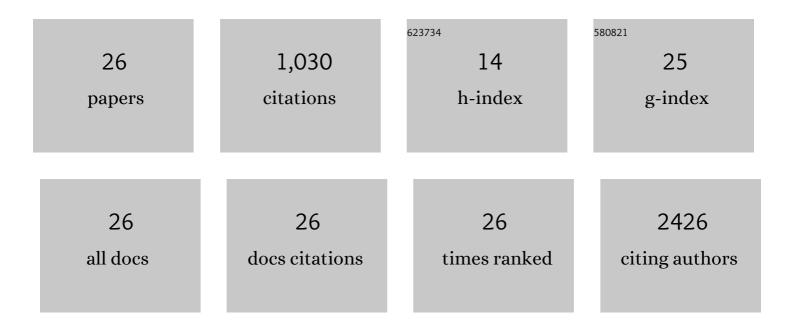
Hannah C Cheung

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4827015/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Multi-site desmoplastic small round cell tumors are genetically related and immune-cold. Npj Precision Oncology, 2022, 6, 21.	5.4	7
2	The androgen receptor is a therapeutic target in desmoplastic small round cell sarcoma. Nature Communications, 2022, 13, .	12.8	14
3	Germline DNMT3A mutation in familial acute myeloid leukaemia. Epigenetics, 2021, 16, 567-576.	2.7	9
4	Metabolic compensation activates pro-survival mTORC1 signaling upon 3-phosphoglycerate dehydrogenase inhibition in osteosarcoma. Cell Reports, 2021, 34, 108678.	6.4	33
5	Divergent clonal evolution of blastic plasmacytoid dendritic cell neoplasm and chronic myelomonocytic leukemia from a shared TET2-mutated origin. Leukemia, 2021, 35, 3299-3303.	7.2	18
6	Enhancer reprogramming in PRC2-deficient malignant peripheral nerve sheath tumors induces a targetable de-differentiated state. Acta Neuropathologica, 2021, 142, 565-590.	7.7	12
7	Bâ€cell lymphoma/leukaemia 11B (BCL11B) expression status helps distinguish early Tâ€cell precursor acute lymphoblastic leukaemia/lymphoma (ETPâ€ALL/LBL) from other subtypes of Tâ€cell ALL/LBL. British Journal of Haematology, 2021, 194, 1034-1038.	2.5	9
8	Single cell T cell landscape and T cell receptor repertoire profiling of AML in context of PD-1 blockade therapy. Nature Communications, 2021, 12, 6071.	12.8	44
9	T(6;14)(q25;q32) involves BCL11B and is highly associated with mixed-phenotype acute leukemia, T/myeloid. Leukemia, 2020, 34, 2509-2512.	7.2	14
10	Immuno-genomic landscape of osteosarcoma. Nature Communications, 2020, 11, 1008.	12.8	143
11	Features of non-activation dendritic state and immune deficiency in blastic plasmacytoid dendritic cell neoplasm (BPDCN). Blood Cancer Journal, 2019, 9, 99.	6.2	26
12	Genomic profiling of dedifferentiated liposarcoma compared to matched well-differentiated liposarcoma reveals higher genomic complexity and a common origin. Journal of Physical Education and Sports Management, 2018, 4, a002386.	1.2	45
13	The SS18-SSX Fusion Oncoprotein Hijacks BAF Complex Targeting and Function to Drive Synovial Sarcoma. Cancer Cell, 2018, 33, 1128-1141.e7.	16.8	169
14	KMT2D/MLL2 inactivation is associated with recurrence in adult-type granulosa cell tumors of the ovary. Nature Communications, 2018, 9, 2496.	12.8	39
15	FusionPathway: Prediction of pathways and therapeutic targets associated with gene fusions in cancer. PLoS Computational Biology, 2018, 14, e1006266.	3.2	8
16	Targeted next generation sequencing of well-differentiated/dedifferentiated liposarcoma reveals novel gene amplifications and mutations. Oncotarget, 2018, 9, 19891-19899.	1.8	28
17	Genomic and immune heterogeneity are associated with differential responses to therapy in melanoma. Npj Genomic Medicine, 2017, 2, .	3.8	120
18	Establishment and characterization of a new human myxoid liposarcoma cell line (DL-221) with the FUS-DDIT3 translocation. Laboratory Investigation, 2016, 96, 885-894.	3.7	17

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#	Article	IF	CITATIONS
19	Implementation of a Pan-Genomic Approach to Investigate Holobiont-Infecting Microbe Interaction: A Case Report of a Leukemic Patient with Invasive Mucormycosis. PLoS ONE, 2015, 10, e0139851.	2.5	47
20	Abstract LB-222: Long-term subclonal evolution of CLL from immune selective pressure after allogeneic stem cell transplant and donor lymphocyte infusion. , 2015, , .		0
21	An S/T-Q cluster domain census unveils new putative targets under Tel1/Mec1 control. BMC Genomics, 2012, 13, 664.	2.8	19
22	PTBP1â€dependent regulation of USP5 alternative RNA splicing plays a role in glioblastoma tumorigenesis. Molecular Carcinogenesis, 2012, 51, 895-906.	2.7	75
23	Constitutional tandem duplication of 9q34 that truncates <i>EHMT1</i> in a child with ganglioglioma. Pediatric Blood and Cancer, 2012, 58, 801-805.	1.5	8
24	Identification of genetic susceptibility to childhood cancer through analysis of genes in parallel. Cancer Genetics, 2011, 204, 19-25.	0.4	14
25	Global analysis of aberrant pre-mRNA splicing in glioblastoma using exon expression arrays. BMC Genomics, 2008, 9, 216.	2.8	66
26	Polypyrimidine tract binding protein and Notch1 are independently re-expressed in glioma. Modern Pathology, 2006, 19, 1034-1041.	5.5	46