

# Minjie Wei

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

28  
papers

2,496  
citations

16  
h-index

31  
g-index

31  
ext. papers

3,635  
ext. citations

12.3  
avg, IF

4.62  
L-index

#	Paper	IF	Citations
28	Immune-Related Long Non-coding RNA Constructs a Prognostic Signature of Ovarian Cancer.. <i>Biological Procedures Online</i> , <b>2021</b> , 23, 24	8.3	1
27	N6-methyladenosine reader IMP2 stabilizes the ZFAS1/OLA1 axis and activates the Warburg effect: implication in colorectal cancer. <i>Journal of Hematology and Oncology</i> , <b>2021</b> , 14, 188	22.4	6
26	lncRNA-Xist/miR-101-3p/KLF6/C/EBP $\beta$ axis promotes TAM polarization to regulate cancer cell proliferation and migration. <i>Molecular Therapy - Nucleic Acids</i> , <b>2021</b> , 23, 536-551	10.7	36
25	Integrated microenvironment-associated genomic profiles identify LRRC15 mediating recurrent glioblastoma-associated macrophages infiltration. <i>Journal of Cellular and Molecular Medicine</i> , <b>2021</b> , 25, 5534-5546	5.6	3
24	NF-B-activated SPRY4-IT1 promotes cancer cell metastasis by downregulating TCEB1 mRNA via Stau1-mediated mRNA decay. <i>Oncogene</i> , <b>2021</b> , 40, 4919-4929	9.2	6
23	LncRNA CBR3-AS1 regulates of breast cancer drug sensitivity as a competing endogenous RNA through the JNK1/MEK4-mediated MAPK signal pathway. <i>Journal of Experimental and Clinical Cancer Research</i> , <b>2021</b> , 40, 41	12.8	8
22	Integrative Analysis of DNA Methylation and Gene Expression to Determine Specific Diagnostic Biomarkers and Prognostic Biomarkers of Breast Cancer. <i>Frontiers in Cell and Developmental Biology</i> , <b>2020</b> , 8, 529386	5.7	6
21	Analysis of immune subtypes based on immunogenomic profiling identifies prognostic signature for cutaneous melanoma. <i>International Immunopharmacology</i> , <b>2020</b> , 89, 107162	5.8	8
20	LNC942 promoting METTL14-mediated mA methylation in breast cancer cell proliferation and progression. <i>Oncogene</i> , <b>2020</b> , 39, 5358-5372	9.2	67
19	Targeting FTO Suppresses Cancer Stem Cell Maintenance and Immune Evasion. <i>Cancer Cell</i> , <b>2020</b> , 38, 79-96.e11	24.3	145
18	Expression signature of six-snoRNA serves as novel non-invasive biomarker for diagnosis and prognosis prediction of renal clear cell carcinoma. <i>Journal of Cellular and Molecular Medicine</i> , <b>2020</b> , 24, 2215-2228	5.6	17
17	Long noncoding RNA ZFAS1 promoting small nucleolar RNA-mediated 2H0-methylation via NOP58 recruitment in colorectal cancer. <i>Molecular Cancer</i> , <b>2020</b> , 19, 95	42.1	35
16	Bioinformatic profiling identifies a platinum-resistant-related risk signature for ovarian cancer. <i>Cancer Medicine</i> , <b>2020</b> , 9, 1242-1253	4.8	6
15	LncRNA SPRY4-IT1 regulates breast cancer cell stemness through competitively binding miR-6882-3p with TCF7L2. <i>Journal of Cellular and Molecular Medicine</i> , <b>2020</b> , 24, 772-784	5.6	31
14	Development of an IFN $\gamma$ response-related signature for predicting the survival of cutaneous melanoma. <i>Cancer Medicine</i> , <b>2020</b> , 9, 8186-8201	4.8	11
13	LncRNA HOTTIP facilitates the stemness of breast cancer via regulation of miR-148a-3p/WNT1 pathway. <i>Journal of Cellular and Molecular Medicine</i> , <b>2020</b> , 24, 6242-6252	5.6	18
12	A five-mRNA signature associated with post-translational modifications can better predict recurrence and survival in cervical cancer. <i>Journal of Cellular and Molecular Medicine</i> , <b>2020</b> , 24, 6283-6297	5.6	10

11	Intrinsic adriamycin resistance in p53-mutated breast cancer is related to the miR-30c/FANCF/REV1-mediated DNA damage response. <i>Cell Death and Disease</i> , <b>2019</b> , 10, 666	9.8	12
10	Long non-coding RNA LUCAT1/miR-5582-3p/TCF7L2 axis regulates breast cancer stemness via Wnt/βcatenin pathway. <i>Journal of Experimental and Clinical Cancer Research</i> , <b>2019</b> , 38, 305	12.8	61
9	Hypoxia-inducible factor-2β directly promotes BCRP expression and mediates the resistance of ovarian cancer stem cells to adriamycin. <i>Molecular Oncology</i> , <b>2019</b> , 13, 403-421	7.9	27
8	Recognition of RNA N-methyladenosine by IGF2BP proteins enhances mRNA stability and translation. <i>Nature Cell Biology</i> , <b>2018</b> , 20, 285-295	23.4	795
7	METTL14 Inhibits Hematopoietic Stem/Progenitor Differentiation and Promotes Leukemogenesis via mRNA mA Modification. <i>Cell Stem Cell</i> , <b>2018</b> , 22, 191-205.e9	18	476
6	R-2HG Exhibits Anti-tumor Activity by Targeting FTO/mA/MYC/CEBPA Signaling. <i>Cell</i> , <b>2018</b> , 172, 90-105.e33	23.3	479
5	High PITX1 expression in lung adenocarcinoma patients is associated with DNA methylation and poor prognosis. <i>Pathology Research and Practice</i> , <b>2018</b> , 214, 2046-2053	3.4	15
4	HIF-2β promotes conversion to a stem cell phenotype and induces chemoresistance in breast cancer cells by activating Wnt and Notch pathways. <i>Journal of Experimental and Clinical Cancer Research</i> , <b>2018</b> , 37, 256	12.8	77
3	miR-302a/b/c/d cooperatively inhibit BCRP expression to increase drug sensitivity in breast cancer cells. <i>Gynecologic Oncology</i> , <b>2016</b> , 141, 592-601	4.9	44
2	MiR-302a/b/c/d cooperatively sensitizes breast cancer cells to adriamycin via suppressing P-glycoprotein(P-gp) by targeting MAP/ERK kinase kinase 1 (MEKK1). <i>Journal of Experimental and Clinical Cancer Research</i> , <b>2016</b> , 35, 25	12.8	60
1	The Hedgehog signalling pathway mediates drug response of MCF-7 mammosphere cells in breast cancer patients. <i>Clinical Science</i> , <b>2015</b> , 129, 809-22	6.5	35