

Mian Wu

List of Publications by Year in descending order

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103
papers

12,420
citations

50170

46
h-index

31759

101
g-index

105
all docs

105
docs citations

105
times ranked

25637
citing authors

#	ARTICLE	IF	CITATIONS
1	The long noncoding RNA glycoLINC assembles a lower glycolytic metabolon to promote glycolysis. <i>Molecular Cell</i> , 2022, 82, 542-554.e6.	4.5	32
2	TRIM27 cooperates with STK38L to inhibit ULK1-mediated autophagy and promote tumorigenesis. <i>EMBO Journal</i> , 2022, 41, .	3.5	18
3	Non-coding RNAs, guardians of the p53 galaxy. <i>Seminars in Cancer Biology</i> , 2021, 75, 72-83.	4.3	27
4	DDIT3 Directs a Dual Mechanism to Balance Glycolysis and Oxidative Phosphorylation during Glutamine Deprivation. <i>Advanced Science</i> , 2021, 8, e2003732.	5.6	15
5	LncRNA GIRGL drives CAPRIN1-mediated phase separation to suppress glutaminase-1 translation under glutamine deprivation. <i>Science Advances</i> , 2021, 7, .	4.7	38
6	PRMT1 Contributes to HNF1A-mediated Regulation Network of CYP3A4 in Huh7 Cells. <i>FASEB Journal</i> , 2021, 35, .	0.2	0
7	lncRNA TRMP-S directs dual mechanisms to regulate p27-mediated cellular senescence. <i>Molecular Therapy - Nucleic Acids</i> , 2021, 24, 971-985.	2.3	13
8	Non-coding RNAs, metabolic stress and adaptive mechanisms in cancer. <i>Cancer Letters</i> , 2020, 491, 60-69.	3.2	10
9	Lnc RNA GUARDIN suppresses cellular senescence through a LRP 130-PGC 1±-FOXO 4±-dependent signaling axis. <i>EMBO Reports</i> , 2020, 21, e48796.	2.0	11
10	SENEBLOC, a long non-coding RNA suppresses senescence via p53-dependent and independent mechanisms. <i>Nucleic Acids Research</i> , 2020, 48, 3089-3102.	6.5	39
11	CircACC1 Regulates Assembly and Activation of AMPK Complex under Metabolic Stress. <i>Cell Metabolism</i> , 2019, 30, 157-173.e7.	7.2	209
12	The lncRNA Neat1 promotes activation of inflammasomes in macrophages. <i>Nature Communications</i> , 2019, 10, 1495.	5.8	323
13	TP53LNC-DB, the database of lncRNAs in the p53 signalling network. <i>Database: the Journal of Biological Databases and Curation</i> , 2019, 2019, .	1.4	7
14	Aurora-A mediated phosphorylation of LDHB promotes glycolysis and tumor progression by relieving the substrate-inhibition effect. <i>Nature Communications</i> , 2019, 10, 5566.	5.8	66
15	TP53, TP53 Target Genes (DRAM, TIGAR), and Autophagy. <i>Advances in Experimental Medicine and Biology</i> , 2019, 1206, 127-149.	0.8	32
16	Tumor-suppressive or tumor-supportive: For p53, that is the question. <i>Molecular and Cellular Oncology</i> , 2018, 5, e1408537.	0.3	3
17	LncRNA IDH1-AS1 links the functions of c-Myc and HIF1± via IDH1 to regulate the Warburg effect. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E1465-E1474.	3.3	93
18	Regulation of the Mdm2-p53 pathway by the ubiquitin E3 ligase <sc>MARCH</sc> 7. <i>EMBO Reports</i> , 2018, 19, 305-319.	2.0	48

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19	GUARDIN is a p53-responsive long non-coding RNA that is essential for genomic stability. <i>Nature Cell Biology</i> , 2018, 20, 492-502.	4.6	239
20	Dual functions for OVAAL in initiation of RAF/MEK/ERK prosurvival signals and evasion of p27-mediated cellular senescence. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E11661-E11670.	3.3	52
21	Fatty acid synthesis is critical for stem cell pluripotency via promoting mitochondrial fission. <i>EMBO Journal</i> , 2017, 36, 1330-1347.	3.5	110
22	The p53-inducible long noncoding RNA TRINGS protects cancer cells from necrosis under glucose starvation. <i>EMBO Journal</i> , 2017, 36, 3483-3500.	3.5	66
23	PHLDA3 impedes somatic cell reprogramming by activating Akt-GSK3 β pathway. <i>Scientific Reports</i> , 2017, 7, 2832.	1.6	12
24	LAST, a c-Myc-inducible long noncoding RNA, cooperates with CNBP to promote CCND1 mRNA stability in human cells. <i>ELife</i> , 2017, 6, .	2.8	67
25	A Novel Sex Chromosome Mosaicism 45,X/45,Y/46,XY/46,YY/47,XYXYY Causing Ambiguous Genitalia. <i>Annals of Clinical and Laboratory Science</i> , 2017, 47, 761-764.	0.2	2
26	Noncoding RNAs Regulating p53 and c-Myc Signaling. <i>Advances in Experimental Medicine and Biology</i> , 2016, 927, 337-365.	0.8	8
27	LncRNA MIF, a c-Myc-activated long noncoding RNA, suppresses glycolysis by promoting Fbxw7-mediated c-Myc degradation. <i>EMBO Reports</i> , 2016, 17, 1204-1220.	2.0	92
28	A redox mechanism underlying nucleolar stress sensing by nucleophosmin. <i>Nature Communications</i> , 2016, 7, 13599.	5.8	94
29	Guidelines for the use and interpretation of assays for monitoring autophagy (3rd edition). <i>Autophagy</i> , 2016, 12, 1-222.	4.3	4,701
30	MIR-181a regulates lipid metabolism via IDH1. <i>Scientific Reports</i> , 2015, 5, 8801.	1.6	53
31	BECN1s, a short splice variant of BECN1, functions in mitophagy. <i>Autophagy</i> , 2015, 11, 2048-2056.	4.3	29
32	miRNA-181 regulates embryo implantation in mice through targeting leukemia inhibitory factor. <i>Journal of Molecular Cell Biology</i> , 2015, 7, 12-22.	1.5	44
33	Mitochondrial E3 ligase March5 maintains stemness of mouse ES cells via suppression of ERK signalling. <i>Nature Communications</i> , 2015, 6, 7112.	5.8	34
34	Pluripotency Activity of Nanog Requires Biochemical Stabilization by Variant Histone Protein H2A.Z. <i>Stem Cells</i> , 2015, 33, 2126-2134.	1.4	10
35	Chaperone-mediated autophagy prevents apoptosis by degrading BBC3/PUMA. <i>Autophagy</i> , 2015, 11, 1623-1635.	4.3	50
36	Orphan nuclear receptor TR3 acts in autophagic cell death via mitochondrial signaling pathway. <i>Nature Chemical Biology</i> , 2014, 10, 133-140.	3.9	193

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37	Reciprocal Regulation of HIF-1 β and LincRNA-p21 Modulates the Warburg Effect. <i>Molecular Cell</i> , 2014, 53, 88-100.	4.5	453
38	Regulation of the pentose phosphate pathway in cancer. <i>Protein and Cell</i> , 2014, 5, 592-602.	4.8	363
39	FITC α -quencher based caspase 3-activatable nanoprobe for effectively sensing caspase 3 in vitro and in cells. <i>Nanoscale</i> , 2013, 5, 8963.	2.8	26
40	Endoplasmic reticulum stress inhibits cell cycle progression via induction of p27 in melanoma cells. <i>Cellular Signalling</i> , 2013, 25, 144-149.	1.7	55
41	Retinoic acid-induced HOXA5 expression is co-regulated by HuR and miR-130a. <i>Cellular Signalling</i> , 2013, 25, 1476-1485.	1.7	32
42	Regulation of α -Threonine Dehydrogenase in Somatic Cell Reprogramming. <i>Stem Cells</i> , 2013, 31, 953-965.	1.4	64
43	XIAP inhibits autophagy via XIAP-Mdm2-p53 signalling. <i>EMBO Journal</i> , 2013, 32, 2204-2216.	3.5	120
44	TAp73 enhances the pentose phosphate pathway and supports cell proliferation. <i>Nature Cell Biology</i> , 2013, 15, 991-1000.	4.6	198
45	Siva1 inhibits p53 function by acting as an ARF E3 ubiquitin ligase. <i>Nature Communications</i> , 2013, 4, 1551.	5.8	47
46	Crystal structure and functional characterization of the human RBM25 PWI domain and its flanking basic region. <i>Biochemical Journal</i> , 2013, 450, 85-94.	1.7	8
47	A new role of p53 in regulating lipid metabolism. <i>Journal of Molecular Cell Biology</i> , 2013, 5, 147-150.	1.5	55
48	DEVD-Based Hydrogelator Minimizes Cellular Apoptosis Induction. <i>Scientific Reports</i> , 2013, 3, 1848.	1.6	18
49	PI(4,5)P2 5-phosphatase A regulates PI3K/Akt signalling and has a tumour suppressive role in human melanoma. <i>Nature Communications</i> , 2013, 4, 1508.	5.8	67
50	c-Myc modulates microRNA processing via the transcriptional regulation of Drosha. <i>Scientific Reports</i> , 2013, 3, 1942.	1.6	57
51	Combinatorial readout of unmodified H3R2 and acetylated H3K14 by the tandem PHD finger of MOZ reveals a regulatory mechanism for <i>HOXA9</i> transcription. <i>Genes and Development</i> , 2012, 26, 1376-1391.	2.7	99
52	p53-Facilitated miR-199a-3p Regulates Somatic Cell Reprogramming. <i>Stem Cells</i> , 2012, 30, 1405-1413.	1.4	65
53	Multifaceted functions of Siva-1: more than an Indian God of Destruction. <i>Protein and Cell</i> , 2012, 3, 117-122.	4.8	14
54	RUVBL2 is a novel repressor of ARF transcription. <i>FEBS Letters</i> , 2012, 586, 435-441.	1.3	11

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55	p53 regulates biosynthesis through direct inactivation of glucose-6-phosphate dehydrogenase. <i>Nature Cell Biology</i> , 2011, 13, 310-316.	4.6	620
56	E2F1 inhibits MDM2 expression in a p53-dependent manner. <i>Cellular Signalling</i> , 2011, 23, 193-200.	1.7	17
57	Recognition of Unmodified Histone H3 by the First PHD Finger of Bromodomain-PHD Finger Protein 2 Provides Insights into the Regulation of Histone Acetyltransferases Monocytic Leukemic Zinc-finger Protein (MOZ) and MOZ-related factor (MORF). <i>Journal of Biological Chemistry</i> , 2011, 286, 36944-36955.	1.6	56
58	Siva1 suppresses epithelial-to-mesenchymal transition and metastasis of tumor cells by inhibiting stathmin and stabilizing microtubules. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 12851-12856.	3.3	86
59	MicroRNA-149*, a p53-responsive microRNA, functions as an oncogenic regulator in human melanoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 15840-15845.	3.3	168
60	p53 Dependent Centrosome Clustering Prevents Multipolar Mitosis in Tetraploid Cells. <i>PLoS ONE</i> , 2011, 6, e27304.	1.1	27
61	Nucleophosmin/B23 Inhibits Eg5-mediated Microtubule Depolymerization by Inactivating Its ATPase Activity. <i>Journal of Biological Chemistry</i> , 2010, 285, 19060-19067.	1.6	17
62	E2F1 represses β -catenin/TCF activity by direct up-regulation of Siah1. <i>Journal of Cellular and Molecular Medicine</i> , 2009, 13, 1719-1727.	1.6	20
63	Biocompatible, Luminescent Silver@Phenol Formaldehyde Resin Core/Shell Nanospheres: Large-Scale Synthesis and Application for In Vivo Bioimaging. <i>Advanced Functional Materials</i> , 2008, 18, 872-879.	7.8	156
64	Sumoylation is critical for DJ-1 to repress p53 transcriptional activity. <i>FEBS Letters</i> , 2008, 582, 1151-1156.	1.3	47
65	Multiple roles for nuclear localization signal (NLS, aa 442-472) of receptor interacting protein 3 (RIP3). <i>Biochemical and Biophysical Research Communications</i> , 2008, 372, 850-855.	1.0	10
66	DJ-1 Decreases Bax Expression through Repressing p53 Transcriptional Activity. <i>Journal of Biological Chemistry</i> , 2008, 283, 4022-4030.	1.6	207
67	Tumor Suppressor ARF Promotes Non-classic Proteasome-independent Polyubiquitination of COMMD1. <i>Journal of Biological Chemistry</i> , 2008, 283, 11453-11460.	1.6	18
68	Domain-swapped Dimerization of the Second PDZ Domain of ZO2 May Provide a Structural Basis for the Polymerization of Claudins. <i>Journal of Biological Chemistry</i> , 2007, 282, 35988-35999.	1.6	30
69	Noxa/Mcl-1 Balance Regulates Susceptibility of Cells to Camptothecin-Induced Apoptosis. <i>Neoplasia</i> , 2007, 9, 871-881.	2.3	47
70	Death effector domain DEDa, a self-cleaved product of caspase-8/Mch5, translocates to the nucleus by binding to ERK1/2 and upregulates procaspase-8 expression via a p53-dependent mechanism. <i>EMBO Journal</i> , 2007, 26, 1068-1080.	3.5	26
71	Phosphorylation of Pirh2 by Calmodulin-dependent kinase II impairs its ability to ubiquitinate p53. <i>EMBO Journal</i> , 2007, 26, 3062-3074.	3.5	43
72	p53 and Bad: remote strangers become close friends. <i>Cell Research</i> , 2007, 17, 283-285.	5.7	35

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73	p45, an ATPase subunit of the 19S proteasome, targets the polyglutamine disease protein ataxin-3 to the proteasome. <i>Journal of Neurochemistry</i> , 2007, 101, 1651-1661.	2.1	17
74	Cleavage of RIP3 inactivates its caspase-independent apoptosis pathway by removal of kinase domain. <i>Cellular Signalling</i> , 2007, 19, 2056-2067.	1.7	388
75	Truncated RIP3 (tRIP3) acts upstream of FADD to induce apoptosis in the human hepatocellular carcinoma cell line QGY-7703. <i>Biochemical and Biophysical Research Communications</i> , 2006, 347, 558-565.	1.0	10
76	Novel link between E2F1 and Smac/DIABLO: proapoptotic Smac/DIABLO is transcriptionally upregulated by E2F1. <i>Nucleic Acids Research</i> , 2006, 34, 2046-2055.	6.5	41
77	Acetylation of p53 at Lysine 373/382 by the Histone Deacetylase Inhibitor Depsipeptide Induces Expression of p21 Waf1/Cip1. <i>Molecular and Cellular Biology</i> , 2006, 26, 2782-2790.	1.1	265
78	The Bad Guy Cooperates with Good Cop p53: Bad Is Transcriptionally Up-Regulated by p53 and Forms a Bad/p53 Complex at the Mitochondria To Induce Apoptosis. <i>Molecular and Cellular Biology</i> , 2006, 26, 9071-9082.	1.1	134
79	Identification of a novel nucleolar localization signal and a degradation signal in Survivin-deltaEx3: a potential link between nucleolus and protein degradation. <i>Oncogene</i> , 2005, 24, 2723-2734.	2.6	51
80	Puma*Mcl-1 interaction is not sufficient to prevent rapid degradation of Mcl-1. <i>Oncogene</i> , 2005, 24, 7224-7237.	2.6	57
81	COQ9, a New Gene Required for the Biosynthesis of Coenzyme Q in <i>Saccharomyces cerevisiae</i> . <i>Journal of Biological Chemistry</i> , 2005, 280, 31397-31404.	1.6	84
82	RIP3 $\hat{1}$ ² and RIP3 $\hat{1}$ ³ , two novel splice variants of receptor-interacting protein 3 (RIP3), downregulate RIP3-induced apoptosis. <i>Biochemical and Biophysical Research Communications</i> , 2005, 332, 181-187.	1.0	26
83	A Single Amino Acid Change (Asp 53 $\hat{1}$ Ala53) Converts Survivin from Anti-apoptotic to Pro-apoptotic. <i>Molecular Biology of the Cell</i> , 2004, 15, 1287-1296.	0.9	43
84	p53-Dependent Apoptotic Mechanism of a New Designer Bimetallic Compound Tri-phenyl Tin Benzimidazolethiol Copper Chloride (TPT-CuCl ₂): In Vivo Studies in Wistar Rats as Well as in Vitro Studies in Human Cervical Cancer Cells. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2004, 311, 22-33.	1.3	40
85	Characterization of Acid-sensing Ion Channels in Dorsal Horn Neurons of Rat Spinal Cord. <i>Journal of Biological Chemistry</i> , 2004, 279, 43716-43724.	1.6	169
86	Nucleocytoplasmic Shuttling of Receptor-interacting Protein 3 (RIP3). <i>Journal of Biological Chemistry</i> , 2004, 279, 38820-38829.	1.6	47
87	Etoposide upregulates Bax-enhancing tumour necrosis factor-related apoptosis inducing ligand-mediated apoptosis in the human hepatocellular carcinoma cell line QGY-7703. <i>FEBS Journal</i> , 2003, 270, 2721-2731.	0.2	22
88	Triphenyl Tin Benzimidazolethiol, a Novel Antitumor Agent, Induces Mitochondrial-Mediated Apoptosis in Human Cervical Cancer Cells via Suppression of HPV-18 Encoded E6. <i>Journal of Biochemistry</i> , 2003, 134, 521-528.	0.9	45
89	Bcl-rambo beta, a special splicing variant with an insertion of an Alu-like cassette, promotes etoposide- and Taxol-induced cell death. <i>FEBS Letters</i> , 2003, 534, 61-68.	1.3	31
90	Direct Interaction between Survivin and Smac/DIABLO Is Essential for the Anti-apoptotic Activity of Survivin during Taxol-induced Apoptosis. <i>Journal of Biological Chemistry</i> , 2003, 278, 23130-23140.	1.6	329

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91	HIV-1 TAT-mediated protein transduction and subcellular localization using novel expression vectors1. FEBS Letters, 2002, 532, 36-44.	1.3	40
92	Site-Specific Integration of the Double-Mutation Glucose Isomerase (GIG138PG247D) Gene in Streptomyces lividans and Its Stable Expression. Current Microbiology, 2002, 44, 18-24.	1.0	1
93	Induction of apoptosis in glioma cell lines by TRAIL/Apo-2l. Journal of Neuroscience Research, 2000, 61, 464-470.	1.3	34
94	MTO1 Codes for a Mitochondrial Protein Required for Respiration in Paromomycin-resistant Mutants of Saccharomyces cerevisiae. Journal of Biological Chemistry, 1998, 273, 27945-27952.	1.6	89
95	FLX1 Codes for a Carrier Protein Involved in Maintaining a Proper Balance of Flavin Nucleotides in Yeast Mitochondria. Journal of Biological Chemistry, 1996, 271, 7392-7397.	1.6	112
96	A novel method employing polymerase chain reaction to disrupt genes lacking convenient restriction enzyme sites in yeast. Molecular Biotechnology, 1995, 3, 72-74.	1.3	0
97	The Saccharomyces cerevisiae homologue of ribosomal protein S26. Gene, 1994, 150, 401-402.	1.0	5
98	YKE2, a yeast nuclear gene encoding a protein showing homology to mouse KE2 and containing a putative leucine-zipper motif. Gene, 1994, 151, 197-201.	1.0	4
99	Nucleotide sequence of a flower-specific MADS box cDNA clone from orchid. Plant Molecular Biology, 1993, 23, 901-904.	2.0	74
100	Nucleotide sequence and in vitro translation of the coat protein gene of cymbidium mosaic virus. Virus Genes, 1993, 7, 157-170.	0.7	11
101	Nucleotide sequence of a Singapore isolate of zucchini yellow mosaic virus coat protein gene revealed an altered DAG motif. Virus Genes, 1993, 7, 381-387.	0.7	13
102	Nucleotide sequence of the 3' half of zucchini yellow mosaic virus (Singapore isolate) genome encoding the 4K protein, protease, polymerase and coat protein. Nucleic Acids Research, 1993, 21, 1317-1317.	6.5	8
103	Nucleotide sequences of the two ORFs upstream to the coat protein gene of cymbidium mosaic virus. Plant Molecular Biology, 1992, 18, 1027-1029.	2.0	5