

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 | 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 |
| 2 | Genetic and Functional Drivers of Diffuse Large B-Cell Lymphoma. <i>Cell</i> , 2017, 171, 481-494.e15. | 28.9 | 804 |
| 3 | The genetic landscape of mutations in Burkitt lymphoma. <i>Nature Genetics</i> , 2012, 44, 1321-1325. | 21.4 | 517 |
| 4 | Genetic heterogeneity of diffuse large B-cell lymphoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 1398-1403. | 7.1 | 494 |
| 5 | 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 |
| 6 | Deep sequencing of the small RNA transcriptome of normal and malignant human B cells identifies hundreds of novel microRNAs. <i>Blood</i> , 2010, 116, e118-e127. | 1.4 | 188 |
| 7 | The Genetic Basis of Hepatosplenic T-cell Lymphoma. <i>Cancer Discovery</i> , 2017, 7, 369-379. | 9.4 | 163 |
| 8 | 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 |
| 9 | A comprehensive joint analysis of the long and short RNA transcriptomes of human erythrocytes. <i>BMC Genomics</i> , 2015, 16, 952. | 2.8 | 90 |
| 10 | 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 |
| 11 | 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 |
| 12 | 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 |
| 13 | Plasmodium parasite exploits host aquaporin-3 during liver stage malaria infection. <i>PLoS Pathogens</i> , 2018, 14, e1007057. | 4.7 | 51 |
| 14 | Gene essentiality landscape and druggable oncogenic dependencies in herpesviral primary effusion lymphoma. <i>Nature Communications</i> , 2018, 9, 3263. | 12.8 | 50 |
| 15 | 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 |
| 16 | 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 |
| 17 | Clinical relevance of molecular characteristics in Burkitt lymphoma differs according to age. <i>Nature Communications</i> , 2022, 13, . | 12.8 | 28 |
| 18 | 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 |

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|----|--|-----|-----------|
| 19 | 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 |
| 20 | 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 |
| 21 | Host Factors for Risk and Survival in Lymphoma. <i>Hematology American Society of Hematology Education Program</i> , 2010, 2010, 255-258. | 2.5 | 9 |
| 22 | 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 |
| 23 | 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 |
| 24 | Oncogenic Integration of Nucleotide Metabolism via Fatty Acid Synthase in Non-Hodgkin Lymphoma. <i>Frontiers in Oncology</i> , 2021, 11, 725137. | 2.8 | 7 |
| 25 | TET2 Deficiency Sets the Stage for B-cell Lymphoma. <i>Cancer Discovery</i> , 2018, 8, 1515-1517. | 9.4 | 4 |
| 26 | 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 |
| 27 | Non-Hodgkin Lymphomas: Malignancies Arising from Mature B Cells. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2021, 11, a034843. | 6.2 | 3 |
| 28 | Whole Exome Sequencing of Type 1 and Type 2 Enteropathy-Associated T Cell Lymphoma Reveals Genetic Basis of EATL Oncogenesis. <i>Blood</i> , 2015, 126, 575-575. | 1.4 | 3 |
| 29 | Burkitt lymphoma: analysis of the genome. <i>Clinical Advances in Hematology and Oncology</i> , 2012, 10, 54-5. | 0.3 | 3 |
| 30 | Genetic convergence of rare lymphomas. <i>Current Opinion in Hematology</i> , 2018, 25, 307-314. | 2.5 | 2 |
| 31 | PI3K Inhibitors Inhibit Lymphoma Growth by Downregulation of MYC-Dependent Proliferation.. <i>Blood</i> , 2009, 114, 1697-1697. | 1.4 | 1 |
| 32 | ID3 Loss In Vivo Synergizes with MYC-Driven Oncogenesis in Burkitt Lymphoma. <i>Blood</i> , 2015, 126, 3906-3906. | 1.4 | 1 |
| 33 | 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 |
| 34 | 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 |
| 35 | Evaluating oncogenic pathway dysregulation in adolescents and young adults with acute myeloid leukemia.. <i>Journal of Clinical Oncology</i> , 2013, 31, 7107-7107. | 1.6 | 0 |
| 36 | The Genetic Landscape Of Mantle Cell Lymphoma and The Epigenetic Origins Of Lineage Specific Mutations. <i>Blood</i> , 2013, 122, 347-347. | 1.4 | 0 |

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|----|--|-----|-----------|
| 37 | Silencing Mutations in NOTCH1 Activate Calcium Signaling in B Cells. Blood, 2014, 124, 1696-1696. | 1.4 | 0 |
| 38 | Integrative Genomics Reveals a Role for GNA13 in Lymphomagenesis within the Germinal Center Niche. Blood, 2014, 124, 782-782. | 1.4 | 0 |
| 39 | Exome-Wide Sequencing Reveals Oncogenic Mutations in Both Progressive and Non-Progressive MBL. Blood, 2015, 126, 361-361. | 1.4 | 0 |
| 40 | 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. Blood, 2015, 126, 558-558. | 1.4 | 0 |
| 41 | Whole Exome Sequencing Defines the Genetic Differences Between Pediatric and Adult Follicular Lymphoma. Blood, 2015, 126, 3636-3636. | 1.4 | 0 |
| 42 | The Landscape of microRNA Expression in HIV and Non-HIV Associated Classical Hodgkin Lymphoma through Next Generation Sequencing. Blood, 2015, 126, 2639-2639. | 1.4 | 0 |
| 43 | Comprehensive Genomic Analysis of Adult Burkitt Lymphoma Identifies the B-Cell Receptor Signaling Pathway As a Potential Therapeutic Target. Blood, 2016, 128, 4095-4095. | 1.4 | 0 |
| 44 | The Atlas of Blood Cancer Genomes (ABCG) Project: A Comprehensive Molecular Characterization of Leukemias and Lymphomas. Blood, 2021, 138, 2213-2213. | 1.4 | 0 |