

Xi-Wen Liao

List of Publications by Year in descending order

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Version: 2024-02-01

58
papers

1,075
citations

361045

20
h-index

476904

29
g-index

58
all docs

58
docs citations

58
times ranked

1256
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification and validation of potential prognostic gene biomarkers for predicting survival in patients with acute myeloid leukemia. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 5243-5254.	1.0	93
2	Genome-scale analysis to identify prognostic markers in patients with early-stage pancreatic ductal adenocarcinoma after pancreaticoduodenectomy. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 4493-4506.	1.0	82
3	Distinct Diagnostic and Prognostic Values of Minichromosome Maintenance Gene Expression in Patients with Hepatocellular Carcinoma. <i>Journal of Cancer</i> , 2018, 9, 2357-2373.	1.2	59
4	Identification of potential prognostic microRNA biomarkers for predicting survival in patients with hepatocellular carcinoma. <i>Cancer Management and Research</i> , 2018, Volume 10, 787-803.	0.9	48
5	Identification of Potential Prognostic Long Non-Coding RNA Biomarkers for Predicting Survival in Patients with Hepatocellular Carcinoma. <i>Cellular Physiology and Biochemistry</i> , 2018, 48, 1854-1869.	1.1	37
6	Aldehyde dehydrogenase 1 (ALDH1) isoform expression and potential clinical implications in hepatocellular carcinoma. <i>PLoS ONE</i> , 2017, 12, e0182208.	1.1	35
7	The prognostic value of CYP2C subfamily genes in hepatocellular carcinoma. <i>Cancer Medicine</i> , 2018, 7, 966-980.	1.3	35
8	Genome-scale analysis to identify prognostic microRNA biomarkers in patients with early stage pancreatic ductal adenocarcinoma after pancreaticoduodenectomy. <i>Cancer Management and Research</i> , 2018, Volume 10, 2537-2551.	0.9	30
9	Genome-Wide Association Study of MKI67 Expression and its Clinical Implications in HBV-Related Hepatocellular Carcinoma in Southern China. <i>Cellular Physiology and Biochemistry</i> , 2017, 42, 1342-1357.	1.1	28
10	Distinct Diagnostic and Prognostic Values of Kinesin Family Member Genes Expression in Patients with Breast Cancer. <i>Medical Science Monitor</i> , 2018, 24, 9442-9464.	0.5	28
11	Comprehensive investigation of key biomarkers and pathways in hepatitis B virus-related hepatocellular carcinoma. <i>Journal of Cancer</i> , 2019, 10, 5689-5704.	1.2	28
12	The prognostic value of differentially expressed CYP3A subfamily members for hepatocellular carcinoma. <i>Cancer Management and Research</i> , 2018, Volume 10, 1713-1726.	0.9	27
13	Identification of prognostic biomarkers for patients with hepatocellular carcinoma after hepatectomy. <i>Oncology Reports</i> , 2019, 41, 1586-1602.	1.2	26
14	Genome-scale analysis to identify potential prognostic microRNA biomarkers for predicting overall survival in patients with colon adenocarcinoma. <i>Oncology Reports</i> , 2018, 40, 1947-1958.	1.2	25
15	Marker of proliferation Ki-67 expression is associated with transforming growth factor beta 1 and can predict the prognosis of patients with hepatic B virus-related hepatocellular carcinoma. <i>Cancer Management and Research</i> , 2018, Volume 10, 679-696.	0.9	25
16	Distinct diagnostic and prognostic values of aminobutyric acid type A receptor family genes in patients with colon adenocarcinoma. <i>Oncology Letters</i> , 2020, 20, 275-291.	0.8	25
17	Identification of key pathways and genes in TP53 mutation acute myeloid leukemia: evidence from bioinformatics analysis. <i>OncoTargets and Therapy</i> , 2018, Volume 11, 163-173.	1.0	24
18	Identification of F5 as a Prognostic Biomarker in Patients with Gastric Cancer. <i>BioMed Research International</i> , 2020, 2020, 1-13.	0.9	24

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19	XRCC1 rs25487 genetic variant and TP53 mutation at codon 249 predict clinical outcomes of hepatitis B virus-related hepatocellular carcinoma after hepatectomy: A cohort study for 10 years follow up. <i>Hepatology Research</i> , 2016, 46, 765-774.	1.8	23
20	Genome-wide investigation of the clinical significance and prospective molecular mechanisms of kinesin family member genes in patients with lung adenocarcinoma. <i>Oncology Reports</i> , 2019, 42, 1017-1034.	1.2	23
21	Integrated analysis of competing endogenous RNA network revealing potential prognostic biomarkers of hepatocellular carcinoma. <i>Journal of Cancer</i> , 2019, 10, 3267-3283.	1.2	23
22	Prognostic value of minichromosome maintenance mRNA expression in early-stage pancreatic ductal adenocarcinoma patients after pancreaticoduodenectomy. <i>Cancer Management and Research</i> , 2018, Volume 10, 3255-3271.	0.9	20
23	Distinct prognostic values of alcohol dehydrogenase mRNA expression in pancreatic adenocarcinoma. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 3719-3732.	1.0	18
24	The diagnosis and prognosis values of WNT mRNA expression in colon adenocarcinoma. <i>Journal of Cellular Biochemistry</i> , 2020, 121, 3145-3161.	1.2	18
25	Distinct prognostic value of dynactin subunit 4 (DCTN4) and diagnostic value of DCTN1, DCTN2, and DCTN4 in colon adenocarcinoma. <i>Cancer Management and Research</i> , 2018, Volume 10, 5807-5824.	0.9	17
26	Use of Genome-Scale Integrated Analysis to Identify Key Genes and Potential Molecular Mechanisms in Recurrence of Lower-Grade Brain Glioma. <i>Medical Science Monitor</i> , 2019, 25, 3716-3727.	0.5	15
27	Prognostic Value of Dynactin mRNA Expression in Cutaneous Melanoma. <i>Medical Science Monitor</i> , 2018, 24, 3752-3763.	0.5	13
28	Polymorphisms of HLA-DQB1 predict survival of hepatitis B virus-related hepatocellular carcinoma patients receiving hepatic resection. <i>Clinics and Research in Hepatology and Gastroenterology</i> , 2016, 40, 739-747.	0.7	12
29	Analysis of clinical significance and prospective molecular mechanism of main elements of the JAK/STAT pathway in hepatocellular carcinoma. <i>International Journal of Oncology</i> , 2019, 55, 805-822.	1.4	12
30	Noteworthy prognostic value of phospholipase C delta genes in early stage pancreatic ductal adenocarcinoma patients after pancreaticoduodenectomy and potential molecular mechanisms. <i>Cancer Medicine</i> , 2020, 9, 859-871.	1.3	12
31	Identify potential clinical significance of long noncoding RNA forkhead box P4 antisense RNA 1 in patients with early stage pancreatic ductal adenocarcinoma. <i>Cancer Medicine</i> , 2020, 9, 2062-2076.	1.3	12
32	CYP2C8 Suppress Proliferation, Migration, Invasion and Sorafenib Resistance of Hepatocellular Carcinoma via PI3K/Akt/p27kip1 Axis. <i>Journal of Hepatocellular Carcinoma</i> , 2021, Volume 8, 1323-1338.	1.8	12
33	Comprehensive investigation of the clinical significance and molecular mechanisms of plasmacytoma variant translocation 1 in sarcoma using genome-wide RNA sequencing data. <i>Journal of Cancer</i> , 2019, 10, 4961-4977.	1.2	11
34	Diagnostic and prognostic values of CXC motif chemokine ligand 3 in patients with colon cancer. <i>Oncology Reports</i> , 2019, 42, 1996-2008.	1.2	11
35	The Perspective of Diagnostic and Prognostic Values of Lipoxygenases mRNA Expression in Colon Adenocarcinoma. <i>OncoTargets and Therapy</i> , 2020, Volume 13, 9389-9405.	1.0	11
36	Identification of potential prognostic small nucleolar RNA biomarkers for predicting overall survival in patients with sarcoma. <i>Cancer Medicine</i> , 2020, 9, 7018-7033.	1.3	11

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37	Genome-wide association pathway analysis to identify candidate single nucleotide polymorphisms and molecular pathways associated with TP53 expression status in HBV-related hepatocellular carcinoma. <i>Cancer Management and Research</i> , 2018, Volume 10, 953-967.	0.9	10
38	Diagnostic and prognostic value of mRNA expression of phospholipase C β family genes in hepatitis B virus-associated hepatocellular carcinoma. <i>Oncology Reports</i> , 2019, 41, 2855-2875.	1.2	10
39	Genome-wide analysis to identify a novel microRNA signature that predicts survival in patients with stomach adenocarcinoma. <i>Journal of Cancer</i> , 2019, 10, 6298-6313.	1.2	9
40	Prognostic value of Glypican family genes in early-stage pancreatic ductal adenocarcinoma after pancreaticoduodenectomy and possible mechanisms. <i>BMC Gastroenterology</i> , 2020, 20, 415.	0.8	9
41	Genome-wide association study identified PLCE1- rs2797992 and EGFR- rs6950826 were associated with TP53 expression in the HBV-related hepatocellular carcinoma of Chinese patients in Guangxi. <i>American Journal of Translational Research (discontinued)</i> , 2016, 8, 1799-812.	0.0	8
42	Identification and verification of the molecular mechanisms and prognostic values of the cadherin gene family in gastric cancer. <i>Scientific Reports</i> , 2021, 11, 23674.	1.6	8
43	Genetic variants in the exon region of versican predict survival of patients with resected early-stage hepatitis B virus-associated hepatocellular carcinoma. <i>Cancer Management and Research</i> , 2018, Volume 10, 1027-1036.	0.9	7
44	Integrative Analysis of a Novel Eleven-Small Nucleolar RNA Prognostic Signature in Patients With Lower Grade Glioma. <i>Frontiers in Oncology</i> , 2021, 11, 650828.	1.3	7
45	High-Mobility Group AT-Hook 1 Served as a Prognosis Biomarker and Associated with Immune Infiltrate in Hepatocellular Carcinoma. <i>International Journal of General Medicine</i> , 2022, Volume 15, 609-621.	0.8	7
46	Prognostic Value of Excision Repair Cross-Complementing mRNA Expression in Gastric Cancer. <i>BioMed Research International</i> , 2018, 2018, 1-16.	0.9	6
47	Prognostic value and potential molecular mechanism of the like-Sm gene family in early-stage pancreatic ductal adenocarcinoma. <i>Translational Cancer Research</i> , 2021, 10, 1744-1760.	0.4	6
48	Cystatin F as a key family 2 cystatin subunit and prognostic biomarker for early-stage pancreatic ductal adenocarcinoma. <i>Oncology Reports</i> , 2019, 42, 79-90.	1.2	5
49	Prognostic value of Kinesin β family genes mRNA expression in early-stage pancreatic ductal adenocarcinoma patients after pancreaticoduodenectomy. <i>Cancer Medicine</i> , 2019, 8, 6487-6502.	1.3	4
50	Clinical significance and prospective molecular mechanism of C motif chemokine receptors in patients with early-stage pancreatic ductal adenocarcinoma after pancreaticoduodenectomy. <i>Oncology Reports</i> , 2019, 42, 1856-1868.	1.2	4
51	Genome-scale integrated analysis to identify prospective molecular mechanisms and therapeutic targets in isocitrate dehydrogenase 2 R140Q-mutated acute myeloid leukemia. <i>Oncology Reports</i> , 2019, 41, 2876-2888.	1.2	4
52	Genome-wide RNA-sequencing dataset reveals the prognostic value and potential molecular mechanisms of lncRNA in non-homologous end joining pathway 1 in early stage Pancreatic Ductal Adenocarcinoma. <i>Journal of Cancer</i> , 2020, 11, 5556-5567.	1.2	4
53	Comprehensive investigation of the clinical significance of long non-coding RNA HOXA-AS2 in acute myeloid leukemia using genome-wide RNA sequencing dataset. <i>Journal of Cancer</i> , 2021, 12, 2151-2164.	1.2	4
54	Investigation and verification of the clinical significance and perspective of natural killer group 2 member D ligands in colon adenocarcinoma. <i>Aging</i> , 2021, 13, 12565-12586.	1.4	4

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55	Integrative genomic analysis of a novel small nucleolar RNAs prognostic signature in patients with acute myelocytic leukemia. <i>Mathematical Biosciences and Engineering</i> , 2022, 19, 2424-2452.	1.0	4
56	LIXT antisense RNA 1 sever as a novel prognostic long non-coding RNA in early stage pancreatic ductal adenocarcinoma patients after receiving pancreaticoduodenectomy. <i>Journal of Cancer</i> , 2021, 12, 2122-2139.	1.2	1
57	Acyl-CoA Binding Domain Containing 4 Polymorphism rs4986172 and Expression Can Serve as Overall Survival Biomarkers for Hepatitis B Virus-Related Hepatocellular Carcinoma Patients After Hepatectomy. <i>Pharmacogenomics and Personalized Medicine</i> , 2022, Volume 15, 277-300.	0.4	1
58	Protein Phosphatase 1 Regulatory Subunit 3: A Prognostic Biomarker in Stomach Adenocarcinoma. <i>International Journal of General Medicine</i> , 2022, Volume 15, 1131-1146.	0.8	0