

Yutao Wang

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

306
citations

1039406

9
h-index

1058022

14
g-index

20
all docs

20
docs citations

20
times ranked

292
citing authors

#	ARTICLE	IF	CITATIONS
1	A novel defined risk signature based on pyroptosis-related genes can predict the prognosis of prostate cancer. <i>BMC Medical Genomics</i> , 2022, 15, 24.	0.7	13
2	Macrophage M2 Co-expression Factors Correlate With the Immune Microenvironment and Predict Outcome of Renal Clear Cell Carcinoma. <i>Frontiers in Genetics</i> , 2021, 12, 615655.	1.1	31
3	M2 Macrophage Co-Expression Factors Correlate With Immune Phenotype and Predict Prognosis of Bladder Cancer. <i>Frontiers in Oncology</i> , 2021, 11, 609334.	1.3	19
4	Coexpressed Genes That Promote the Infiltration of M2 Macrophages in Melanoma Can Evaluate the Prognosis and Immunotherapy Outcome. <i>Journal of Immunology Research</i> , 2021, 2021, 1-15.	0.9	15
5	Gene Instability-Related lncRNA Prognostic Model of Melanoma Patients via Machine Learning Strategy. <i>Journal of Oncology</i> , 2021, 2021, 1-22.	0.6	8
6	Genome instability-related long non-coding RNA in clear renal cell carcinoma determined using computational biology. <i>BMC Cancer</i> , 2021, 21, 727.	1.1	12
7	9-Gene Signature Correlated With CD8+ T Cell Infiltration Activated by IFN- γ : A Biomarker of Immune Checkpoint Therapy Response in Melanoma. <i>Frontiers in Immunology</i> , 2021, 12, 622563.	2.2	26
8	Wnt pathway-related three-mRNA clinical outcome signature in bladder urothelial carcinoma: computational biology and experimental analyses. <i>Journal of Translational Medicine</i> , 2021, 19, 409.	1.8	5
9	Identification of a Prognosis-Related Risk Signature for Bladder Cancer to Predict Survival and Immune Landscapes. <i>Journal of Immunology Research</i> , 2021, 2021, 1-26.	0.9	14
10	RNA m6A Methylation Regulators Multi-Omics Analysis in Prostate Cancer. <i>Frontiers in Genetics</i> , 2021, 12, 768041.	1.1	10
11	Treatment of Poor Sperm Quality and Erectile Dysfunction With Oral Pentoxifylline: A Systematic Review. <i>Frontiers in Pharmacology</i> , 2021, 12, 789787.	1.6	8
12	A robust 11-genes prognostic model can predict overall survival in bladder cancer patients based on five cohorts. <i>Cancer Cell International</i> , 2020, 20, 402.	1.8	14
13	CD8+ T Cell Co-Expressed Genes Correlate With Clinical Phenotype and Microenvironments of Urothelial Cancer. <i>Frontiers in Oncology</i> , 2020, 10, 553399.	1.3	26
14	Identification of a Robust Five-Gene Risk Model in Prostate Cancer: A Robust Likelihood-Based Survival Analysis. <i>International Journal of Genomics</i> , 2020, 2020, 1-23.	0.8	15
15	Identification of biomarkers related to CD8+ T cell infiltration with gene co-expression network in clear cell renal cell carcinoma. <i>Aging</i> , 2020, 12, 3694-3712.	1.4	51
16	Three-gene risk model in papillary renal cell carcinoma: a robust likelihood-based survival analysis. <i>Aging</i> , 2020, 12, 21854-21873.	1.4	19
17	Identification of core genes associated with prostate cancer progression and outcome via bioinformatics analysis in multiple databases. <i>PeerJ</i> , 2020, 8, e8786.	0.9	20