

Hu Zhou

List of Publications by Citations

Source: <https://exaly.com/author-pdf/960065/hu-zhou-publications-by-citations.pdf>

Version: 2024-04-27

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

124
papers

4,623
citations

36
h-index

64
g-index

130
ext. papers

6,198
ext. citations

8.9
avg, IF

5.19
L-index

#	Paper	IF	Citations
124	Meclofenamic acid selectively inhibits FTO demethylation of m6A over ALKBH5. <i>Nucleic Acids Research</i> , 2015 , 43, 373-84	20.1	278
123	Small-Molecule Targeting of Oncogenic FTO Demethylase in Acute Myeloid Leukemia. <i>Cancer Cell</i> , 2019 , 35, 677-691.e10	24.3	239
122	Integrated Proteogenomic Characterization of HBV-Related Hepatocellular Carcinoma. <i>Cell</i> , 2019 , 179, 561-577.e22	56.2	232
121	Long noncoding RNA LINC00336 inhibits ferroptosis in lung cancer by functioning as a competing endogenous RNA. <i>Cell Death and Differentiation</i> , 2019 , 26, 2329-2343	12.7	204
120	Altered intestinal microbiota-host mitochondria crosstalk in new onset Crohn's disease. <i>Nature Communications</i> , 2016 , 7, 13419	17.4	189
119	YTHDF1 links hypoxia adaptation and non-small cell lung cancer progression. <i>Nature Communications</i> , 2019 , 10, 4892	17.4	140
118	Lipidomics era: accomplishments and challenges. <i>Mass Spectrometry Reviews</i> , 2010 , 29, 877-929	11	139
117	Multi-omic measurements of heterogeneity in HeLa cells across laboratories. <i>Nature Biotechnology</i> , 2019 , 37, 314-322	44.5	129
116	pFind: a novel database-searching software system for automated peptide and protein identification via tandem mass spectrometry. <i>Bioinformatics</i> , 2005 , 21, 3049-50	7.2	116
115	Acetylation of PGK1 promotes liver cancer cell proliferation and tumorigenesis. <i>Hepatology</i> , 2017 , 65, 515-528	11.2	113
114	EGLN1/c-Myc Induced Lymphoid-Specific Helicase Inhibits Ferroptosis through Lipid Metabolic Gene Expression Changes. <i>Theranostics</i> , 2017 , 7, 3293-3305	12.1	106
113	Proteomic analysis of hepatitis B virus-associated hepatocellular carcinoma: Identification of potential tumor markers. <i>Proteomics</i> , 2005 , 5, 1125-39	4.8	103
112	ERK kinase phosphorylates and destabilizes the tumor suppressor FBW7 in pancreatic cancer. <i>Cell Research</i> , 2015 , 25, 561-73	24.7	88
111	Conformational states of the full-length glucagon receptor. <i>Nature Communications</i> , 2015 , 6, 7859	17.4	86
110	Characterization of the 3a protein of SARS-associated coronavirus in infected vero E6 cells and SARS patients. <i>Journal of Molecular Biology</i> , 2004 , 341, 271-9	6.5	83
109	Missense mutation in APOC3 within the C-terminal lipid binding domain of human ApoC-III results in impaired assembly and secretion of triacylglycerol-rich very low density lipoproteins: evidence that ApoC-III plays a major role in the formation of lipid precursors within the microsomal lumen. <i>Journal of Biological Chemistry</i> , 2011 , 286, 27769-80	5.4	75
108	A complex structure of arrestin-2 bound to a G protein-coupled receptor. <i>Cell Research</i> , 2019 , 29, 971-983	24.7	71

107	Quantitative analysis of severe acute respiratory syndrome (SARS)-associated coronavirus-infected cells using proteomic approaches: implications for cellular responses to virus infection. <i>Molecular and Cellular Proteomics</i> , 2005 , 4, 902-13	7.6	69
106	Proteomic analysis with integrated multiple dimensional liquid chromatography/mass spectrometry based on elution of ion exchange column using pH steps. <i>Analytical Chemistry</i> , 2005 , 77, 5793-9	7.8	68
105	Advancements in top-down proteomics. <i>Analytical Chemistry</i> , 2012 , 84, 720-34	7.8	67
104	A redox mechanism underlying nucleolar stress sensing by nucleophosmin. <i>Nature Communications</i> , 2016 , 7, 13599	17.4	66
103	A high-throughput approach for subcellular proteome: identification of rat liver proteins using subcellular fractionation coupled with two-dimensional liquid chromatography tandem mass spectrometry and bioinformatic analysis. <i>Molecular and Cellular Proteomics</i> , 2004 , 3, 441-55	7.6	66
102	Phosphoproteome analysis of mouse liver using immobilized metal affinity purification and linear ion trap mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2004 , 18, 2169-76	2.2	61
101	Ruthenium-Promoted C-H Activation Reactions between DNA-Conjugated Acrylamide and Aromatic Acids. <i>Organic Letters</i> , 2018 , 20, 4764-4768	6.2	55
100	Localized-statistical quantification of human serum proteome associated with type 2 diabetes. <i>PLoS ONE</i> , 2008 , 3, e3224	3.7	54
99	Nonsynonymous mutations within APOB in human familial hypobetalipoproteinemia: evidence for feedback inhibition of lipogenesis and postendoplasmic reticulum degradation of apolipoprotein B. <i>Journal of Biological Chemistry</i> , 2010 , 285, 6453-64	5.4	52
98	The differential protein and lipid compositions of noncaveolar lipid microdomains and caveolae. <i>Cell Research</i> , 2009 , 19, 497-506	24.7	52
97	Rare cell proteomic reactor applied to stable isotope labeling by amino acids in cell culture (SILAC)-based quantitative proteomics study of human embryonic stem cell differentiation. <i>Molecular and Cellular Proteomics</i> , 2011 , 10, M110.000679	7.6	50
96	The mTOR-S6K pathway links growth signalling to DNA damage response by targeting RNF168. <i>Nature Cell Biology</i> , 2018 , 20, 320-331	23.4	48
95	Heterogeneous immunogenomic features and distinct escape mechanisms in multifocal hepatocellular carcinoma. <i>Journal of Hepatology</i> , 2020 , 72, 896-908	13.4	48
94	Functional analysis of the missense APOC3 mutation Ala23Thr associated with human hypotriglyceridemia. <i>Journal of Lipid Research</i> , 2010 , 51, 1524-34	6.3	46
93	A fully automated 2-D LC-MS method utilizing online continuous pH and RP gradients for global proteome analysis. <i>Electrophoresis</i> , 2007 , 28, 4311-9	3.6	43
92	Proteomics: from technology developments to biological applications. <i>Analytical Chemistry</i> , 2009 , 81, 4585-99	7.8	40
91	The liver connexin32 interactome is a novel plasma membrane-mitochondrial signaling nexus. <i>Journal of Proteome Research</i> , 2013 , 12, 2597-610	5.6	38
90	Palladium-Promoted DNA-Compatible Heck Reaction. <i>Organic Letters</i> , 2019 , 21, 719-723	6.2	37

89	Large-scale identification of human biliary proteins from a cholesterol stone patient using a proteomic approach. <i>Rapid Communications in Mass Spectrometry</i> , 2005 , 19, 3569-78	2.2	36
88	Technological developments in lipidomics. <i>Briefings in Functional Genomics & Proteomics</i> , 2008 , 7, 395-409		33
87	Quantitative proteomic analysis reveals the neuroprotective effects of huperzine A for amyloid beta treated neuroblastoma N2a cells. <i>Proteomics</i> , 2013 , 13, 1314-24	4.8	32
86	Proteomic reactors and their applications in biology. <i>FEBS Journal</i> , 2011 , 278, 3796-806	5.7	32
85	Non-transition Metal-Mediated Diverse Aryl-Heteroatom Bond Formation of Arylammonium Salts. <i>IScience</i> , 2019 , 15, 307-315	6.1	31
84	Triptolide Induces Cell Killing in Multidrug-Resistant Tumor Cells via CDK7/RPB1 Rather than XPB or p44. <i>Molecular Cancer Therapeutics</i> , 2016 , 15, 1495-503	6.1	31
83	Metaproteomics: A strategy to study the taxonomy and functionality of the gut microbiota. <i>Journal of Proteomics</i> , 2020 , 219, 103737	3.9	29
82	A novel USP9X substrate TTK contributes to tumorigenesis in non-small-cell lung cancer. <i>Theranostics</i> , 2018 , 8, 2348-2360	12.1	29
81	Regulation of septin dynamics by the <i>Saccharomyces cerevisiae</i> lysine acetyltransferase NuA4. <i>PLoS ONE</i> , 2011 , 6, e25336	3.7	29
80	Glycoproteomic reactor for human plasma. <i>Journal of Proteome Research</i> , 2009 , 8, 556-66	5.6	29
79	Improved recovery and identification of membrane proteins from rat hepatic cells using a centrifugal proteomic reactor. <i>Molecular and Cellular Proteomics</i> , 2011 , 10, O111.008425	7.6	29
78	Label-free proteomics uncovers energy metabolism and focal adhesion regulations responsive for endometrium receptivity. <i>Journal of Proteome Research</i> , 2015 , 14, 1831-42	5.6	28
77	mChIP-KAT-MS, a method to map protein interactions and acetylation sites for lysine acetyltransferases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, E1641-50	11.5	28
76	Real-Time Analysis on Drug-Antibody Ratio of Antibody-Drug Conjugates for Synthesis, Process Optimization, and Quality Control. <i>Scientific Reports</i> , 2017 , 7, 7763	4.9	27
75	Microsome-associated luminal lipid droplets in the regulation of lipoprotein secretion. <i>Current Opinion in Lipidology</i> , 2013 , 24, 160-70	4.4	27
74	Lyso-form fragment ions facilitate the determination of stereospecificity of diacyl glycerophospholipids. <i>Rapid Communications in Mass Spectrometry</i> , 2011 , 25, 205-17	2.2	27
73	Proteomic profiling of regionalized proteins in rat epididymis indicates consistency between specialized distribution and protein functions. <i>Journal of Proteome Research</i> , 2006 , 5, 299-307	5.6	27
72	Analytical aspects of proteomics: 2009-2010. <i>Analytical Chemistry</i> , 2011 , 83, 4407-26	7.8	26

71	Two-dimensional gel electrophoresis maps of the proteome and phosphoproteome of primitively cultured rat mesangial cells. <i>Electrophoresis</i> , 2005 , 26, 4540-62	3.6	26
70	Analysis of the subcellular phosphoproteome using a novel phosphoproteomic reactor. <i>Journal of Proteome Research</i> , 2010 , 9, 1279-88	5.6	25
69	GIAT4RA functions as a tumor suppressor in non-small cell lung cancer by counteracting Uchl3-mediated deubiquitination of LSH. <i>Oncogene</i> , 2019 , 38, 7133-7145	9.2	24
68	Quantitative analysis of secretome from adipocytes regulated by insulin. <i>Acta Biochimica Et Biophysica Sinica</i> , 2009 , 41, 910-21	2.8	24
67	SARS-CoV-2 envelope protein causes acute respiratory distress syndrome (ARDS)-like pathological damages and constitutes an antiviral target. <i>Cell Research</i> , 2021 , 31, 847-860	24.7	24
66	Prefractionation of proteome by liquid isoelectric focusing prior to two-dimensional liquid chromatography mass spectrometric identification. <i>Journal of Proteome Research</i> , 2005 , 4, 1256-64	5.6	23
65	Gsy, a novel glucansucrase from <i>Leuconostoc mesenteroides</i> , mediates the formation of cell aggregates in response to oxidative stress. <i>Scientific Reports</i> , 2016 , 6, 38122	4.9	23
64	Characterization of Gain-of-Function Mutant Provides New Insights into ClpP Structure. <i>ACS Chemical Biology</i> , 2016 , 11, 1964-72	4.9	23
63	Activation of AhR with nuclear IKK β regulates cancer stem-like properties in the occurrence of radioresistance. <i>Cell Death and Disease</i> , 2018 , 9, 490	9.8	22
62	Identification of Glutamyl Cyclase isoenzyme isoQC as a regulator of SIRPECD47 axis. <i>Cell Research</i> , 2019 , 29, 502-505	24.7	21
61	An intramolecular disulfide bond designed in myoglobin fine-tunes both protein structure and peroxidase activity. <i>Archives of Biochemistry and Biophysics</i> , 2016 , 600, 47-55	4.1	19
60	TGF- β /p65/MAT2A pathway regulates liver fibrogenesis via intracellular SAM. <i>EBioMedicine</i> , 2019 , 42, 458-469	8.8	18
59	Identification of a Novel Function of Adipocyte Plasma Membrane-Associated Protein (APMAP) in Gestational Diabetes Mellitus by Proteomic Analysis of Omental Adipose Tissue. <i>Journal of Proteome Research</i> , 2016 , 15, 628-37	5.6	18
58	Structural basis of rifampin inactivation by rifampin phosphotransferase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 3803-8	11.5	18
57	Global and Site-Specific Effect of Phosphorylation on Protein Turnover. <i>Developmental Cell</i> , 2021 , 56, 111-124.e6	10.2	16
56	Leflunomide Increases Hepatic Exposure to Methotrexate and Its Metabolite by Differentially Regulating Multidrug Resistance-Associated Protein Mrp2/3/4 Transporters via Peroxisome Proliferator-Activated Receptor Activation. <i>Molecular Pharmacology</i> , 2018 , 93, 563-574	4.3	15
55	Highly sensitive detection of S-nitrosylated proteins by capillary gel electrophoresis with laser induced fluorescence. <i>Journal of Chromatography A</i> , 2011 , 1218, 6756-62	4.5	15
54	Isoform-resolved correlation analysis between mRNA abundance regulation and protein level degradation. <i>Molecular Systems Biology</i> , 2020 , 16, e9170	12.2	14

53	Discovery of a Potential Plasma Protein Biomarker Panel for Acute-on-Chronic Liver Failure Induced by Hepatitis B Virus. <i>Frontiers in Physiology</i> , 2017 , 8, 1009	4.6	14
52	Analysis of low-abundance proteins using the proteomic reactor with pH fractionation. <i>Talanta</i> , 2010 , 80, 1526-31	6.2	14
51	Proteomic analysis of minute amount of colonic biopsies by enteroscopy sampling. <i>Biochemical and Biophysical Research Communications</i> , 2016 , 476, 286-292	3.4	14
50	Selective N-glycan editing on living cell surfaces to probe glycoconjugate function. <i>Nature Chemical Biology</i> , 2020 , 16, 766-775	11.7	13
49	Dissociative role for dorsal hippocampus in mediating heroin self-administration and relapse through CDK5 and RhoB signaling revealed by proteomic analysis. <i>Addiction Biology</i> , 2017 , 22, 1731-1742	4.6	13
48	Standardization and harmonization of distributed multi-center proteotype analysis supporting precision medicine studies. <i>Nature Communications</i> , 2020 , 11, 5248	17.4	13
47	Quantitative Proteomic Study Reveals Up-Regulation of cAMP Signaling Pathway-Related Proteins in Mild Traumatic Brain Injury. <i>Journal of Proteome Research</i> , 2018 , 17, 858-869	5.6	13
46	Combination of online enzyme digestion with stable isotope labeling for high-throughput quantitative proteome analysis. <i>Proteomics</i> , 2012 , 12, 3129-37	4.8	12
45	The cross-talk between methylation and phosphorylation in lymphoid-specific helicase drives cancer stem-like properties. <i>Signal Transduction and Targeted Therapy</i> , 2020 , 5, 197	21	12
44	Mouse embryonic stem cells have increased capacity for replication fork restart driven by the specific FliA-Floped protein complex. <i>Cell Research</i> , 2018 , 28, 69-89	24.7	12
43	DNA methylation modifier LSH inhibits p53 ubiquitination and transactivates p53 to promote lipid metabolism. <i>Epigenetics and Chromatin</i> , 2019 , 12, 59	5.8	11
42	Recent technological developments in proteomics shed new light on translational research on diabetic microangiopathy. <i>FEBS Journal</i> , 2013 , 280, 5668-81	5.7	11
41	Quantitative proteomic analysis of mice corneal tissues reveals angiogenesis-related proteins involved in corneal neovascularization. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2016 , 1864, 787-93	4	11
40	S100A11 Promotes Liver Steatosis via FOXO1-Mediated Autophagy and Lipogenesis. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2021 , 11, 697-724	7.9	11
39	Iron-induced energy supply deficiency and mitochondrial fragmentation in neurons. <i>Journal of Neurochemistry</i> , 2018 , 147, 816-830	6	11
38	Andrographolide Sulfonate Attenuates Acute Lung Injury by Reducing Expression of Myeloperoxidase and Neutrophil-Derived Proteases in Mice. <i>Frontiers in Physiology</i> , 2018 , 9, 939	4.6	11
37	Development of Online pH Gradient-Eluted Strong Cation Exchange Nano-electrospray-Tandem Mass Spectrometry for Proteomic Analysis Facilitating Basic and Histidine-Containing Peptides Identification. <i>Analytical Chemistry</i> , 2016 , 88, 583-91	7.8	10
36	New ammunition for the proteomic reactor: strong anion exchange beads and multiple enzymes enhance protein identification and sequence coverage. <i>Analytical and Bioanalytical Chemistry</i> , 2010 , 397, 3421-30	4.4	10

35	Analysis of microdissected cells by two-dimensional LC-MS approaches. <i>Methods in Molecular Biology</i> , 2008 , 428, 193-208	1.4	10
34	pVHL mediates K63-linked ubiquitination of IKK β leading to IKK β inactivation. <i>Cancer Letters</i> , 2016 , 383, 1-8	9.9	10
33	SCF/GSK3 β -Mediated GFI1 Degradation Suppresses Proliferation of Gastric Cancer Cells. <i>Cancer Research</i> , 2019 , 79, 4387-4398	10.1	9
32	Extracellular Signal-regulated Kinases (ERKs) Phosphorylate Lin28a Protein to Modulate P19 Cell Proliferation and Differentiation. <i>Journal of Biological Chemistry</i> , 2017 , 292, 3970-3976	5.4	8
31	SILAC-based quantitative proteomic analysis of the livers of spontaneous obese and diabetic rhesus monkeys. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2018 , 315, E294-E306	6	8
30	ITCH nuclear translocation and H1.2 polyubiquitination negatively regulate the DNA damage response. <i>Nucleic Acids Research</i> , 2019 , 47, 824-842	20.1	8
29	Identification of a USP9X Substrate NFX1-123 by SILAC-Based Quantitative Proteomics. <i>Journal of Proteome Research</i> , 2019 , 18, 2654-2665	5.6	7
28	A novel short-term high-lactose culture approach combined with a matrix-assisted laser desorption ionization-time of flight mass spectrometry assay for differentiating <i>Escherichia coli</i> and <i>Shigella</i> species using artificial neural networks. <i>PLoS ONE</i> , 2019 , 14, e0222636	3.7	7
27	Proteogenomic characterization identifies clinically relevant subgroups of intrahepatic cholangiocarcinoma. <i>Cancer Cell</i> , 2021 ,	24.3	7
26	LOXL1 confers antiapoptosis and promotes gliomagenesis through stabilizing BAG2. <i>Cell Death and Differentiation</i> , 2020 , 27, 3021-3036	12.7	6
25	Allosteric Regulation of Hsp90 α Activity by Small Molecules Targeting the Middle Domain of the Chaperone. <i>iScience</i> , 2020 , 23, 100857	6.1	5
24	Enhancing Membrane Protein Identification Using a Simplified Centrifugation and Detergent-Based Membrane Extraction Approach. <i>Analytical Chemistry</i> , 2018 , 90, 2434-2439	7.8	5
23	Quantitative Proteomics Reveals the Protective Effects of Huangqi Decoction Against Acute Cholestatic Liver Injury by Inhibiting the NF- κ B/IL-6/STAT3 Signaling Pathway. <i>Journal of Proteome Research</i> , 2020 , 19, 677-687	5.6	5
22	Discovery of a subtype-selective, covalent inhibitor against palmitoylation pocket of TEAD3. <i>Acta Pharmaceutica Sinica B</i> , 2021 , 11, 3206-3219	15.5	5
21	Quantitative proteomics reveals Shexiang Baoxin Pill exerts cardioprotective effects by preserving energy metabolism in a rat model of myocardial infarction. <i>Journal of Ethnopharmacology</i> , 2021 , 266, 113460	5	5
20	Identification of Highly Selective Lipoprotein-Associated Phospholipase A2 (Lp-PLA2) Inhibitors by a Covalent Fragment-Based Approach. <i>Journal of Medicinal Chemistry</i> , 2020 , 63, 7052-7065	8.3	4
19	Deep Phosphoproteomic Measurements Pinpointing Drug Induced Protective Mechanisms in Neuronal Cells. <i>Frontiers in Physiology</i> , 2016 , 7, 635	4.6	4
18	The proteomic study of serially passaged human skin fibroblast cells uncovers down-regulation of the chromosome condensin complex proteins involved in replicative senescence. <i>Biochemical and Biophysical Research Communications</i> , 2018 , 505, 1112-1120	3.4	4

17	Comparative Evaluation of Small Molecular Additives and Their Effects on Peptide/Protein Identification. <i>Analytical Chemistry</i> , 2017 , 89, 5784-5792	7.8	3
16	Suppression of asparagine synthetase enhances the antitumor potency of ART and artemalogue SOMCL-14-221 in non-small cell lung cancer. <i>Cancer Letters</i> , 2020 , 475, 22-33	9.9	3
15	Solution structure of extracellular loop of human β subunit of BK channel and its biological implication on ChTX sensitivity. <i>Scientific Reports</i> , 2018 , 8, 4571	4.9	3
14	Label-free quantitative proteomic analysis identifies CTNNB1 as a direct target of FOXP3 in gastric cancer cells. <i>Oncology Letters</i> , 2018 , 15, 7655-7660	2.6	3
13	Uncovering kappa-opioid receptor agonist-induced PAK1/2 phosphorylation by quantitative phosphoproteomics. <i>Biochemical and Biophysical Research Communications</i> , 2019 , 516, 320-326	3.4	3
12	Quantitative proteomic Analysis Reveals up-regulation of caveolin-1 in FOXP3-overexpressed human gastric cancer cells. <i>Scientific Reports</i> , 2017 , 7, 14460	4.9	2
11	Targeting the RT loop of Src SH3 in Platelets Prevents Thrombosis without Compromising Hemostasis.. <i>Advanced Science</i> , 2022 , e2103228	13.6	2
10	Dynamics of Post-Translational Modification Inspires Drug Design in the Kinase Family. <i>Journal of Medicinal Chemistry</i> , 2021 , 64, 15111-15125	8.3	2
9	The Active Constituent From <i>Gynostemma Pentaphyllum</i> Prevents Liver Fibrosis Through Regulation of the TGF- β /NDRG2/MAPK Axis. <i>Frontiers in Genetics</i> , 2020 , 11, 594824	4.5	2
8	Exploring biological basis of Syndrome differentiation in coronary heart disease patients with two distinct Syndromes by integrated multi-omics and network pharmacology strategy. <i>Chinese Medicine</i> , 2021 , 16, 109	4.7	1
7	Helix Matrix Transformation Combined With Convolutional Neural Network Algorithm for Matrix-Assisted Laser Desorption Ionization-Time of Flight Mass Spectrometry-Based Bacterial Identification. <i>Frontiers in Microbiology</i> , 2020 , 11, 565434	5.7	1
6	Revelation of AbfR in regulation of mismatch repair and energy metabolism in <i>S. epidermidis</i> by integrated proteomic and metabolomic analysis. <i>Journal of Proteomics</i> , 2020 , 226, 103900	3.9	1
5	PGE2-JNK signaling axis non-canonically promotes Gli activation by protecting Gli2 from ubiquitin-proteasomal degradation. <i>Cell Death and Disease</i> , 2021 , 12, 707	9.8	1
4	Quantitative proteomic analysis uncovers inhibition of melanin synthesis by silk fibroin via MITF/tyrosinase axis in B16 melanoma cells. <i>Life Sciences</i> , 2021 , 284, 119930	6.8	1
3	Establishment of a pseudovirus neutralization assay based on SARS-CoV-2 S protein incorporated into lentiviral particles.. <i>Biosafety and Health</i> , 2022 , 4, 38-38	4.7	0
2	A novel lncRNA Discn fine-tunes replication protein A (RPA) availability to promote genomic stability. <i>Nature Communications</i> , 2021 , 12, 5572	17.4	0
1	Serum proteomic analysis reveals the cardioprotective effects of Shexiang Baoxin Pill and Suxiao Jiuxin Pill in a rat model of acute myocardial infarction.. <i>Journal of Ethnopharmacology</i> , 2022 , 115279	5	0