

# Chen Ding

## List of Publications by Year in descending order

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Version: 2024-02-01

49  
papers

2,696  
citations

279798

23  
h-index

197818

49  
g-index

52  
all docs

52  
docs citations

52  
times ranked

4934  
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteogenomic characterization of cholangiocarcinoma. <i>Hepatology</i> , 2023, 77, 411-429.	7.3	12
2	TRIM21 regulates pyroptotic cell death by promoting Gasdermin D oligomerization. <i>Cell Death and Differentiation</i> , 2022, 29, 439-450.	11.2	33
3	Gut microbiome alterations and gut barrier dysfunction are associated with host immune homeostasis in COVID-19 patients. <i>BMC Medicine</i> , 2022, 20, 24.	5.5	83
4	Integrative proteomic characterization of trace FFPE samples in early-stage gastrointestinal cancer. <i>Proteome Science</i> , 2022, 20, 5.	1.7	2
5	A proteogenomic analysis of clear cell renal cell carcinoma in a Chinese population. <i>Nature Communications</i> , 2022, 13, 2052.	12.8	48
6	Proteomic analysis reveals key differences between squamous cell carcinomas and adenocarcinomas across multiple tissues. <i>Nature Communications</i> , 2022, 13, .	12.8	11
7	Tumor-derived neomorphic mutations in ASXL1 impairs the BAP1-ASXL1-FOXK1/K2 transcription network. <i>Protein and Cell</i> , 2021, 12, 557-577.	11.0	14
8	Histone H3K27 methyltransferase EZH2 and demethylase JMJD3 regulate hepatic stellate cells activation and liver fibrosis. <i>Theranostics</i> , 2021, 11, 361-378.	10.0	48
9	Deep Dive on the Proteome of Human Body Fluids: A Valuable Data Resource for Biomarker Discovery. <i>Cancer Genomics and Proteomics</i> , 2021, 18, 549-568.	2.0	10
10	Overexpressed NEDD8 as a potential therapeutic target in esophageal squamous cell carcinoma. <i>Cancer Biology and Medicine</i> , 2021, 19, 504-517.	3.0	3
11	Polypeptide <i>N</i> -acetylgalactosaminyltransferase 18 retains in endoplasmic reticulum depending on its luminal regions interacting with ER resident UGGT1, PLOD3 and LPCAT1. <i>Glycobiology</i> , 2021, 31, 947-958.	2.5	3
12	Demethylation of the <i>SFRP4</i> Promoter Drives Gastric Cancer Progression via the Wnt Pathway. <i>Molecular Cancer Research</i> , 2021, 19, 1454-1464.	3.4	9
13	E3 ligase TRIM25 ubiquitinates RIP3 to inhibit TNF induced cell necrosis. <i>Cell Death and Differentiation</i> , 2021, 28, 2888-2899.	11.2	25
14	Establishment of an Immune Cell Infiltration Score to Help Predict the Prognosis and Chemotherapy Responsiveness of Gastric Cancer Patients. <i>Frontiers in Oncology</i> , 2021, 11, 650673.	2.8	26
15	Proteome-Wide Profiling of Readers for DNA Modification. <i>Advanced Science</i> , 2021, 8, e2101426.	11.2	7
16	METTL3-mediated m6A RNA methylation promotes the anti-tumour immunity of natural killer cells. <i>Nature Communications</i> , 2021, 12, 5522.	12.8	96
17	Proteome-wide profiling of transcriptional machinery on accessible chromatin with biotinylated transposons. <i>Science Advances</i> , 2021, 7, eabh1022.	10.3	3
18	Protumorigenic Role of Elevated Levels of DNA Polymerase Epsilon Predicts an Immune-Suppressive Microenvironment in Clear Cell Renal Cell Carcinoma. <i>Frontiers in Genetics</i> , 2021, 12, 751977.	2.3	6

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19	Dosage effect of multiple genes accounts for multisystem disorder of myotonic dystrophy type 1. <i>Cell Research</i> , 2020, 30, 133-145.	12.0	21
20	Combinations of proteasome inhibitors with obatoclox are effective for small cell lung cancer. <i>Acta Pharmacologica Sinica</i> , 2020, 42, 1298-1310.	6.1	5
21	Blood molecular markers associated with COVID-19 immunopathology and multi-organ damage. <i>EMBO Journal</i> , 2020, 39, e105896.	7.8	123
22	Myeloid PTEN promotes chemotherapy-induced NLRP3-inflammasome activation and antitumour immunity. <i>Nature Cell Biology</i> , 2020, 22, 716-727.	10.3	70
23	Time- and Dose-Resolved Proteome of PM2.5-Exposure-Induced Lung Injury and Repair in Rats. <i>Journal of Proteome Research</i> , 2020, 19, 3162-3175.	3.7	8
24	A time-resolved multi-omic atlas of the developing mouse liver. <i>Genome Research</i> , 2020, 30, 263-275.	5.5	22
25	Proteomic Investigations of Transcription Factors Critical in Geniposide-Mediated Suppression of Alcoholic Steatosis and in Overdose-Induced Hepatotoxicity on Liver in Rats. <i>Journal of Proteome Research</i> , 2019, 18, 3821-3830.	3.7	10
26	Phosphoproteomics Enables Molecular Subtyping and Nomination of Kinase Candidates for Individual Patients of Diffuse-Type Gastric Cancer. <i>IScience</i> , 2019, 22, 44-57.	4.1	16
27	Proteomics and Precision Medicine. <i>Small Methods</i> , 2019, 3, 1900075.	8.6	5
28	Transketolase (TKT) activity and nuclear localization promote hepatocellular carcinoma in a metabolic and a non-metabolic manner. <i>Journal of Experimental and Clinical Cancer Research</i> , 2019, 38, 154.	8.6	54
29	An alternatively transcribed <i>TAZ</i> variant negatively regulates <i>JAK</i> - <i>STAT</i> signaling. <i>EMBO Reports</i> , 2019, 20, .	4.5	14
30	Allele-selective lowering of mutant HTT protein by HTT-LC3 linker compounds. <i>Nature</i> , 2019, 575, 203-209.	27.8	288
31	A region-resolved mucosa proteome of the human stomach. <i>Nature Communications</i> , 2019, 10, 39.	12.8	43
32	OTUB2 Promotes Cancer Metastasis via Hippo-Independent Activation of YAP and TAZ. <i>Molecular Cell</i> , 2019, 73, 7-21.e7.	9.7	112
33	A proteomic landscape of diffuse-type gastric cancer. <i>Nature Communications</i> , 2018, 9, 1012.	12.8	175
34	A proteomics landscape of circadian clock in mouse liver. <i>Nature Communications</i> , 2018, 9, 1553.	12.8	115
35	3' UTR lengthening as a novel mechanism in regulating cellular senescence. <i>Genome Research</i> , 2018, 28, 285-294.	5.5	90
36	A time-resolved multi-omic atlas of the developing mouse stomach. <i>Nature Communications</i> , 2018, 9, 4910.	12.8	31

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37	Differential proteomics profiling identifies LDPs and biological functions in high-fat diet-induced fatty livers. <i>Journal of Lipid Research</i> , 2017, 58, 681-694.	4.2	20
38	Firmiana: towards a one-stop proteomic cloud platform for data processing and analysis. <i>Nature Biotechnology</i> , 2017, 35, 409-412.	17.5	80
39	Proteome-wide Mapping of Endogenous SUMOylation Sites in Mouse Testis. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 717-727.	3.8	15
40	TRIM65-catalyzed ubiquitination is essential for MDA5-mediated antiviral innate immunity. <i>Journal of Experimental Medicine</i> , 2017, 214, 459-473.	8.5	120
41	An integrated approach to identify critical transcription factors in the protection against hydrogen peroxide-induced oxidative stress by Danhong injection. <i>Free Radical Biology and Medicine</i> , 2017, 112, 480-493.	2.9	17
42	Multidimensional Proteomics Reveals a Role of UHRF2 in the Regulation of Epithelial-Mesenchymal Transition (EMT). <i>Molecular and Cellular Proteomics</i> , 2016, 15, 2263-2278.	3.8	26
43	Destabilization of Fatty Acid Synthase by Acetylation Inhibits <i>De Novo</i> Lipogenesis and Tumor Cell Growth. <i>Cancer Research</i> , 2016, 76, 6924-6936.	0.9	92
44	Transcription Factor Response Elements on Tip: A Sensitive Approach for Large-Scale Endogenous Transcription Factor Quantitative Identification. <i>Analytical Chemistry</i> , 2016, 88, 11990-11994.	6.5	8
45	A Cell-type-resolved Liver Proteome. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 3190-3202.	3.8	83
46	Proteome-wide profiling of activated transcription factors with a concatenated tandem array of transcription factor response elements. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 6771-6776.	7.1	91
47	A Fast Workflow for Identification and Quantification of Proteomes. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 2370-2380.	3.8	94
48	Analysis of the Human Endogenous Coregulator Complexome. <i>Cell</i> , 2011, 145, 787-799.	28.9	383
49	Hepatocytes proteomic alteration and seroproteome analysis of HBV-transgenic mice. <i>Proteomics</i> , 2009, 9, 87-105.	2.2	16