

Chung H Kok

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

52
papers

953
citations

14
h-index

30
g-index

57
ext. papers

1,134
ext. citations

4.9
avg, IF

3.25
L-index

#	Paper	IF	Citations
52	Heritable GATA2 mutations associated with familial myelodysplastic syndrome and acute myeloid leukemia. <i>Nature Genetics</i> , 2011 , 43, 1012-7	36.3	424
51	Ecotopic viral integration site 1 (EVI1) regulates multiple cellular processes important for cancer and is a synergistic partner for FOS protein in invasive tumors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 2168-73	11.5	61
50	Expression profiling of a hemopoietic cell survival transcriptome implicates osteopontin as a functional prognostic factor in AML. <i>Blood</i> , 2009 , 114, 4859-70	2.2	51
49	HoxA9 regulated Bcl-2 expression mediates survival of myeloid progenitors and the severity of HoxA9-dependent leukemia. <i>Oncotarget</i> , 2013 , 4, 1933-47	3.3	40
48	Long-term treatment-free remission of chronic myeloid leukemia with falling levels of residual leukemic cells. <i>Leukemia</i> , 2018 , 32, 2572-2579	10.7	37
47	High prevalence of relapse in children with Philadelphia-like acute lymphoblastic leukemia despite risk-adapted treatment. <i>Haematologica</i> , 2017 , 102, e490-e493	6.6	36
46	Genetic regulators of myelopoiesis and leukemic signaling identified by gene profiling and linear modeling. <i>Journal of Leukocyte Biology</i> , 2006 , 80, 433-47	6.5	28
45	The granulocyte-associated transcription factor Krüppel-like factor 5 is silenced by hypermethylation in acute myeloid leukemia. <i>Leukemia Research</i> , 2012 , 36, 110-6	2.7	23
44	PTTG1 expression is associated with hyperproliferative disease and poor prognosis in multiple myeloma. <i>Journal of Hematology and Oncology</i> , 2015 , 8, 106	22.4	23
43	Lineage of measurable residual disease in patients with chronic myeloid leukemia in treatment-free remission. <i>Leukemia</i> , 2020 , 34, 1052-1061	10.7	23
42	Successful treatment-free remission in chronic myeloid leukaemia and its association with reduced immune suppressors and increased natural killer cells. <i>British Journal of Haematology</i> , 2020 , 191, 433-441	4.5	22
41	Gene expression signature that predicts early molecular response failure in chronic-phase CML patients on frontline imatinib. <i>Blood Advances</i> , 2019 , 3, 1610-1621	7.8	20
40	Differential expression of MUC4, GPR110 and IL2RA defines two groups of CRLF2-rearranged acute lymphoblastic leukemia patients with distinct secondary lesions. <i>Cancer Letters</i> , 2017 , 408, 92-101	9.9	17
39	The GM-CSF receptor utilizes Ecatenin and Tcf4 to specify macrophage lineage differentiation. <i>Differentiation</i> , 2012 , 83, 47-59	3.5	16
38	Pre-B acute lymphoblastic leukaemia recurrent fusion, EP300-ZNF384, is associated with a distinct gene expression. <i>British Journal of Cancer</i> , 2018 , 118, 1000-1004	8.7	14
37	Methylation of KLF5 contributes to reduced expression in acute myeloid leukaemia and is associated with poor overall survival. <i>British Journal of Haematology</i> , 2013 , 161, 884-8	4.5	13
36	Interleukin-3-mediated regulation of Ecatenin in myeloid transformation and acute myeloid leukemia. <i>Journal of Leukocyte Biology</i> , 2014 , 96, 83-91	6.5	11

35	COVID-19 in Patients (pts) with Chronic Myeloid Leukemia (CML): Results from the International CML Foundation (iCMLF) CML and COVID-19 (CANDID) Study. <i>Blood</i> , 2020 , 136, 46-47	2.2	10
34	The preferential occurrence of FLT3-TKD mutations in inv(16) AML and impact on survival outcome: a combined analysis of 1053 core-binding factor AML patients. <i>British Journal of Haematology</i> , 2013 , 160, 557-9	4.5	9
33	A novel somatic JAK2 kinase-domain mutation in pediatric acute lymphoblastic leukemia with rapid on-treatment development of LOH. <i>Cancer Genetics</i> , 2017 , 216-217, 86-90	2.3	8
32	GATA2 is a New Predisposition Gene for Familial Myelodysplastic Syndrome (MDS) and Acute Myeloid Leukemia (AML). <i>Blood</i> , 2010 , 116, LBA-3-LBA-3	2.2	8
31	Increased peroxisome proliferator-activated receptor α activity reduces imatinib uptake and efficacy in chronic myeloid leukemia mononuclear cells. <i>Haematologica</i> , 2017 , 102, 843-853	6.6	7
30	A Method for Next-Generation Sequencing of Paired Diagnostic and Remission Samples to Detect Mitochondrial DNA Mutations Associated with Leukemia. <i>Journal of Molecular Diagnostics</i> , 2017 , 19, 711-721	5.1	7
29	miR-155 as a potential target of IL-3 signaling in primary AML cells. <i>Leukemia Research</i> , 2017 , 57, 57-59	2.7	6
28	p53-Dependent transcriptional responses to interleukin-3 signaling. <i>PLoS ONE</i> , 2012 , 7, e31428	3.7	6
27	Modelling ponatinib resistance in tyrosine kinase inhibitor-naïve and dasatinib resistant + cell lines. <i>Oncotarget</i> , 2018 , 9, 34735-34747	3.3	6
26	Integrated Bioinformatics Analysis Reveals Key Candidate Genes and Pathways Associated With Clinical Outcome in Hepatocellular Carcinoma. <i>Frontiers in Genetics</i> , 2020 , 11, 814	4.5	5
25	Genome-wide gene expression profiling identifies overlap with malignant adrenocortical tumours and novel mechanisms of inefficient steroidogenesis in familial ACTH-independent macronodular adrenal hyperplasia. <i>Endocrine-Related Cancer</i> , 2012 , 19, L19-23	5.7	4
24	The Natural History of NPM1MUT Measurable Residual Disease (MRD) Positivity after Completion of Chemotherapy in Acute Myeloid Leukemia (AML). <i>Blood</i> , 2020 , 136, 25-27	2.2	3
23	Polycomb Factor PHF19 Controls Cell Growth and Differentiation Toward Erythroid Pathway in Chronic Myeloid Leukemia Cells. <i>Frontiers in Cell and Developmental Biology</i> , 2021 , 9, 655201	5.7	3
22	Clinical impact of NPM1-mutant molecular persistence after chemotherapy for acute myeloid leukemia. <i>Blood Advances</i> , 2021 , 5, 5107-5111	7.8	2
21	Modeling Ponatinib Resistance in BCR-ABL1+ Cell Lines: Implications for Ponatinib Resistance in TKI-Resistant and TKI-naïve Patients. <i>Blood</i> , 2014 , 124, 4515-4515	2.2	1
20	Identification of an Epithelial-to-Mesenchymal Transition (EMT)-like Programme in t(4;14)-Positive Multiple Myeloma Reveals Novel Targets for Therapeutic Intervention. <i>Blood</i> , 2014 , 124, 647-647	2.2	1
19	High Prevalence of Relapse in Australian Children with Ph-like Acute Lymphoblastic Leukemia Despite Risk Adapted Treatment. <i>Blood</i> , 2015 , 126, 1419-1419	2.2	1
18	A 20 Gene Expression Signature That Predicts Early Molecular Response Failure in Chronic Phase CML Patients Treated with Frontline Imatinib. <i>Blood</i> , 2015 , 126, 596-596	2.2	1

17	In Vitro Modeling of Ph-like ALL Fusions Identifies Novel Kinase-Domain Mutations As Mode of TKI-Resistance - Implications for Targeted Therapy. <i>Blood</i> , 2016 , 128, 3957-3957	2.2	1
16	Integration of Multiple Bioassays Using Machine Learning to Identify High-Risk CP-CML Patients Treated with Frontline Imatinib. <i>Blood</i> , 2018 , 132, 1728-1728	2.2	1
15	Differential Lineage Involvement Between Very Low and Higher OCT-1 Activity Chronic-Phase CML Patients. <i>Blood</i> , 2011 , 118, 1675-1675	2.2	1
14	COVID-19 in Patients with Chronic Myeloid Leukemia: Poor Outcomes for Patients with Comorbidities, Older Age, Advanced Phase Disease, and Those from Low-Income Countries: An Update of the Candid Study. <i>Blood</i> , 2021 , 138, 634-634	2.2	0
13	Co-fuse: a new class discovery analysis tool to identify and prioritize recurrent fusion genes from RNA-sequencing data. <i>Molecular Genetics and Genomics</i> , 2018 , 293, 1217-1229	3.1	
12	Highly Sensitive Droplet Digital PCR to Identify CML Patients with a High Probability of Achieving Treatment-Free Remission. <i>Blood</i> , 2021 , 138, 2559-2559	2.2	
11	A Combination of CD302 gene Expression and 3-Months BCR-ABL1 Level Predicts Inferior Achievement of Deep Molecular Response in CP-CML Patients Treated with Imatinib. <i>Blood</i> , 2019 , 134, 663-663	2.2	
10	Whole Exome Sequencing of Acute Myeloid Leukaemia Patients Identifies Somatic and Germline Mutations in Fanconi Anaemia Genes. <i>Blood</i> , 2014 , 124, 698-698	2.2	
9	High Plasma Levels of TGF- β and IL-6 at Diagnosis Predict Early Molecular Response Failure and Transformation in CML. <i>Blood</i> , 2014 , 124, 1788-1788	2.2	
8	High Peroxisome Proliferator-Activated Receptor-Gamma (PPAR γ) Transcriptional Activity Reduces Active Influx of Imatinib and Kinase Inhibition in CML Cells. <i>Blood</i> , 2015 , 126, 2770-2770	2.2	
7	Non-Steroidal Anti-Inflammatory Drugs and Imatinib; Drug Interactions That May Impact Efficacy,. <i>Blood</i> , 2011 , 118, 3501-3501	2.2	
6	Methylation of a Single CpG in the GADD45A Proximal Promoter Is Associated with Poor Survival in Acute Myeloid Leukemia,. <i>Blood</i> , 2011 , 118, 3540-3540	2.2	
5	Global DNA Methylation Analysis Identifies Key Pathway Differences Between Poor (Low OCT-1 Activity) and Standard Risk CP-CML Patients At Diagnosis. <i>Blood</i> , 2012 , 120, 3730-3730	2.2	
4	Methylation of the Proximal Promoter of GADD45A Is Common in Acute Myeloid Leukemia and Is Associated with Poor Survival.. <i>Blood</i> , 2012 , 120, 2396-2396	2.2	
3	Role Of Peroxisome Proliferator-Activated Receptor Gamma (PPAR γ) and Its Ligands In The Regulation Of Functional OCT-1 Activity In CML Cells. <i>Blood</i> , 2013 , 122, 1470-1470	2.2	
2	CKLF and IL1B transcript levels at diagnosis are predictive of relapse in children with pre-B-cell acute lymphoblastic leukaemia. <i>British Journal of Haematology</i> , 2021 , 193, 171-175	4.5	
1	Highly sensitive droplet digital polymerase chain reaction for BCR :: ABL1 messenger RNA identifies patients with chronic myeloid leukaemia with a low probability of achieving treatment-free remission. <i>British Journal of Haematology</i> ,	4.5	