

Jyoti Nangalia

List of Publications by Year in descending order

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Version: 2024-02-01

43
papers

3,670
citations

201674

27
h-index

289244

40
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48
all docs

48
docs citations

48
times ranked

5076
citing authors

#	ARTICLE	IF	CITATIONS
1	Life histories of myeloproliferative neoplasms inferred from phylogenies. <i>Nature</i> , 2022, 602, 162-168.	27.8	140
2	Bayesian networks elucidate complex genomic landscapes in cancer. <i>Communications Biology</i> , 2022, 5, 306.	4.4	5
3	The longitudinal dynamics and natural history of clonal haematopoiesis. <i>Nature</i> , 2022, 606, 335-342.	27.8	136
4	Clonal dynamics of haematopoiesis across the human lifespan. <i>Nature</i> , 2022, 606, 343-350.	27.8	160
5	Clonal hematopoiesis and therapy-related myeloid neoplasms following neuroblastoma treatment. <i>Blood</i> , 2021, 137, 2992-2997.	1.4	19
6	Lineage tracing of human development through somatic mutations. <i>Nature</i> , 2021, 595, 85-90.	27.8	79
7	Unmet clinical needs in the management of CALR-mutated essential thrombocythaemia: a consensus-based proposal from the European LeukemiaNet. <i>Lancet Haematology</i> , 2021, 8, e658-e665.	4.6	17
8	Caught in the antiviral crossfire: Ganciclovir-associated mutagenesis in HSC transplant recipients. <i>Cell Stem Cell</i> , 2021, 28, 1683-1685.	11.1	1
9	Changes in Clonal Architecture Inform MPN Disease Course in Advance of Phenotypic Manifestations. <i>Blood</i> , 2021, 138, 3590-3590.	1.4	1
10	Inherited Blood Cancer Predisposition through Altered Transcription Elongation. <i>Blood</i> , 2021, 138, 629-629.	1.4	0
11	Long-Term Clonal Dynamics upon Allogeneic Haematopoietic Stem Cell Transplantation Revealed Using Somatic Mutations As Clonal Tracking Marks in HSC Donor and Recipient Pairs. <i>Blood</i> , 2021, 138, 3813-3813.	1.4	0
12	Genomic heterogeneity in myeloproliferative neoplasms and applications to clinical practice. <i>Blood Reviews</i> , 2020, 42, 100708.	5.7	10
13	The mutational landscape of normal human endometrial epithelium. <i>Nature</i> , 2020, 580, 640-646.	27.8	338
14	Extensive heterogeneity in somatic mutation and selection in the human bladder. <i>Science</i> , 2020, 370, 75-82.	12.6	195
15	Methylation age as a correlate for allele burden, disease status, and clinical response in myeloproliferative neoplasm patients treated with vorinostat. <i>Experimental Hematology</i> , 2019, 79, 26-34.	0.4	8
16	Clonal approaches to understanding the impact of mutations on hematologic disease development. <i>Blood</i> , 2019, 133, 1436-1445.	1.4	14
17	Genome Sequencing during a Patient's Journey through Cancer. <i>New England Journal of Medicine</i> , 2019, 381, 2145-2156.	27.0	50
18	A guideline for the management of specific situations in polycythaemia vera and secondary erythrocytosis. <i>British Journal of Haematology</i> , 2019, 184, 161-175.	2.5	76

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19	A guideline for the diagnosis and management of polycythaemia vera. A British Society for Haematology Guideline. <i>British Journal of Haematology</i> , 2019, 184, 176-191.	2.5	102
20	Hydroxycarbamide Plus Aspirin Versus Aspirin Alone in Patients With Essential Thrombocythemia Age 40 to 59 Years Without High-Risk Features. <i>Journal of Clinical Oncology</i> , 2018, 36, 3361-3369.	1.6	54
21	Classification and Personalized Prognosis in Myeloproliferative Neoplasms. <i>New England Journal of Medicine</i> , 2018, 379, 1416-1430.	27.0	442
22	Pacritinib versus best available therapy for the treatment of myelofibrosis irrespective of baseline cytopenias (PERSIST-1): an international, randomised, phase 3 trial. <i>Lancet Haematology</i> , 2017, 4, e225-e236.	4.6	224
23	Molecular determinants of pathogenesis and clinical phenotype in myeloproliferative neoplasms. <i>Haematologica</i> , 2017, 102, 7-17.	3.5	74
24	Myeloproliferative neoplasms: from origins to outcomes. <i>Blood</i> , 2017, 130, 2475-2483.	1.4	107
25	Pathogenesis of Myeloproliferative Disorders. <i>Annual Review of Pathology: Mechanisms of Disease</i> , 2016, 11, 101-126.	22.4	38
26	Megakaryocytic hyperplasia in myeloproliferative neoplasms is driven by disordered proliferative, apoptotic and epigenetic mechanisms. <i>Journal of Clinical Pathology</i> , 2016, 69, 155-163.	2.0	25
27	Relationship of JAK2V617F Allelic Burden (AB) to Demographics, Disease Characteristics, and Response to Therapy in Persist-1, a Randomized Phase III Study of Pacritinib (PAC) Versus Best Available Therapy (BAT) in Patients (pts) with Primary and Secondary Myelofibrosis (MF). <i>Blood</i> , 2016, 128, 3131-3131.	1.4	2
28	RECQL5 Suppresses Oncogenic JAK2-Induced Replication Stress and Genomic Instability. <i>Cell Reports</i> , 2015, 13, 2345-2352.	6.4	28
29	DNMT3A mutations occur early or late in patients with myeloproliferative neoplasms and mutation order influences phenotype. <i>Haematologica</i> , 2015, 100, e438-e442.	3.5	105
30	Effect of Mutation Order on Myeloproliferative Neoplasms. <i>New England Journal of Medicine</i> , 2015, 372, 601-612.	27.0	467
31	Frequent somatic transfer of mitochondrial DNA into the nuclear genome of human cancer cells. <i>Genome Research</i> , 2015, 25, 814-824.	5.5	69
32	Genetic variation at MECOM, TERT, JAK2 and HBS1L-MYB predisposes to myeloproliferative neoplasms. <i>Nature Communications</i> , 2015, 6, 6691.	12.8	145
33	Evaluation of methods to detect CALR mutations in myeloproliferative neoplasms. <i>Leukemia Research</i> , 2015, 39, 82-87.	0.8	55
34	The evolving genomic landscape of myeloproliferative neoplasms. <i>Hematology American Society of Hematology Education Program</i> , 2014, 2014, 287-296.	2.5	62
35	JAK2V617F homozygosity drives a phenotypic switch in myeloproliferative neoplasms, but is insufficient to sustain disease. <i>Blood</i> , 2014, 123, 3139-3151.	1.4	77
36	<i>CALR</i> mutations in myeloproliferative neoplasms: Hidden behind the reticulum. <i>American Journal of Hematology</i> , 2014, 89, 453-456.	4.1	34

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37	Inactivating CUX1 mutations promote tumorigenesis. <i>Nature Genetics</i> , 2014, 46, 33-38.	21.4	111
38	JAK2V617F promotes replication fork stalling with disease-restricted impairment of the intra-S checkpoint response. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 15190-15195.	7.1	35
39	Nongenetic stochastic expansion of JAK2V617F-homozygous subclones in polycythemia vera?. <i>Blood</i> , 2014, 124, 3332-3334.	1.4	3
40	Cooperativity of imprinted genes inactivated by acquired chromosome 20q deletions. <i>Journal of Clinical Investigation</i> , 2013, 123, 2169-2182.	8.2	36
41	Outcome of Refractory Anemia with Ringed Sideroblasts Associated with Marked Thrombocytosis (RARS-T) In a Large Cohort of Patients. <i>Blood</i> , 2010, 116, 4113-4113.	1.4	1
42	Isolated neutropenia during ABVD chemotherapy for Hodgkin lymphoma does not require growth factor support. <i>Leukemia and Lymphoma</i> , 2008, 49, 1530-1536.	1.3	11
43	Molecular basis of thrombin recognition by protein C inhibitor revealed by the 1.6-Å structure of the heparin-bridged complex. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 4661-4666.	7.1	59