Jinjuan Yao

List of Publications by Year in descending order

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840776 752698 5,293 21 11 20 h-index citations g-index papers 22 22 22 10897 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Mutational landscape of metastatic cancer revealed from prospective clinical sequencing of 10,000 patients. Nature Medicine, 2017, 23, 703-713.	30.7	2,473
2	Memorial Sloan Kettering-Integrated Mutation Profiling of Actionable Cancer Targets (MSK-IMPACT). Journal of Molecular Diagnostics, 2015, 17, 251-264.	2.8	1,566
3	Genetic characterization of TET1, TET2, and TET3 alterations in myeloid malignancies. Blood, 2009, 114, 144-147.	1.4	661
4	Genetic Analysis of Transforming Events That Convert Chronic Myeloproliferative Neoplasms to Leukemias. Cancer Research, 2010, 70, 447-452.	0.9	279
5	Enhanced specificity of clinical high-sensitivity tumor mutation profiling in cell-free DNA via paired normal sequencing using MSK-ACCESS. Nature Communications, 2021, 12, 3770.	12.8	68
6	Recurrent somatic JAK3 mutations in NK-cell enteropathy. Blood, 2019, 134, 986-991.	1.4	42
7	Myeloid ELF1-like Factor Is a Potent Activator of Interleukin-8 Expression in Hematopoietic Cells. Journal of Biological Chemistry, 2004, 279, 6395-6400.	3.4	38
8	Depletion of Core Needle Biopsy Cellularity and DNA Content as a Result of Vigorous Touch Preparations. Archives of Pathology and Laboratory Medicine, 2015, 139, 907-912.	2.5	35
9	The ETS Protein MEF Is Regulated by Phosphorylation-Dependent Proteolysis via the Protein-Ubiquitin Ligase SCF Skp2. Molecular and Cellular Biology, 2006, 26, 3114-3123.	2.3	34
10	JAK2, PD-L1, and PD-L2 (9p24.1) amplification in metastatic mucosal and cutaneous melanomas with durable response to immunotherapy. Human Pathology, 2019, 88, 87-91.	2.0	20
11	Routine Evaluation of Minimal Residual Disease in Myeloma Using Next-Generation Sequencing Clonality Testing. Journal of Molecular Diagnostics, 2021, 23, 181-199.	2.8	19
12	Novel patient-derived models of desmoplastic small round cell tumor confirm a targetable dependency on ERBB signaling. DMM Disease Models and Mechanisms, 2022, 15, .	2.4	11
13	Pan-Cancer Biomarkers: Changing the Landscape of Molecular Testing. Archives of Pathology and Laboratory Medicine, 2021, 145, 692-698.	2.5	10
14	Divergent clonal evolution of a common precursor to mantle cell lymphoma and classic Hodgkin lymphoma. Journal of Physical Education and Sports Management, 2019, 5, a004259.	1.2	8
15	Myeloid/lymphoid neoplasms with eosinophilia/ basophilia and ETV6-ABL1 fusion: cell-of-origin and response to tyrosine kinase inhibition. Haematologica, 2021, 106, 614-618.	3 . 5	8
16	A case of acute myeloid leukemia with e6a2 BCR-ABL fusion transcript acquired after progressing from chronic myelomonocytic leukemia. Leukemia Research Reports, 2017, 7, 17-19.	0.4	7
17	A case ofKMT2A–SEPT9fusion–associated acute megakaryoblastic leukemia. Journal of Physical Education and Sports Management, 2018, 4, a003426.	1.2	5
18	Rare and novel <scp><i>RUNX1</i></scp> fusions in myeloid neoplasms: A singleâ€institute experience. Genes Chromosomes and Cancer, 2021, 60, 100-107.	2.8	3

#	Article	IF	CITATIONS
19	A Brief Overview and Update on Major Molecular Genomic Alterations in Solid, Bone and Soft Tissue Tumors, and Hematopoietic As Well As Lymphoid Malignancies. Archives of Pathology and Laboratory Medicine, 2021, 145, 1358-1366.	2.5	2
20	Quantitative Off-Target Detection of Epstein-Barr Virus–Derived DNA in Routine Molecular Profiling of Hematopoietic Neoplasms by Panel-Based Hybrid-Capture Next-Generation Sequencing. Journal of Molecular Diagnostics, 2021, , .	2.8	2
21	Clonally-Related CD5+ CLL/SLL and CD10+ high grade B-cell lymphoma suggests common neoplastic progenitor with branched disease evolution, with therapeutic implications. Leukemia and Lymphoma, 2020, 61, 460-464.	1.3	0