MYD88 in intestinal and breast epithelia and oncogenesis. Nature Cell Biology, 2014, 16, 1238-1248.	10.3	106
57	Hedgehog Signaling Restrains Bladder Cancer Progression by Eliciting Stromal Production of Urothelial Differentiation Factors. Cancer Cell, 2014, 26, 521-533.	16.8	164
58	Discriminating cellular heterogeneity using microwell-based RNA cytometry. Nature Communications, 2014, 5, 3451.	12.8	49
59	Clonal precursor of bone, cartilage, and hematopoietic niche stromal cells. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 12643-12648.	7.1	116
60	The Power of Boolean Implication Networks. Frontiers in Physiology, 2012, 3, 276.	2.8	14
61	Three differentiation states risk-stratify bladder cancer into distinct subtypes. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 2078-2083.	7.1	232
62	Gene Expression Commons: An Open Platform for Absolute Gene Expression Profiling. PLoS ONE, 2012, 7, e40321.	2.5	227
63	Human bone marrow hematopoietic stem cells are increased in frequency and myeloid-biased with age. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 20012-20017.	7.1	730
64	Single-cell dissection of transcriptional heterogeneity in human colon tumors. Nature Biotechnology, 2011, 29, 1120-1127.	17.5	658
65	MiDReG: A method of mining developmentally regulated genes using Boolean implications. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 5732-5737.	7.1	57
66	Ly6d marks the earliest stage of B-cell specification and identifies the branchpoint between B-cell and T-cell development. Genes and Development, 2009, 23, 2376-2381.	5.9	254
67	Boolean implication networks derived from large scale, whole genome microarray datasets. Genome Biology, 2008, 9, R157.	9.6	108
68	Extracting binary signals from microarray time-course data. Nucleic Acids Research, 2007, 35, 3705-3712.	14.5	145
69	Genome Wide Association Study of Pregnancy in Childhood Cancer Survivors: A Report from the Childhood Cancer Survivor Study. Cancer Epidemiology Biomarkers and Prevention, 0, , .	2.5	0
70	<i>LiteCON</i> : An All-Photonic Neuromorphic Accelerator for Energy-efficient Deep Learning. Transactions on Architecture and Code Optimization, 0, , .	2.0	0