

Xiangqian Guo

List of Publications by Year in descending order

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Version: 2024-02-01

83
papers

3,569
citations

236925

25
h-index

144013

57
g-index

83
all docs

83
docs citations

83
times ranked

6149
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification and Validation of an Annexin-Related Prognostic Signature and Therapeutic Targets for Bladder Cancer: Integrative Analysis. <i>Biology</i> , 2022, 11, 259.	2.8	7
2	OShnscc: a novel user-friendly online survival analysis tool for head and neck squamous cell carcinoma based on RNA expression profiles and long-term survival information. <i>Journal of Zhejiang University: Science B</i> , 2022, 23, 249-257.	2.8	1
3	OSgc: A Web Portal to Assess the Performance of Prognostic Biomarkers in Gastric Cancer. <i>Frontiers in Oncology</i> , 2022, 12, 856988.	2.8	2
4	OSov: An Interactive Web Server to Evaluate Prognostic Biomarkers for Ovarian Cancer. <i>Biology</i> , 2022, 11, 23.	2.8	0
5	The Emerging Role of Long Non-Coding RNAs in Esophageal Cancer: Functions in Tumorigenesis and Clinical Implications. <i>Frontiers in Pharmacology</i> , 2022, 13, .	3.5	4
6	Comment on "The expression landscape of cachexia-inducing factors in human cancers" by Freire et al. <i>Journal of Cachexia, Sarcopenia and Muscle</i> , 2021, 12, 523-524.	7.3	1
7	The Abnormal Glycopatterns of Salivary Glycoproteins in Esophageal Squamous Cell Carcinoma Patients. <i>Frontiers in Chemistry</i> , 2021, 9, 637730.	3.6	7
8	OSchol: an online consensus survival web server for cholangiocarcinoma prognosis analysis. <i>Hpb</i> , 2021, 23, 545-550.	0.3	3
9	Sex differences in changes in BMI and blood pressure in Chinese school-aged children during the COVID-19 quarantine. <i>International Journal of Obesity</i> , 2021, 45, 2132-2136.	3.4	24
10	Identification of Clinical Relevant Molecular Subtypes of Pheochromocytoma. <i>Frontiers in Endocrinology</i> , 2021, 12, 605797.	3.5	2
11	Identification of Distinct Molecular Subtypes of Endometrioid Adenocarcinoma. <i>Frontiers in Genetics</i> , 2021, 12, 568779.	2.3	0
12	Sex-Specific Differences in Related Indicators of Blood Pressure in School-Age Children With Overweight and Obesity: A Cross-Sectional Study. <i>Frontiers in Pediatrics</i> , 2021, 9, 674504.	1.9	1
13	Editorial: Big Data and Machine Learning in Cancer Genomics. <i>Frontiers in Genetics</i> , 2021, 12, 749584.	2.3	2
14	Comprehensive analysis of lncRNAs as biomarkers for diagnosis, prognosis, and treatment response in clear cell renal cell carcinoma. <i>Molecular Therapy - Oncolytics</i> , 2021, 22, 209-218.	4.4	6
15	The translational values of TRIM family in pan-cancers: From functions and mechanisms to clinics. <i>Frontiers in Molecular and Cellular Oncology</i> , 2021, 227, 107881.		27
16	The Identification of RNA Modification Gene PUS7 as a Potential Biomarker of Ovarian Cancer. <i>Biology</i> , 2021, 10, 1130.	2.8	13
17	OSumv: An interactive online consensus survival tool for uveal melanoma prognosis analysis. <i>Molecular Carcinogenesis</i> , 2020, 59, 56-61.	2.7	18
18	OSdblcl: An online consensus survival analysis web server based on gene expression profiles of diffuse large B-cell lymphoma. <i>Cancer Medicine</i> , 2020, 9, 1790-1797.	2.8	10

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19	OSpaad: An online tool to perform survival analysis by integrating gene expression profiling and long-term follow-up data of 1319 pancreatic carcinoma patients. <i>Molecular Carcinogenesis</i> , 2020, 59, 304-310.	2.7	16
20	OSlgg: An Online Prognostic Biomarker Analysis Tool for Low-Grade Glioma. <i>Frontiers in Oncology</i> , 2020, 10, 1097.	2.8	6
21	Systematic Review of Prognostic Gene Signature in Gastric Cancer Patients. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020, 8, 805.	4.1	13
22	Separation and characterization of extracellular vesicles from human plasma by asymmetrical flow field-flow fractionation. <i>Analytica Chimica Acta</i> , 2020, 1127, 234-245.	5.4	41
23	Gene expression profile identifies distinct molecular subtypes and potential therapeutic genes in Merkel cell carcinoma. <i>Translational Oncology</i> , 2020, 13, 100816.	3.7	4
24	Editorial: Bioinformatics Tools (and Web Server) for Cancer Biomarker Development. <i>Frontiers in Oncology</i> , 2020, 10, 599085.	2.8	2
25	Risk factors for in-hospital mortality in patients with cancer and COVID-19. <i>Lancet Oncology</i> , The, 2020, 21, e407.	10.7	1
26	OSucs: An Online Prognostic Biomarker Analysis Tool for Uterine Carcinosarcoma. <i>Genes</i> , 2020, 11, 1040.	2.4	4
27	Systematic Analysis of microRNA Biomarkers for Diagnosis, Prognosis, and Therapy in Patients With Clear Cell Renal Cell Carcinoma. <i>Frontiers in Oncology</i> , 2020, 10, 543817.	2.8	10
28	OSmfs: An Online Interactive Tool to Evaluate Prognostic Markers for Myxofibrosarcoma. <i>Genes</i> , 2020, 11, 1523.	2.4	3
29	OSskcm: an online survival analysis webserver for skin cutaneous melanoma based on 1085 transcriptomic profiles. <i>Cancer Cell International</i> , 2020, 20, 176.	4.1	16
30	OSlihc: An Online Prognostic Biomarker Analysis Tool for Hepatocellular Carcinoma. <i>Frontiers in Pharmacology</i> , 2020, 11, 875.	3.5	15
31	The Application of Deep Learning in Cancer Prognosis Prediction. <i>Cancers</i> , 2020, 12, 603.	3.7	218
32	Identification of hub genes and pathways in adrenocortical carcinoma by integrated bioinformatic analysis. <i>Journal of Cellular and Molecular Medicine</i> , 2020, 24, 4428-4438.	3.6	16
33	OSeac: An Online Survival Analysis Tool for Esophageal Adenocarcinoma. <i>Frontiers in Oncology</i> , 2020, 10, 315.	2.8	7
34	High TXLNA Expression Predicts Favourable Outcome for Pancreatic Adenocarcinoma Patients. <i>BioMed Research International</i> , 2020, 2020, 1-9.	1.9	6
35	Comprehensive Review of Web Servers and Bioinformatics Tools for Cancer Prognosis Analysis. <i>Frontiers in Oncology</i> , 2020, 10, 68.	2.8	76
36	Systematic Comparison of Two Animal-to-Human Transmitted Human Coronaviruses: SARS-CoV-2 and SARS-CoV. <i>Viruses</i> , 2020, 12, 244.	3.3	552

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37	Low preoperative serum ALB level is independently associated with poor overall survival in endometrial cancer patients. <i>Future Oncology</i> , 2020, 16, 307-316.	2.4	20
38	Cover Image, Volume 59, Issue 3. <i>Molecular Carcinogenesis</i> , 2020, 59, i.	2.7	0
39	Systemic analysis of the expression and prognostic significance of PAKs in breast cancer. <i>Genomics</i> , 2020, 112, 2433-2444.	2.9	13
40	OSluca: An Interactive Web Server to Evaluate Prognostic Biomarkers for Lung Cancer. <i>Frontiers in Genetics</i> , 2020, 11, 420.	2.3	25
41	Blood transcriptome and clonal T-cell correlates of response and non-response to eltrombopag therapy in a cohort of patients with chronic immune thrombocytopenia. <i>Haematologica</i> , 2020, 105, e129-e132.	3.5	11
42	Autophagy-related gene P4HB: a novel diagnosis and prognosis marker for kidney renal clear cell carcinoma. <i>Aging</i> , 2020, 12, 1828-1842.	3.1	57
43	Genomic, epigenomic, and immune subtype analysis of CTSL/B and SARS-CoV-2 receptor ACE2 in pan-cancer. <i>Aging</i> , 2020, 12, 22370-22389.	3.1	12
44	Preparation and preliminary evaluation of hepatitis B core antigen virus like nanoparticles loaded with indocyanine green. <i>Annals of Translational Medicine</i> , 2020, 8, 1661-1661.	1.7	3
45	Interactive online consensus survival tool for esophageal squamous cell carcinoma prognosis analysis. <i>Oncology Letters</i> , 2019, 18, 1199-1206.	1.8	51
46	<p>Toxicity and serum metabolomics investigation of Mn-doped ZnS quantum dots in mice</p>. <i>International Journal of Nanomedicine</i> , 2019, Volume 14, 6297-6311.	6.7	10
47	OSkirc: a web tool for identifying prognostic biomarkers in kidney renal clear cell carcinoma. <i>Future Oncology</i> , 2019, 15, 3103-3110.	2.4	63
48	Novel Model for Comprehensive Assessment of Robust Prognostic Gene Signature in Ovarian Cancer Across Different Independent Datasets. <i>Frontiers in Genetics</i> , 2019, 10, 931.	2.3	2
49	OScc: an online survival analysis web server to evaluate the prognostic value of biomarkers in cervical cancer. <i>Future Oncology</i> , 2019, 15, 3693-3699.	2.4	22
50	OSblca: A Web Server for Investigating Prognostic Biomarkers of Bladder Cancer Patients. <i>Frontiers in Oncology</i> , 2019, 9, 466.	2.8	55
51	Gene Expression Profiling Reveals Distinct Molecular Subtypes of Esophageal Squamous Cell Carcinoma in Asian Populations. <i>Neoplasia</i> , 2019, 21, 571-581.	5.3	21
52	OSlms: A Web Server to Evaluate the Prognostic Value of Genes in Leiomyosarcoma. <i>Frontiers in Oncology</i> , 2019, 9, 190.	2.8	51
53	High FAM83A Expression is a Novel Prognostic Indicator of Poor Overall Survival in Lung Adenocarcinoma. , 2019, , .		1
54	OSbrca: A Web Server for Breast Cancer Prognostic Biomarker Investigation With Massive Data From Tens of Cohorts. <i>Frontiers in Oncology</i> , 2019, 9, 1349.	2.8	28

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55	<p>OSacc: Gene Expression-Based Survival Analysis Web Tool For Adrenocortical Carcinoma</p>. Cancer Management and Research, 2019, Volume 11, 9145-9152.	1.9	33
56	High KRT8 Expression Independently Predicts Poor Prognosis for Lung Adenocarcinoma Patients. Genes, 2019, 10, 36.	2.4	49
57	A novel approach for drug response prediction in cancer cell lines via network representation learning. Bioinformatics, 2019, 35, 1527-1535.	4.1	58
58	OSgbm: An Online Consensus Survival Analysis Web Server for Glioblastoma. Frontiers in Genetics, 2019, 10, 1378.	2.3	21
59	A quantitative method for measuring the transfection efficiency of CD19-directed chimeric antigen receptor in target cells. Advances in Clinical and Experimental Medicine, 2019, 28, 159-164.	1.4	0
60	High Sensitive and Non-invasive ctDNAs Sequencing Facilitate Clinical Diagnosis And Clinical Guidance of Non-small Cell Lung Cancer Patient: A Time Course Study. Frontiers in Oncology, 2018, 8, 491.	2.8	5
61	The Application of Sandwich Method in Experimental Teaching of Molecular Biology Based on qPCR Technology. , 2018, , .		0
62	Distinct esophageal adenocarcinoma molecular subtype has subtype-specific gene expression and mutation patterns. BMC Genomics, 2018, 19, 769.	2.8	8
63	Regulatory Role of N ⁶ -methyladenosine (m ⁶ A) Methylation in RNA Processing and Human Diseases. Journal of Cellular Biochemistry, 2017, 118, 2534-2543.	2.6	127
64	Exomic and Epigenomic Analyses in a Pair of Monozygotic Twins Discordant for Cryptorchidism. Twin Research and Human Genetics, 2017, 20, 349-354.	0.6	8
65	Expression of <scp>PTEN</scp>â€long mediated by <scp>CRISPR</scp>/Cas9 can repress U87 cell proliferation. Journal of Cellular and Molecular Medicine, 2017, 21, 3337-3346.	3.6	6
66	Distinct molecular subtypes of uterine leiomyosarcoma respond differently to chemotherapy treatment. BMC Cancer, 2017, 17, 639.	2.6	19
67	Identification of distinct molecular subtypes of uterine carcinosarcoma. Oncotarget, 2017, 8, 15878-15886.	1.8	9
68	CDX2 as a Prognostic Biomarker in Stage II and Stage III Colon Cancer. New England Journal of Medicine, 2016, 374, 211-222.	27.0	388
69	Differential expression of miRNAs related to caste differentiation in the honey bee, Apis mellifera. Apidologie, 2016, 47, 495-508.	2.0	18
70	Progressive loss of myogenic differentiation in leiomyosarcoma has prognostic value. Histopathology, 2015, 66, 627-638.	2.9	29
71	Clinically Relevant Molecular Subtypes in Leiomyosarcoma. Clinical Cancer Research, 2015, 21, 3501-3511.	7.0	129
72	Molecular subtyping of leiomyosarcoma with 3â€² end RNA sequencing. Genomics Data, 2015, 5, 366-367.	1.3	6

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73	Molecular pathological analysis of sarcomas using paraffin-embedded tissue: current limitations and future possibilities. <i>Histopathology</i> , 2014, 64, 163-170.	2.9	17
74	Dystrophin is a tumor suppressor in human cancers with myogenic programs. <i>Nature Genetics</i> , 2014, 46, 601-606.	21.4	142
75	A shared transcriptional program in early breast neoplasias despite genetic and clinical distinctions. <i>Genome Biology</i> , 2014, 15, R71.	9.6	30
76	Genome evolution during progression to breast cancer. <i>Genome Research</i> , 2013, 23, 1097-1108.	5.5	98
77	Next generation sequencing-based expression profiling identifies signatures from benign stromal proliferations that define stromal components of breast cancer. <i>Breast Cancer Research</i> , 2013, 15, R117.	5.0	14
78	Recipe for a Busy Bee: MicroRNAs in Honey Bee Caste Determination. <i>PLoS ONE</i> , 2013, 8, e81661.	2.5	60
79	14-3-3 fusion oncogenes in high-grade endometrial stromal sarcoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 929-934.	7.1	239
80	The novel long non-coding RNA CRC regulates <i>Drosophila</i> locomotor behavior. <i>Nucleic Acids Research</i> , 2012, 40, 11714-11727.	14.5	53
81	Cyclin D1 as a Diagnostic Immunomarker for Endometrial Stromal Sarcoma With YWHAE-FAM22 Rearrangement. <i>American Journal of Surgical Pathology</i> , 2012, 36, 1562-1570.	3.7	184
82	Transcriptional profiling of long non-coding RNAs and novel transcribed regions across a diverse panel of archived human cancers. <i>Genome Biology</i> , 2012, 13, R75.	9.6	221
83	Experimental RNomics and genomic comparative analysis reveal a large group of species-specific small non-message RNAs in the silkworm <i>Bombyx mori</i> . <i>Nucleic Acids Research</i> , 2011, 39, 3792-3805.	14.5	19