

# Hai Yan

## List of Publications by Year in descending order

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Version: 2024-02-01

44  
papers

8,748  
citations

279487  
23  
h-index

301761  
39  
g-index

44  
all docs

44  
docs citations

44  
times ranked

13819  
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>IDH1</i> and <i>IDH2</i> Mutations in Gliomas. <i>New England Journal of Medicine</i> , 2009, 360, 765-773.	13.9	5,285
2	<i>TERT</i> promoter mutations occur frequently in gliomas and a subset of tumors derived from cells with low rates of self-renewal. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 6021-6026.	3.3	1,202
3	Isocitrate dehydrogenase mutations in gliomas. <i>Neuro-Oncology</i> , 2016, 18, 16-26.	0.6	221
4	The implications of IDH mutations for cancer development and therapy. <i>Nature Reviews Clinical Oncology</i> , 2021, 18, 645-661.	12.5	155
5	Recurrent TERT promoter mutations identified in a large-scale study of multiple tumour types are associated with increased TERT expression and telomerase activation. <i>European Journal of Cancer</i> , 2015, 51, 969-976.	1.3	150
6	Exome sequencing identifies somatic gain-of-function PPM1D mutations in brainstem gliomas. <i>Nature Genetics</i> , 2014, 46, 726-730.	9.4	148
7	The genome-wide mutational landscape of pituitary adenomas. <i>Cell Research</i> , 2016, 26, 1255-1259.	5.7	137
8	Mutant Metabolic Enzymes Are at the Origin of Gliomas. <i>Cancer Research</i> , 2009, 69, 9157-9159.	0.4	132
9	DNA hypermethylation within TERT promoter upregulates TERT expression in cancer. <i>Journal of Clinical Investigation</i> , 2018, 129, 223-229.	3.9	130
10	Detection of early-stage hepatocellular carcinoma in asymptomatic HBsAg-seropositive individuals by liquid biopsy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 6308-6312.	3.3	127
11	The genomic landscape of TERT promoter wildtype-IDH wildtype glioblastoma. <i>Nature Communications</i> , 2018, 9, 2087.	5.8	124
12	Molecular profiling of tumors of the brainstem by sequencing of CSF-derived circulating tumor DNA. <i>Acta Neuropathologica</i> , 2019, 137, 297-306.	3.9	109
13	Identification of recurrent USP48 and BRAF mutations in Cushing's disease. <i>Nature Communications</i> , 2018, 9, 3171.	5.8	106
14	The H3.3 K27M mutation results in a poorer prognosis in brainstem gliomas than thalamic gliomas in adults. <i>Human Pathology</i> , 2015, 46, 1626-1632.	1.1	88
15	Germline Mutations in CDH23, Encoding Cadherin-Related 23, Are Associated with Both Familial and Sporadic Pituitary Adenomas. <i>American Journal of Human Genetics</i> , 2017, 100, 817-823.	2.6	57
16	Clonality analysis of multifocal papillary thyroid carcinoma by using genetic profiles. <i>Journal of Pathology</i> , 2016, 239, 72-83.	2.1	56
17	Genome-Wide CRISPR-Cas9 Screen Reveals Selective Vulnerability of <i>ATR</i> -Mutant Cancers to WEE1 Inhibition. <i>Cancer Research</i> , 2020, 80, 510-523.	0.4	52
18	Allelic variations in gene expression. <i>Current Opinion in Oncology</i> , 2004, 16, 39-43.	1.1	50

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19	The integrated genomic and epigenomic landscape of brainstem glioma. <i>Nature Communications</i> , 2020, 11, 3077.	5.8	50
20	<i>Cic</i> Loss Promotes Gliomagenesis via Aberrant Neural Stem Cell Proliferation and Differentiation. <i>Cancer Research</i> , 2017, 77, 6097-6108.	0.4	46
21	A machine learning-based prediction model of H3K27M mutations in brainstem gliomas using conventional MRI and clinical features. <i>Radiotherapy and Oncology</i> , 2019, 130, 172-179.	0.3	42
22	Mutant IDH1 Disrupts the Mouse Subventricular Zone and Alters Brain Tumor Progression. <i>Molecular Cancer Research</i> , 2017, 15, 507-520.	1.5	41
23	TERT promoter mutations contribute to IDH mutations in predicting differential responses to adjuvant therapies in WHO grade II and III diffuse gliomas. <i>Oncotarget</i> , 2015, 6, 24871-24883.	0.8	34
24	MTAP Loss Promotes Stemness in Glioblastoma and Confers Unique Susceptibility to Purine Starvation. <i>Cancer Research</i> , 2019, 79, 3383-3394.	0.4	30
25	A PRMT5-RNF168-SMURF2 Axis Controls H2AX Proteostasis. <i>Cell Reports</i> , 2019, 28, 3199-3211.e5.	2.9	27
26	Sensitive and rapid detection of <i>TERT</i> promoter and <i>IDH</i> mutations in diffuse gliomas. <i>Neuro-Oncology</i> , 2019, 21, 440-450.	0.6	27
27	Patient-derived DIPG cells preserve stem-like characteristics and generate orthotopic tumors. <i>Oncotarget</i> , 2017, 8, 76644-76655.	0.8	27
28	Synthesis and evaluation of radiolabeled AGI-5198 analogues as candidate radiotracers for imaging mutant IDH1 expression in tumors. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2018, 28, 694-699.	1.0	18
29	Synthesis and Evaluation of a <sup>18</sup> F-Labeled Triazinediamine Analogue for Imaging Mutant IDH1 Expression in Gliomas by PET. <i>ACS Medicinal Chemistry Letters</i> , 2018, 9, 606-611.	1.3	17
30	Radiolabeled inhibitors as probes for imaging mutant IDH1 expression in gliomas: Synthesis and preliminary evaluation of labeled butyl-phenyl sulfonamide analogs. <i>European Journal of Medicinal Chemistry</i> , 2016, 119, 218-230.	2.6	13
31	Dual role of allele-specific DNA hypermethylation within the TERT promoter in cancer. <i>Journal of Clinical Investigation</i> , 2021, 131, .	3.9	11
32	Functional requirement of a wild-type allele for mutant IDH1 to suppress anchorage-independent growth through redox homeostasis. <i>Acta Neuropathologica</i> , 2018, 135, 285-298.	3.9	10
33	Chromatin Accessibility Mapping Identifies Mediators of Basal Transcription and Retinoid-Induced Repression of OTX2 in Medulloblastoma. <i>PLoS ONE</i> , 2014, 9, e107156.	1.1	8
34	Mutant allele quantification reveals a genetic basis for TP53 mutation-driven castration resistance in prostate cancer cells. <i>Scientific Reports</i> , 2018, 8, 12507.	1.6	5
35	ATIM-27. TUMOR MUTATIONAL BURDEN PREDICTS RESPONSE TO ONCOLYTIC POLIO/RHINOVIRUS RECOMBINANT (PVSRIPO) IN MALIGNANT GLIOMA PATIENTS: ASSESSMENT OF TRANSCRIPTIONAL AND IMMUNOLOGICAL CORRELATES. <i>Neuro-Oncology</i> , 2019, 21, vi7-vi7.	0.6	5
36	Hereditary brain tumor with a homozygous germline mutation in PMS2: pedigree analysis and prenatal screening in a family with constitutional mismatch repair deficiency (CMMRD) syndrome. <i>Familial Cancer</i> , 2019, 18, 261-265.	0.9	3

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37	Improved grading of IDH-mutated astrocytic gliomas. <i>Nature Reviews Neurology</i> , 2018, 14, 383-384.	4.9	2
38	Digital karyotyping: a powerful tool for cancer gene discovery. , 2006, , .		1
39	IMMU-34. ATRX MUTATIONS PREDICT RESPONSE TO INNATE BASED THERAPY IN GLIOMA. <i>Neuro-Oncology</i> , 2019, 21, vi126-vi126.	0.6	1
40	Hitting Gliomas When They Are Down: Exploiting IDH-Mutant Metabolic Vulnerabilities. <i>Cancer Discovery</i> , 2020, 10, 1629-1631.	7.7	1
41	GENE-42. THE GENOMIC LANDSCAPE OF TRIPLE-NEGATIVE GLIOBLASTOMA. <i>Neuro-Oncology</i> , 2018, 20, vi112-vi112.	0.6	0
42	TMOD-33. ESTABLISHMENT AND PRELIMINARY EVALUATION OF BEVACIZUMAB-RESISTANT GLIOMA XENOGRAFT MODELS. <i>Neuro-Oncology</i> , 2018, 20, vi275-vi275.	0.6	0
43	GENE-01. THE GENOMIC LANDSCAPE OF TRIPLE-NEGATIVE GLIOBLASTOMA. <i>Neuro-Oncology</i> , 2018, 20, vi102-vi103.	0.6	0
44	ATIM-31. SAFETY OF TUMOR-SPECIFIC PEPTIDE VACCINE TARGETING ISOCITRATE DEHYDROGENASE 1 MUTATION IN RECURRENT RESECTABLE LOW GRADE GLIOMA PATIENTS. <i>Neuro-Oncology</i> , 2019, 21, vi8-vi8.	0.6	0