

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	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
2	CDX2 as a Prognostic Biomarker in Stage II and Stage III Colon Cancer. <i>New England Journal of Medicine</i> , 2016, 374, 211-222.	27.0	388
3	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
4	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
5	The Application of Deep Learning in Cancer Prognosis Prediction. <i>Cancers</i> , 2020, 12, 603.	3.7	218
6	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
7	Dystrophin is a tumor suppressor in human cancers with myogenic programs. <i>Nature Genetics</i> , 2014, 46, 601-606.	21.4	142
8	Clinically Relevant Molecular Subtypes in Leiomyosarcoma. <i>Clinical Cancer Research</i> , 2015, 21, 3501-3511.	7.0	129
9	Regulatory Role of N ⁶ -methyladenosine (m ⁶ A) Methylation in RNA Processing and Human Diseases. <i>Journal of Cellular Biochemistry</i> , 2017, 118, 2534-2543.	2.6	127
10	Genome evolution during progression to breast cancer. <i>Genome Research</i> , 2013, 23, 1097-1108.	5.5	98
11	Comprehensive Review of Web Servers and Bioinformatics Tools for Cancer Prognosis Analysis. <i>Frontiers in Oncology</i> , 2020, 10, 68.	2.8	76
12	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
13	Recipe for a Busy Bee: MicroRNAs in Honey Bee Caste Determination. <i>PLoS ONE</i> , 2013, 8, e81661.	2.5	60
14	A novel approach for drug response prediction in cancer cell lines via network representation learning. <i>Bioinformatics</i> , 2019, 35, 1527-1535.	4.1	58
15	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
16	OSblca: A Web Server for Investigating Prognostic Biomarkers of Bladder Cancer Patients. <i>Frontiers in Oncology</i> , 2019, 9, 466.	2.8	55
17	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
18	Interactive online consensus survival tool for esophageal squamous cell carcinoma prognosis analysis. <i>Oncology Letters</i> , 2019, 18, 1199-1206.	1.8	51

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19	OSlms: A Web Server to Evaluate the Prognostic Value of Genes in Leiomyosarcoma. <i>Frontiers in Oncology</i> , 2019, 9, 190.	2.8	51
20	High KRT8 Expression Independently Predicts Poor Prognosis for Lung Adenocarcinoma Patients. <i>Genes</i> , 2019, 10, 36.	2.4	49
21	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
22	<p>OSacc: Gene Expression-Based Survival Analysis Web Tool For Adrenocortical Carcinoma</p>. <i>Cancer Management and Research</i> , 2019, Volume 11, 9145-9152.	1.9	33
23	A shared transcriptional program in early breast neoplasias despite genetic and clinical distinctions. <i>Genome Biology</i> , 2014, 15, R71.	9.6	30
24	Progressive loss of myogenic differentiation in leiomyosarcoma has prognostic value. <i>Histopathology</i> , 2015, 66, 627-638.	2.9	29
25	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
26	The translational values of TRIM family in pan-cancers: From functions and mechanisms to clinics. , 2021, 227, 107881.		27
27	OSluca: An Interactive Web Server to Evaluate Prognostic Biomarkers for Lung Cancer. <i>Frontiers in Genetics</i> , 2020, 11, 420.	2.3	25
28	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
29	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
30	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
31	OSgbm: An Online Consensus Survival Analysis Web Server for Glioblastoma. <i>Frontiers in Genetics</i> , 2019, 10, 1378.	2.3	21
32	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
33	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
34	Distinct molecular subtypes of uterine leiomyosarcoma respond differently to chemotherapy treatment. <i>BMC Cancer</i> , 2017, 17, 639.	2.6	19
35	Differential expression of miRNAs related to caste differentiation in the honey bee, <i>Apis mellifera</i> . <i>Apidologie</i> , 2016, 47, 495-508.	2.0	18
36	OSumv: An interactive online consensus survival tool for uveal melanoma prognosis analysis. <i>Molecular Carcinogenesis</i> , 2020, 59, 56-61.	2.7	18

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37	Molecular pathological analysis of sarcomas using paraffin-embedded tissue: current limitations and future possibilities. <i>Histopathology</i> , 2014, 64, 163-170.	2.9	17
38	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
39	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
40	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
41	OSlihc: An Online Prognostic Biomarker Analysis Tool for Hepatocellular Carcinoma. <i>Frontiers in Pharmacology</i> , 2020, 11, 875.	3.5	15
42	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
43	Systematic Review of Prognostic Gene Signature in Gastric Cancer Patients. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020, 8, 805.	4.1	13
44	Systemic analysis of the expression and prognostic significance of PAKs in breast cancer. <i>Genomics</i> , 2020, 112, 2433-2444.	2.9	13
45	The Identification of RNA Modification Gene PUS7 as a Potential Biomarker of Ovarian Cancer. <i>Biology</i> , 2021, 10, 1130.	2.8	13
46	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
47	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
48	<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
49	OSdlbcl: 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
50	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
51	Identification of distinct molecular subtypes of uterine carcinosarcoma. <i>Oncotarget</i> , 2017, 8, 15878-15886.	1.8	9
52	Exomic and Epigenomic Analyses in a Pair of Monozygotic Twins Discordant for Cryptorchidism. <i>Twin Research and Human Genetics</i> , 2017, 20, 349-354.	0.6	8
53	Distinct esophageal adenocarcinoma molecular subtype has subtype-specific gene expression and mutation patterns. <i>BMC Genomics</i> , 2018, 19, 769.	2.8	8
54	OSeac: An Online Survival Analysis Tool for Esophageal Adenocarcinoma. <i>Frontiers in Oncology</i> , 2020, 10, 315.	2.8	7

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55	The Abnormal Glycopatterns of Salivary Glycoproteins in Esophageal Squamous Cell Carcinoma Patients. <i>Frontiers in Chemistry</i> , 2021, 9, 637730.	3.6	7
56	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
57	Molecular subtyping of leiomyosarcoma with 3â€² end RNA sequencing. <i>Genomics Data</i> , 2015, 5, 366-367.	1.3	6
58	Expression of <sc>PTEN</sc>â€”long mediated by <sc>CRISPR</sc>/Cas9 can repress U87 cell proliferation. <i>Journal of Cellular and Molecular Medicine</i> , 2017, 21, 3337-3346.	3.6	6
59	OSlgg: An Online Prognostic Biomarker Analysis Tool for Low-Grade Glioma. <i>Frontiers in Oncology</i> , 2020, 10, 1097.	2.8	6
60	High TXLNA Expression Predicts Favourable Outcome for Pancreatic Adenocarcinoma Patients. <i>BioMed Research International</i> , 2020, 2020, 1-9.	1.9	6
61	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
62	High Sensitive and Non-invasive ctDNAs Sequencing Facilitate Clinical Diagnosis And Clinical Guidance of Non-small Cell Lung Cancer Patient: A Time Course Study. <i>Frontiers in Oncology</i> , 2018, 8, 491.	2.8	5
63	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
64	OSucs: An Online Prognostic Biomarker Analysis Tool for Uterine Carcinosarcoma. <i>Genes</i> , 2020, 11, 1040.	2.4	4
65	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
66	OSmfs: An Online Interactive Tool to Evaluate Prognostic Markers for Myxofibrosarcoma. <i>Genes</i> , 2020, 11, 1523.	2.4	3
67	OSchol: an online consensus survival web server for cholangiocarcinoma prognosis analysis. <i>Hpb</i> , 2021, 23, 545-550.	0.3	3
68	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
69	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
70	Editorial: Bioinformatics Tools (and Web Server) for Cancer Biomarker Development. <i>Frontiers in Oncology</i> , 2020, 10, 599085.	2.8	2
71	Identification of Clinical Relevant Molecular Subtypes of Pheochromocytoma. <i>Frontiers in Endocrinology</i> , 2021, 12, 605797.	3.5	2
72	Editorial: Big Data and Machine Learning in Cancer Genomics. <i>Frontiers in Genetics</i> , 2021, 12, 749584.	2.3	2

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73	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
74	High FAM83A Expression is a Novel Prognostic Indicator of Poor Overall Survival in Lung Adenocarcinoma. , 2019, , .		1
75	Risk factors for in-hospital mortality in patients with cancer and COVID-19. <i>Lancet Oncology</i> , The, 2020, 21, e407.	10.7	1
76	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
77	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
78	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
79	The Application of Sandwich Method in Experimental Teaching of Molecular Biology Based on qPCR Technology. , 2018, , .		0
80	Cover Image, Volume 59, Issue 3. <i>Molecular Carcinogenesis</i> , 2020, 59, i.	2.7	0
81	Identification of Distinct Molecular Subtypes of Endometrioid Adenocarcinoma. <i>Frontiers in Genetics</i> , 2021, 12, 568779.	2.3	0
82	A quantitative method for measuring the transfection efficiency of CD19-directed chimeric antigen receptor in target cells. <i>Advances in Clinical and Experimental Medicine</i> , 2019, 28, 159-164.	1.4	0
83	OSov: An Interactive Web Server to Evaluate Prognostic Biomarkers for Ovarian Cancer. <i>Biology</i> , 2022, 11, 23.	2.8	0