

Sandeep S Dave

List of Publications by Year in descending order

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Version: 2024-02-01

44
papers

4,144
citations

394421

19
h-index

395702

33
g-index

46
all docs

46
docs citations

46
times ranked

6332
citing authors

#	ARTICLE	IF	CITATIONS
1	Aspirin effects on platelet gene expression are associated with a paradoxical, increase in platelet function. <i>British Journal of Clinical Pharmacology</i> , 2022, 88, 2074-2083.	2.4	4
2	The 5th edition of the World Health Organization Classification of Haematolymphoid Tumours: Lymphoid Neoplasms. <i>Leukemia</i> , 2022, 36, 1720-1748.	7.2	1,023
3	Clinical relevance of molecular characteristics in Burkitt lymphoma differs according to age. <i>Nature Communications</i> , 2022, 13, .	12.8	28
4	Non-Hodgkin Lymphomas: Malignancies Arising from Mature B Cells. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2021, 11, a034843.	6.2	3
5	Monocarboxylate transporter antagonism reveals metabolic vulnerabilities of viral-driven lymphomas. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	34
6	Oncogenic Integration of Nucleotide Metabolism via Fatty Acid Synthase in Non-Hodgkin Lymphoma. <i>Frontiers in Oncology</i> , 2021, 11, 725137.	2.8	7
7	ALK-Negative Anaplastic Large Cell Lymphomas Encompass Distinct Subgroups Including an ALK-Positive-like Subgroup with Favorable Prognosis. <i>Blood</i> , 2021, 138, 2403-2403.	1.4	1
8	The Atlas of Blood Cancer Genomes (ABCG) Project: A Comprehensive Molecular Characterization of Leukemias and Lymphomas. <i>Blood</i> , 2021, 138, 2213-2213.	1.4	0
9	Genome-defined African ancestry is associated with distinct mutations and worse survival in patients with diffuse large B-cell lymphoma. <i>Cancer</i> , 2020, 126, 3493-3503.	4.1	15
10	PD-L1 expression is low in large B-cell lymphoma with MYC or double-hit translocation. <i>Hematological Oncology</i> , 2019, 37, 375-382.	1.7	8
11	Genetic convergence of rare lymphomas. <i>Current Opinion in Hematology</i> , 2018, 25, 307-314.	2.5	2
12	TET2 Deficiency Sets the Stage for B-cell Lymphoma. <i>Cancer Discovery</i> , 2018, 8, 1515-1517.	9.4	4
13	Gene essentiality landscape and druggable oncogenic dependencies in herpesviral primary effusion lymphoma. <i>Nature Communications</i> , 2018, 9, 3263.	12.8	50
14	Plasmodium parasite exploits host aquaporin-3 during liver stage malaria infection. <i>PLoS Pathogens</i> , 2018, 14, e1007057.	4.7	51
15	The Genetic Basis of Hepatosplenic T-cell Lymphoma. <i>Cancer Discovery</i> , 2017, 7, 369-379.	9.4	163
16	Human Mesenchymal Stem Cell-educated Macrophages Are a Distinct High IL-6-producing Subset that Confer Protection in Graft-versus-Host-Disease and Radiation Injury Models. <i>Biology of Blood and Marrow Transplantation</i> , 2017, 23, 897-905.	2.0	49
17	Enteropathy-associated T cell lymphoma subtypes are characterized by loss of function of SETD2. <i>Journal of Experimental Medicine</i> , 2017, 214, 1371-1386.	8.5	144
18	Genetic and Functional Drivers of Diffuse Large B-Cell Lymphoma. <i>Cell</i> , 2017, 171, 481-494.e15.	28.9	804

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19	Adult high-grade B-cell lymphoma with Burkitt lymphoma signature: genomic features and potential therapeutic targets. <i>Blood</i> , 2017, 130, 1819-1831.	1.4	62
20	Generation and comparison of CRISPR-Cas9 and Cre-mediated genetically engineered mouse models of sarcoma. <i>Nature Communications</i> , 2017, 8, 15999.	12.8	53
21	Epstein-Barr Virus Induces Adhesion Receptor CD226 (DNAM-1) Expression during Primary B-Cell Transformation into Lymphoblastoid Cell Lines. <i>MSphere</i> , 2017, 2, .	2.9	8
22	GNA13 loss in germinal center B cells leads to impaired apoptosis and promotes lymphoma in vivo. <i>Blood</i> , 2016, 127, 2723-2731.	1.4	52
23	Comprehensive Genomic Analysis of Adult Burkitt Lymphoma Identifies the B-Cell Receptor Signaling Pathway As a Potential Therapeutic Target. <i>Blood</i> , 2016, 128, 4095-4095.	1.4	0
24	SMRT Sequencing for Parallel Analysis of Multiple Targets and Accurate SNP Phasing. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 2801-2808.	1.8	25
25	A comprehensive joint analysis of the long and short RNA transcriptomes of human erythrocytes. <i>BMC Genomics</i> , 2015, 16, 952.	2.8	90
26	The Role of EBV in the Pathogenesis of Diffuse Large B Cell Lymphoma. <i>Current Topics in Microbiology and Immunology</i> , 2015, 390, 315-337.	1.1	23
27	ID3 Loss In Vivo Synergizes with MYC-Driven Oncogenesis in Burkitt Lymphoma. <i>Blood</i> , 2015, 126, 3906-3906.	1.4	1
28	Whole Exome Sequencing of Type 1 and Type 2 Enteropathy-Associated T Cell Lymphoma Reveals Genetic Basis of Eat1 Oncogenesis. <i>Blood</i> , 2015, 126, 575-575.	1.4	3
29	Exome-Wide Sequencing Reveals Oncogenic Mutations in Both Progressive and Non-Progressive MBL. <i>Blood</i> , 2015, 126, 361-361.	1.4	0
30	Recurrent Mutations in Focal Adhesion Genes Including the RhoA Signaling Axis Are a Defining Feature of Germinal Center (GC) Derived B Cell Lymphomas and Promote Aberrant Migration and Cellular Adhesion within the GC Niche. <i>Blood</i> , 2015, 126, 558-558.	1.4	0
31	Whole Exome Sequencing Defines the Genetic Differences Between Pediatric and Adult Follicular Lymphoma. <i>Blood</i> , 2015, 126, 3636-3636.	1.4	0
32	The Landscape of microRNA Expression in HIV and Non-HIV Associated Classical Hodgkin Lymphoma through Next Generation Sequencing. <i>Blood</i> , 2015, 126, 2639-2639.	1.4	0
33	The genomic landscape of mantle cell lymphoma is related to the epigenetically determined chromatin state of normal B cells. <i>Blood</i> , 2014, 123, 2988-2996.	1.4	224
34	Chemical Genomics Reveals JAK STAT Activation As a Mechanism of Resistance to HDAC Inhibitors in B Cell Lymphomas. <i>Blood</i> , 2014, 124, 271-271.	1.4	1
35	Silencing Mutations in NOTCH1 Activate Calcium Signaling in B Cells. <i>Blood</i> , 2014, 124, 1696-1696.	1.4	0
36	Integrative Genomics Reveals a Role for GNA13 in Lymphomagenesis within the Germinal Center Niche. <i>Blood</i> , 2014, 124, 782-782.	1.4	0

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37	Genetic heterogeneity of diffuse large B-cell lymphoma. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 1398-1403.	7.1	494
38	Evaluating oncogenic pathway dysregulation in adolescents and young adults with acute myeloid leukemia.. Journal of Clinical Oncology, 2013, 31, 7107-7107.	1.6	0
39	The Genetic Landscape Of Mantle Cell Lymphoma and The Epigenetic Origins Of Lineage Specific Mutations. Blood, 2013, 122, 347-347.	1.4	0
40	The genetic landscape of mutations in Burkitt lymphoma. Nature Genetics, 2012, 44, 1321-1325.	21.4	517
41	Burkitt lymphoma: analysis of the genome. Clinical Advances in Hematology and Oncology, 2012, 10, 54-5.	0.3	3
42	Host Factors for Risk and Survival in Lymphoma. Hematology American Society of Hematology Education Program, 2010, 2010, 255-258.	2.5	9
43	Deep sequencing of the small RNA transcriptome of normal and malignant human B cells identifies hundreds of novel microRNAs. Blood, 2010, 116, e118-e127.	1.4	188
44	PI3K Inhibitors Inhibit Lymphoma Growth by Downregulation of MYC-Dependent Proliferation.. Blood, 2009, 114, 1697-1697.	1.4	1