

Na Liu

List of Publications by Year in descending order

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78
papers

4,203
citations

147566

31
h-index

123241

61
g-index

85
all docs

85
docs citations

85
times ranked

4903
citing authors

#	ARTICLE	IF	CITATIONS
1	WTAP-mediated m6A modification of lncRNA DIAPH1-AS1 enhances its stability to facilitate nasopharyngeal carcinoma growth and metastasis. <i>Cell Death and Differentiation</i> , 2022, 29, 1137-1151.	5.0	66
2	USP44 regulates irradiation-induced DNA double-strand break repair and suppresses tumorigenesis in nasopharyngeal carcinoma. <i>Nature Communications</i> , 2022, 13, 501.	5.8	32
3	Expression Profiles and Prognostic Value of Multiple Inhibitory Checkpoints in Head and Neck Lymphoepithelioma-Like Carcinoma. <i>Frontiers in Immunology</i> , 2022, 13, 818411.	2.2	1
4	The immune modulation effects of gemcitabine plus cisplatin induction chemotherapy in nasopharyngeal carcinoma. <i>Cancer Medicine</i> , 2022, , .	1.3	3
5	Protein C receptor maintains cancer stem cell properties via activating lipid synthesis in nasopharyngeal carcinoma. <i>Signal Transduction and Targeted Therapy</i> , 2022, 7, 46.	7.1	9
6	A lncRNA signature associated with tumor immune heterogeneity predicts distant metastasis in locoregionally advanced nasopharyngeal carcinoma. <i>Nature Communications</i> , 2022, 13, .	5.8	31
7	Final Overall Survival Analysis of Gemcitabine and Cisplatin Induction Chemotherapy in Nasopharyngeal Carcinoma: A Multicenter, Randomized Phase III Trial. <i>Journal of Clinical Oncology</i> , 2022, 40, 2420-2425.	0.8	44
8	Association of Intratumoral Microbiota With Prognosis in Patients With Nasopharyngeal Carcinoma From 2 Hospitals in China. <i>JAMA Oncology</i> , 2022, 8, 1301.	3.4	44
9	Multi-Omics Integration Reveals the Crucial Role of <i>Fusobacterium</i> in the Inflammatory Immune Microenvironment in Head and Neck Squamous Cell Carcinoma. <i>Microbiology Spectrum</i> , 2022, 10, .	1.2	9
10	A Gene-Expression Predictor for Efficacy of Induction Chemotherapy in Locoregionally Advanced Nasopharyngeal Carcinoma. <i>Journal of the National Cancer Institute</i> , 2021, 113, 471-480.	3.0	17
11	Unraveling tumour microenvironment heterogeneity in nasopharyngeal carcinoma identifies biologically distinct immune subtypes predicting prognosis and immunotherapy responses. <i>Molecular Cancer</i> , 2021, 20, 14.	7.9	48
12	Prognostic Value of Pretreatment Serum Cystatin C Level in Nasopharyngeal Carcinoma Patients in the Intensity-modulated Radiotherapy Era. <i>OncoTargets and Therapy</i> , 2021, Volume 14, 29-37.	1.0	1
13	Chemotherapeutic and targeted agents can modulate the tumor microenvironment and increase the efficacy of immune checkpoint blockades. <i>Molecular Cancer</i> , 2021, 20, 27.	7.9	54
14	Gemcitabine synergizes with cisplatin to inhibit nasopharyngeal carcinoma cell proliferation and tumor growth. <i>FASEB Journal</i> , 2021, 35, e21885.	0.2	4
15	A gene expression-based immune content predictor for survival and postoperative radiotherapy response in head and neck cancer. <i>Molecular Therapy - Oncolytics</i> , 2021, 22, 380-387.	2.0	0
16	WIPI-1 inhibits metastasis and tumour growth via the WIPI-1-TRIM21 axis and MYC regulation in nasopharyngeal carcinoma. <i>Oral Oncology</i> , 2021, 122, 105576.	0.8	2
17	Spatial heterogeneity of immune infiltration predicts the prognosis of nasopharyngeal carcinoma patients. <i>Oncolimmunology</i> , 2021, 10, 1976439.	2.1	8
18	Prognostic Implication of Metabolic Syndrome in Patients with Nasopharyngeal Carcinoma: A Large Institution-Based Cohort Study from an Endemic Area. <i>Cancer Management and Research</i> , 2021, Volume 13, 9355-9366.	0.9	1

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19	<i>ZNF582</i> hypermethylation promotes metastasis of nasopharyngeal carcinoma by regulating the transcription of adhesion molecules <i>Nectin3</i> and <i>NRXN3</i> . <i>Cancer Communications</i> , 2020, 40, 721-737.	3.7	18
20	Long Noncoding RNA TINCR-Mediated Regulation of Acetyl-CoA Metabolism Promotes Nasopharyngeal Carcinoma Progression and Chemoresistance. <i>Cancer Research</i> , 2020, 80, 5174-5188.	0.4	63
21	Single-cell transcriptomics reveals regulators underlying immune cell diversity and immune subtypes associated with prognosis in nasopharyngeal carcinoma. <i>Cell Research</i> , 2020, 30, 1024-1042.	5.7	182
22	AR-induced long non-coding RNA LINC01503 facilitates proliferation and metastasis via the SFPQ-FOSL1 axis in nasopharyngeal carcinoma. <i>Oncogene</i> , 2020, 39, 5616-5632.	2.6	24
23	Prognostic value of immune score in nasopharyngeal carcinoma using digital pathology. , 2020, 8, e000334.		21
24	Plasma protein-based signature predicts distant metastasis and induction chemotherapy benefit in Nasopharyngeal Carcinoma. <i>Theranostics</i> , 2020, 10, 9767-9778.	4.6	14
25	Serum Calcium Levels Before Antitumour Therapy Predict Clinical Outcomes in Patients with Nasopharyngeal Carcinoma. <i>OncoTargets and Therapy</i> , 2020, Volume 13, 13111-13119.	1.0	6
26	Hypermethylation of UCHL1 Promotes Metastasis of Nasopharyngeal Carcinoma by Suppressing Degradation of Cortactin (CTTN). <i>Cells</i> , 2020, 9, 559.	1.8	23
27	Circular RNA CRIM1 functions as a ceRNA to promote nasopharyngeal carcinoma metastasis and docetaxel chemoresistance through upregulating FOXQ1. <i>Molecular Cancer</i> , 2020, 19, 33.	7.9	128
28	TRIM21-SERPINB5 aids GMPs repression to protect nasopharyngeal carcinoma cells from radiation-induced apoptosis. <i>Journal of Biomedical Science</i> , 2020, 27, 30.	2.6	30
29	<i>FNDC3B</i> 3' UTR shortening escapes from microRNA-mediated gene repression and promotes nasopharyngeal carcinoma progression. <i>Cancer Science</i> , 2020, 111, 1991-2003.	1.7	22
30	Higher vs. Lower DP for Ventilated Patients with Acute Respiratory Distress Syndrome: A Systematic Review and Meta-Analysis. <i>Emergency Medicine International</i> , 2019, 2019, 1-12.	0.3	5
31	Long Noncoding RNA FAM225A Promotes Nasopharyngeal Carcinoma Tumorigenesis and Metastasis by Acting as ceRNA to Sponge miR-590-3p/miR-1275 and Upregulate ITGB3. <i>Cancer Research</i> , 2019, 79, 4612-4626.	0.4	250
32	Astragaloligosides reduces lipopolysaccharide-induced acute lung injury in rats via induction of heme oxygenase-1. <i>Archives of Pharmacal Research</i> , 2019, 42, 704-711.	2.7	15
33	Gemcitabine and Cisplatin Induction Chemotherapy in Nasopharyngeal Carcinoma. <i>New England Journal of Medicine</i> , 2019, 381, 1124-1135.	13.9	573
34	NFAT1 Hypermethylation Promotes Epithelial-Mesenchymal Transition and Metastasis in Nasopharyngeal Carcinoma by Activating ITGA6 Transcription. <i>Neoplasia</i> , 2019, 21, 311-321.	2.3	18
35	Development and validation of an immune checkpoint-based signature to predict prognosis in nasopharyngeal carcinoma using computational pathology analysis. , 2019, 7, 298.		40
36	Hypermethylation of <i>SHISA3</i> Promotes Nasopharyngeal Carcinoma Metastasis by Reducing <i>SGSM1</i> Stability. <i>Cancer Research</i> , 2019, 79, 747-759.	0.4	35

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37	ARNTL hypermethylation promotes tumorigenesis and inhibits cisplatin sensitivity by activating CDK5 transcription in nasopharyngeal carcinoma. <i>Journal of Experimental and Clinical Cancer Research</i> , 2019, 38, 11.	3.5	41
38	EZH2-DNMT1-mediated epigenetic silencing of miR-142-3p promotes metastasis through targeting ZEB2 in nasopharyngeal carcinoma. <i>Cell Death and Differentiation</i> , 2019, 26, 1089-1106.	5.0	52
39	Prognostic significance of tumor-infiltrating lymphocytes in nondisseminated nasopharyngeal carcinoma: A large-scale cohort study. <i>International Journal of Cancer</i> , 2018, 142, 2558-2566.	2.3	73
40	Development and validation of a gene expression-based signature to predict distant metastasis in locoregionally advanced nasopharyngeal carcinoma: a retrospective, multicentre, cohort study. <i>Lancet Oncology</i> , The, 2018, 19, 382-393.	5.1	232
41	MicroRNA-101 inhibits invasion and angiogenesis through targeting ITGA3 and its systemic delivery inhibits lung metastasis in nasopharyngeal carcinoma. <i>Cell Death and Disease</i> , 2018, 8, e2566-e2566.	2.7	48
42	Long non-coding RNA DANCR stabilizes HIF-1 α and promotes metastasis by interacting with NF90/NF45 complex in nasopharyngeal carcinoma. <i>Theranostics</i> , 2018, 8, 5676-5689.	4.6	102
43	m6A-mediated ZNF750 repression facilitates nasopharyngeal carcinoma progression. <i>Cell Death and Disease</i> , 2018, 9, 1169.	2.7	83
44	Famitinib in combination with concurrent chemoradiotherapy in patients with locoregionally advanced nasopharyngeal carcinoma: a phase 1, open-label, dose-escalation Study. <i>Cancer Communications</i> , 2018, 38, 1-13.	3.7	20
45	Differential genome-wide profiling of alternative polyadenylation sites in nasopharyngeal carcinoma by high-throughput sequencing. <i>Journal of Biomedical Science</i> , 2018, 25, 74.	2.6	7
46	TIP3 hypermethylation correlates with worse prognosis and promotes tumor progression in nasopharyngeal carcinoma. <i>Journal of Experimental and Clinical Cancer Research</i> , 2018, 37, 227.	3.5	17
47	Molecular subtyping of nasopharyngeal carcinoma (NPC) and a microRNA-based prognostic model for distant metastasis. <i>Journal of Biomedical Science</i> , 2018, 25, 16.	2.6	38
48	<i>RAB37</i> Hypermethylation Regulates Metastasis and Resistance to Docetaxel-Based Induction Chemotherapy in Nasopharyngeal Carcinoma. <i>Clinical Cancer Research</i> , 2018, 24, 6495-6508.	3.2	25
49	HOPX hypermethylation promotes metastasis via activating SNAIL transcription in nasopharyngeal carcinoma. <i>Nature Communications</i> , 2017, 8, 14053.	5.8	95
50	The immune molecular landscape of the B7 and TNFR immunoregulatory ligand-receptor families in head and neck cancer: A comprehensive overview and the immunotherapeutic implications. <i>Oncotmmunology</i> , 2017, 6, e1288329.	2.1	16
51	Genomic Analysis of Tumor Microenvironment Immune Types across 14 Solid Cancer Types: Immunotherapeutic Implications. <i>Theranostics</i> , 2017, 7, 3585-3594.	4.6	214
52	Epigenetic mediated zinc finger protein 671 downregulation promotes cell proliferation and tumorigenicity in nasopharyngeal carcinoma by inhibiting cell cycle arrest. <i>Journal of Experimental and Clinical Cancer Research</i> , 2017, 36, 147.	3.5	23
53	Microarray Expression Profiling of Long Non-Coding RNAs Involved in Nasopharyngeal Carcinoma Metastasis. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1956.	1.8	31
54	Overexpression of Mitochondria Mediator Gene TRIAP1 by miR-320b Loss Is Associated with Progression in Nasopharyngeal Carcinoma. <i>PLoS Genetics</i> , 2016, 12, e1006183.	1.5	48

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55	Role of SFRP1 in NPC Metastasis Response. <i>Cancer Prevention Research</i> , 2016, 9, 416-416.	0.7	1
56	YPEL3 suppresses epithelial-mesenchymal transition and metastasis of nasopharyngeal carcinoma cells through the Wnt/ β -catenin signaling pathway. <i>Journal of Experimental and Clinical Cancer Research</i> , 2016, 35, 109.	3.5	41
57	miR-16 targets fibroblast growth factor 2 to inhibit NPC cell proliferation and invasion via PI3K/AKT and MAPK signaling pathways. <i>Oncotarget</i> , 2016, 7, 3047-3058.	0.8	52
58	CXCL12 genetic variants as prognostic markers in nasopharyngeal carcinoma. <i>OncoTargets and Therapy</i> , 2015, 8, 2835.	1.0	12
59	Efficacy of the Additional Neoadjuvant Chemotherapy to Concurrent Chemoradiotherapy for Patients with Locoregionally Advanced Nasopharyngeal Carcinoma: a Bayesian Network Meta-analysis of Randomized Controlled Trials. <i>Journal of Cancer</i> , 2015, 6, 883-892.	1.2	68
60	High expression of Talin-1 is associated with poor prognosis in patients with nasopharyngeal carcinoma. <i>BMC Cancer</i> , 2015, 15, 332.	1.1	21
61	Identification of miR-143 as a tumour suppressor in nasopharyngeal carcinoma based on microRNA expression profiling. <i>International Journal of Biochemistry and Cell Biology</i> , 2015, 61, 120-128.	1.2	30
62	MicroRNA-93 promotes cell growth and invasion in nasopharyngeal carcinoma by targeting disabled homolog-2. <i>Cancer Letters</i> , 2015, 363, 146-155.	3.2	54
63	Genome-Wide Identification of a Methylation Gene Panel as a Prognostic Biomarker in Nasopharyngeal Carcinoma. <i>Molecular Cancer Therapeutics</i> , 2015, 14, 2864-2873.	1.9	80
64	Low SFRP1 Expression Correlates with Poor Prognosis and Promotes Cell Invasion by Activating the Wnt/ β -Catenin Signaling Pathway in NPC. <i>Cancer Prevention Research</i> , 2015, 8, 968-977.	0.7	33
65	MiR-145 Inhibits Metastasis by Targeting Fascin Actin-Bundling Protein 1 in Nasopharyngeal Carcinoma. <i>PLoS ONE</i> , 2015, 10, e0122228.	1.1	24
66	Prognostic value of MET protein overexpression and gene amplification in locoregionally advanced nasopharyngeal carcinoma. <i>Oncotarget</i> , 2015, 6, 13309-13319.	0.8	19
67	Effect of latent membrane protein 1 expression on overall survival in Epstein-Barr virus-associated cancers: a literature-based meta-analysis. <i>Oncotarget</i> , 2015, 6, 29311-29323.	0.8	37
68	A four-miRNA signature identified from genome-wide serum miRNA profiling predicts survival in patients with nasopharyngeal carcinoma. <i>International Journal of Cancer</i> , 2014, 134, 1359-1368.	2.3	95
69	Overexpression of CIP2A is an independent prognostic indicator in nasopharyngeal carcinoma and its depletion suppresses cell proliferation and tumor growth. <i>Molecular Cancer</i> , 2014, 13, 111.	7.9	21
70	Hotspot mutations in common oncogenes are infrequent in nasopharyngeal carcinoma. <i>Oncology Reports</i> , 2014, 32, 1661-1669.	1.2	17
71	5-Azacytidine Enhances the Radiosensitivity of CNE2 and SUNE1 Cells In Vitro and In Vivo Possibly by Altering DNA Methylation. <i>PLoS ONE</i> , 2014, 9, e93273.	1.1	30
72	Outcomes in patients with non-ST-elevation acute coronary syndrome randomly assigned to invasive versus conservative treatment strategies: A meta-analysis. <i>Clinics</i> , 2014, 69, 398-404.	0.6	8

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73	MiR-451 inhibits cell growth and invasion by targeting MIF and is associated with survival in nasopharyngeal carcinoma. <i>Molecular Cancer</i> , 2013, 12, 123.	7.9	104
74	MiR-29c suppresses invasion and metastasis by targeting TIAM1 in nasopharyngeal carcinoma. <i>Cancer Letters</i> , 2013, 329, 181-188.	3.2	118
75	Reduced expression of Dicer11 is associated with poor prognosis in patients with nasopharyngeal carcinoma. <i>Medical Oncology</i> , 2013, 30, 360.	1.2	10
76	Prognostic value of a microRNA signature in nasopharyngeal carcinoma: a microRNA expression analysis. <i>Lancet Oncology</i> , The, 2012, 13, 633-641.	5.1	274
77	Low BRMS1 expression promotes nasopharyngeal carcinoma metastasis in vitro and in vivo and is associated with poor patient survival. <i>BMC Cancer</i> , 2012, 12, 376.	1.1	20
78	Nuclear overexpression of metastasis-associated protein 1 correlates significantly with poor survival in nasopharyngeal carcinoma. <i>Journal of Translational Medicine</i> , 2012, 10, 78.	1.8	22