

Hannah C Cheung

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

1,030
citations

623734

14
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580821

25
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all docs

26
docs citations

26
times ranked

2426
citing authors

#	ARTICLE	IF	CITATIONS
1	Multi-site desmoplastic small round cell tumors are genetically related and immune-cold. <i>Npj Precision Oncology</i> , 2022, 6, 21.	5.4	7
2	The androgen receptor is a therapeutic target in desmoplastic small round cell sarcoma. <i>Nature Communications</i> , 2022, 13, .	12.8	14
3	Germline DNMT3A mutation in familial acute myeloid leukaemia. <i>Epigenetics</i> , 2021, 16, 567-576.	2.7	9
4	Metabolic compensation activates pro-survival mTORC1 signaling upon 3-phosphoglycerate dehydrogenase inhibition in osteosarcoma. <i>Cell Reports</i> , 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. <i>Leukemia</i> , 2021, 35, 3299-3303.	7.2	18
6	Enhancer reprogramming in PRC2-deficient malignant peripheral nerve sheath tumors induces a targetable de-differentiated state. <i>Acta Neuropathologica</i> , 2021, 142, 565-590.	7.7	12
7	B α cell lymphoma/leukaemia 11B (BCL11B) expression status helps distinguish early T α cell precursor acute lymphoblastic leukaemia/lymphoma (ETP α ALL/LBL) from other subtypes of T α cell ALL/LBL. <i>British Journal of Haematology</i> , 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. <i>Nature Communications</i> , 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. <i>Leukemia</i> , 2020, 34, 2509-2512.	7.2	14
10	Immuno-genomic landscape of osteosarcoma. <i>Nature Communications</i> , 2020, 11, 1008.	12.8	143
11	Features of non-activation dendritic state and immune deficiency in blastic plasmacytoid dendritic cell neoplasm (BPDCN). <i>Blood Cancer Journal</i> , 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. <i>Journal of Physical Education and Sports Management</i> , 2018, 4, a002386.	1.2	45
13	The SS18-SSX Fusion Oncoprotein Hijacks BAF Complex Targeting and Function to Drive Synovial Sarcoma. <i>Cancer Cell</i> , 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. <i>Nature Communications</i> , 2018, 9, 2496.	12.8	39
15	FusionPathway: Prediction of pathways and therapeutic targets associated with gene fusions in cancer. <i>PLoS Computational Biology</i> , 2018, 14, e1006266.	3.2	8
16	Targeted next generation sequencing of well-differentiated/dedifferentiated liposarcoma reveals novel gene amplifications and mutations. <i>Oncotarget</i> , 2018, 9, 19891-19899.	1.8	28
17	Genomic and immune heterogeneity are associated with differential responses to therapy in melanoma. <i>Npj Genomic Medicine</i> , 2017, 2, .	3.8	120
18	Establishment and characterization of a new human myxoid liposarcoma cell line (DL-221) with the FUS-DDIT3 translocation. <i>Laboratory Investigation</i> , 2016, 96, 885-894.	3.7	17

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19	Implementation of a Pan-Genomic Approach to Investigate Holobiont-Infected 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	PTBP1-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