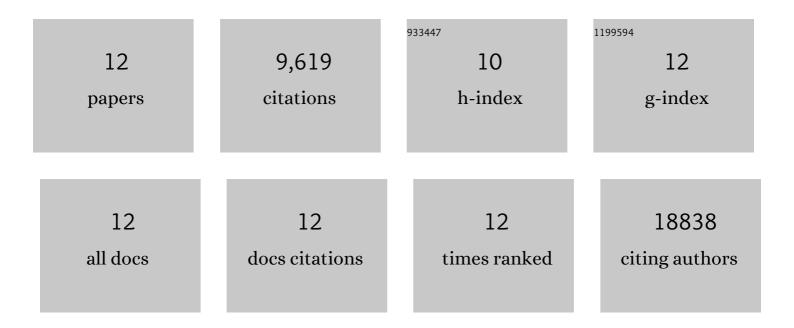
Hongxin Zhang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8253417/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Genome Nexus: A Comprehensive Resource for the Annotation and Interpretation of Genomic Variants in Cancer. JCO Clinical Cancer Informatics, 2022, 6, e2100144.	2.1	4
2	Base editing sensor libraries for high-throughput engineering and functional analysis of cancer-associated single nucleotide variants. Nature Biotechnology, 2022, 40, 862-873.	17.5	44
3	OncoTree: A Cancer Classification System for Precision Oncology. JCO Clinical Cancer Informatics, 2021, 5, 221-230.	2.1	51
4	Tumor fraction-guided cell-free DNA profiling in metastatic solid tumor patients. Genome Medicine, 2021, 13, 96.	8.2	26
5	Therapeutic Implications of Germline Testing in Patients With Advanced Cancers. Journal of Clinical Oncology, 2021, 39, 2698-2709.	1.6	83
6	Annotation of Somatic Genomic Variants in Hematologic Diseases Using OncoKB, a Precision Oncology Knowledgebase. Blood, 2019, 134, 2148-2148.	1.4	3
7	Comprehensive Characterization of Cancer Driver Genes and Mutations. Cell, 2018, 173, 371-385.e18.	28.9	1,670
8	Cell-of-Origin Patterns Dominate the Molecular Classification of 10,000 Tumors from 33 Types of Cancer. Cell, 2018, 173, 291-304.e6.	28.9	1,718
9	Oncogenic Signaling Pathways in The Cancer Genome Atlas. Cell, 2018, 173, 321-337.e10.	28.9	2,111
10	3D clusters of somatic mutations in cancer reveal numerous rare mutations as functional targets. Genome Medicine, 2017, 9, 4.	8.2	170
11	Mutational landscape of metastatic cancer revealed from prospective clinical sequencing of 10,000 patients. Nature Medicine, 2017, 23, 703-713.	30.7	2,473
12	OncoKB: A Precision Oncology Knowledge Base. JCO Precision Oncology, 2017, 2017, 1-16.	3.0	1,266