

Jiaheng Xie

List of Publications by Year in descending order

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Version: 2024-02-01

25
papers

349
citations

933264

10
h-index

940416

16
g-index

27
all docs

27
docs citations

27
times ranked

61
citing authors

#	ARTICLE	IF	CITATIONS
1	ADM-assisted prepectoral breast reconstruction is not associated with high complication rate as before: a Meta-analysis. <i>Journal of Plastic Surgery and Hand Surgery</i> , 2023, 57, 7-15.	0.4	10
2	Construction and validation of the diagnostic model of keloid based on weighted gene co-expression network analysis (WGCNA) and differential expression analysis. <i>Journal of Plastic Surgery and Hand Surgery</i> , 2022, , 1-9.	0.4	6
3	A novel signature based on autophagy-related lncRNA for prognostic prediction and candidate drugs for lung adenocarcinoma. <i>Translational Cancer Research</i> , 2022, 11, 14-28.	0.4	14
4	A novel regulatory mechanism network mediated by lncRNA TUG1 that induces the impairment of spiral artery remodeling in preeclampsia. <i>Molecular Therapy</i> , 2022, 30, 1692-1705.	3.7	18
5	A Necroptosis-Related Prognostic Model of Uveal Melanoma Was Constructed by Single-Cell Sequencing Analysis and Weighted Co-Expression Network Analysis Based on Public Databases. <i>Frontiers in Immunology</i> , 2022, 13, 847624.	2.2	34
6	Pan-cancer analysis revealed the significance of the GTPBP family in cancer. <i>Aging</i> , 2022, 14, 2558-2573.	1.4	2
7	Transcriptome Analysis of Solute Carrier-Associated Genes in Hepatocellular Carcinoma: Friend or Foe?. <i>Frontiers in Genetics</i> , 2022, 13, 856393.	1.1	3
8	Integrative analysis of TRPV family to prognosis and immune infiltration in renal clear cell carcinoma. <i>Channels</i> , 2022, 16, 84-96.	1.5	5
9	An immune subtype-related prognostic signature of hepatocellular carcinoma based on single-cell sequencing analysis. <i>Aging</i> , 2022, 14, 3276-3292.	1.4	34
10	Single-Cell Sequencing Analysis Based on Public Databases for Constructing a Metastasis-Related Prognostic Model for Gastric Cancer. <i>Applied Bionics and Biomechanics</i> , 2022, 2022, 1-19.	0.5	5
11	Head and Neck Squamous Cell Carcinoma: NT5E Could Be a Prognostic Biomarker. <i>Applied Bionics and Biomechanics</i> , 2022, 2022, 1-14.	0.5	1
12	Solute carrier transporter superfamily member SLC16A1 is a potential prognostic biomarker and associated with immune infiltration in skin cutaneous melanoma. <i>Channels</i> , 2021, 15, 483-495.	1.5	19
13	Pan-cancer analysis and single-cell analysis revealed the role of ABCC5 transporter in hepatocellular carcinoma. <i>Channels</i> , 2021, 15, 541-554.	1.5	9
14	Database mining analysis revealed the role of the putative H ⁺ /sugar transporter solute carrier family 45 in skin cutaneous melanoma. <i>Channels</i> , 2021, 15, 496-506.	1.5	18
15	A pan-cancer analysis revealed the role of the SLC16 family in cancer. <i>Channels</i> , 2021, 15, 528-540.	1.5	18
16	A Novel Pyroptosis-Related lncRNA Signature for Predicting the Prognosis of Skin Cutaneous Melanoma. <i>International Journal of General Medicine</i> , 2021, Volume 14, 6517-6527.	0.8	46
17	Identification of ABCC5 Among ATP-Binding Cassette Transporter Family as a New Biomarker for Hepatocellular Carcinoma Based on Bioinformatics Analysis. <i>International Journal of General Medicine</i> , 2021, Volume 14, 7235-7246.	0.8	34
18	Clinical Value for Diagnosis and Prognosis of Signal Sequence Receptor 1 (SSR1) and Its Potential Mechanism in Hepatocellular Carcinoma: A Comprehensive Study Based on High-Throughput Data Analysis. <i>International Journal of General Medicine</i> , 2021, Volume 14, 7435-7451.	0.8	3

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19	ANXA1 (Annexin A1) regulated by MYC (MYC proto-oncogene) promotes the growth of papillary thyroid carcinoma. <i>Bioengineered</i> , 2021, 12, 9251-9265.	1.4	10
20	Prognostic Value of SLC16A3(MCT4) in Lung Adenocarcinoma and Its Clinical Significance. <i>International Journal of General Medicine</i> , 2021, Volume 14, 8413-8425.	0.8	14
21	Construction and Validation of a Novel Pyroptosis-Related Gene Signature to Predict the Prognosis of Uveal Melanoma. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 761350.	1.8	19
22	Transcriptomic Alterations Induced By Vemurafenib after Treatment of Melanoma: A Comprehensive Bioinformatics Analysis. , 2021, 83, .		0
23	Single-Cell Sequencing Analysis and Weighted Co-Expression Network Analysis Based on Public Databases Identified That TNC Is a Novel Biomarker for Keloid. <i>Frontiers in Immunology</i> , 2021, 12, 783907.	2.2	23
24	Integrated Analysis of Genomic and Transcriptomic Profiles Identified the Role of GTP Binding Protein-4 (GTPBP4) in Breast Cancer. <i>Frontiers in Pharmacology</i> , 0, 13, .	1.6	3
25	Response: Commentary: Single-Cell Sequencing Analysis and Weighted CoExpression Network Analysis Based on Public Databases Identified That TNC Is a Novel Biomarker for Keloid. <i>Frontiers in Immunology</i> , 0, 13, .	2.2	1