

# Linlin Wang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/433978/publications.pdf>

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9  
papers

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citations

1307594

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1372567

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178  
citing authors

#	ARTICLE	IF	CITATIONS
1	Weighted gene expression profiles identify diagnostic and prognostic genes for lung adenocarcinoma and squamous cell carcinoma. <i>Journal of International Medical Research</i> , 2020, 48, 030006051989383.	1.0	7
2	Diagnostic Value of Combination of MicroRNA-192 in Urinary Sediment and B-Ultrasound for Bladder Cancer. <i>Technology in Cancer Research and Treatment</i> , 2020, 19, 153303381989457.	1.9	11
3	Integrated Analysis of Three Publicly Available Gene Expression Profiles Identified Genes and Pathways Associated with Clear Cell Renal Cell Carcinoma. <i>Medical Science Monitor</i> , 2020, 26, e919965.	1.1	2
4	Construction of subtype-specific prognostic gene signatures for early-stage non-small cell lung cancer using meta feature selection methods. <i>Oncology Letters</i> , 2019, 18, 2366-2375.	1.8	0
5	miR-5590-3p-CY1 feedback loop promotes the proliferation and migration of triple-negative breast cancer cells. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 18415-18424.	2.6	14
6	Chrysin ameliorates ANU-induced pulmonary edema and pulmonary arterial hypertension via modulation of VEGF and eNOs. <i>Journal of Biochemical and Molecular Toxicology</i> , 2019, 33, e22332.	3.0	20
7	miR-127 suppresses gastric cancer cell migration and invasion via targeting Wnt7a. <i>Oncology Letters</i> , 2019, 17, 3219-3226.	1.8	14
8	Classification of Non-Small Cell Lung Cancer Using Significance Analysis of Microarray-Gene Set Reduction Algorithm. <i>BioMed Research International</i> , 2016, 2016, 1-8.	1.9	14
9	Identification of Genes Discriminating Multiple Sclerosis Patients from Controls by Adapting a Pathway Analysis Method. <i>PLoS ONE</i> , 2016, 11, e0165543.	2.5	4