Weidong Zhou

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.

38 832 14 28 g-index

41 1,090 5.6 avg, IF L-index

#	Paper	IF	Citations
38	Extracellular vesicles from HTLV-1 infected cells modulate target cells and viral spread. <i>Retrovirology</i> , 2021 , 18, 6	3.6	10
37	Proteomic Discovery of VEEV E2-Host Partner Interactions Identifies GRP78 Inhibitor HA15 as a Potential Therapeutic for Alphavirus Infections. <i>Pathogens</i> , 2021 , 10,	4.5	3
36	Use of magnetic nanotrap particles in capturing Yersinia pestis virulence factors, nucleic acids and bacteria. <i>Journal of Nanobiotechnology</i> , 2021 , 19, 186	9.4	
35	Exosomes originating from infection with the cytoplasmic single-stranded RNA virus Rift Valley fever virus (RVFV) protect recipient cells by inducing RIG-I mediated IFN-B response that leads to activation of autophagy <i>Cell and Bioscience</i> , 2021 , 11, 220	9.8	0
34	Evaluation of pathogen specific urinary peptides in tick-borne illnesses. <i>Scientific Reports</i> , 2020 , 10, 193	40 9	2
33	Salivary proteomic profile of patients with renal cell carcinoma <i>Journal of Clinical Oncology</i> , 2020 , 38, 622-622	2.2	
32	Shotgun proteomics coupled to nanoparticle-based biomarker enrichment reveals a novel panel of extracellular matrix proteins as candidate serum protein biomarkers for early-stage breast cancer detection. <i>Breast Cancer Research</i> , 2020 , 22, 135	8.3	6
31	Venezuelan Equine Encephalitis Virus nsP3 Phosphorylation Can Be Mediated by IKK[Kinase Activity and Abrogation of Phosphorylation Inhibits Negative-Strand Synthesis. <i>Viruses</i> , 2020 , 12,	6.2	10
30	Stem Cell Extracellular Vesicles and their Potential to Contribute to the Repair of Damaged CNS Cells. <i>Journal of NeuroImmune Pharmacology</i> , 2020 , 15, 520-537	6.9	16
29	Tumor-Draining Lymph Secretome En Route to the Regional Lymph Node in Breast Cancer Metastasis. <i>Breast Cancer: Targets and Therapy</i> , 2020 , 12, 57-67	3.9	2
28	An Omics Approach to Extracellular Vesicles from HIV-1 Infected Cells. <i>Cells</i> , 2019 , 8,	7.9	9
27	FABP7 is a key metabolic regulator in HER2+ breast cancer brain metastasis. <i>Oncogene</i> , 2019 , 38, 6445-6	6 46 0	29
26	Dynamic Regulation of Caveolin-1 Phosphorylation and Caveolae Formation by Mammalian Target of Rapamycin Complex 2 in Bladder Cancer Cells. <i>American Journal of Pathology</i> , 2019 , 189, 1846-1862	5.8	7
25	Gasdermin D Exerts Anti-inflammatory Effects by Promoting Neutrophil Death. <i>Cell Reports</i> , 2018 , 22, 2924-2936	10.6	148
24	Phosphoproteomic Analysis Identifies Dynamic Regulation of Caveolin-1 Phosphorylation and Caveolae Formation by mTORC2 in Bladder Cancer Cells. <i>FASEB Journal</i> , 2018 , 32, 660.4	0.9	
23	An exploratory study examining how nano-liquid chromatography-mass spectrometry and phosphoproteomics can differentiate patients with advanced fibrosis and higher percentage collagen in non-alcoholic fatty liver disease. <i>BMC Medicine</i> , 2018 , 16, 170	11.4	5
22	Ebola Virus VP40 Modulates Cell Cycle and Biogenesis of Extracellular Vesicles. <i>Journal of Infectious Diseases</i> , 2018 , 218, S365-S387	7	28

(2012-2018)

21	Protein Phosphatase 1Interacts with Venezuelan Equine Encephalitis Virus Capsid Protein and Regulates Viral Replication through Modulation of Capsid Phosphorylation. <i>Journal of Virology</i> , 2018 , 92,	6.6	9	
20	Antiretroviral Drugs Alter the Content of Extracellular Vesicles from HIV-1-Infected Cells. <i>Scientific Reports</i> , 2018 , 8, 7653	4.9	43	
19	Mass Spectrometry-Based Biomarker Discovery. <i>Methods in Molecular Biology</i> , 2017 , 1606, 297-311	1.4	14	
18	Proteomic analysis of cerebrospinal fluid from children with central nervous system tumors identifies candidate proteins relating to tumor metastatic spread. <i>Oncotarget</i> , 2017 , 8, 46177-46190	3.3	15	
17	Positive Regulation of Interleukin-1 Bioactivity by Physiological ROS-Mediated Cysteine S-Glutathionylation. <i>Cell Reports</i> , 2017 , 20, 224-235	10.6	28	
16	The Warburg Effect and Mass Spectrometry-based Proteomic Analysis. <i>Cancer Genomics and Proteomics</i> , 2017 , 14, 211-218	3.3	12	
15	Targeting the Warburg effect in cancer cells through ENO1 knockdown rescues oxidative phosphorylation and induces growth arrest. <i>Oncotarget</i> , 2016 , 7, 5598-612	3.3	74	
14	Identification of novel candidate circulating biomarkers for malignant soft tissue sarcomas: Correlation with metastatic progression. <i>Proteomics</i> , 2016 , 16, 689-97	4.8	8	
13	Proteins that mediate protein aggregation and cytotoxicity distinguish Alzheimer hippocampus from normal controls. <i>Aging Cell</i> , 2016 , 15, 924-39	9.9	31	
12	Cancer metabolism and mass spectrometry-based proteomics. <i>Cancer Letters</i> , 2015 , 356, 176-83	9.9	25	
11	Nitric oxide as a regulator of B. anthracis pathogenicity. Frontiers in Microbiology, 2015, 6, 921	5.7	9	
10	Whole proteome analysis of mouse lymph nodes in cutaneous anthrax. <i>PLoS ONE</i> , 2014 , 9, e110873	3.7	9	
9	Mass spectrometric analysis reveals O-methylation of pyruvate kinase from pancreatic cancer cells. <i>Analytical and Bioanalytical Chemistry</i> , 2013 , 405, 4937-43	4.4	6	
8	Proteomic analysis reveals Warburg effect and anomalous metabolism of glutamine in pancreatic cancer cells. <i>Journal of Proteome Research</i> , 2012 , 11, 554-63	5.6	65	
7	MS analysis reveals O-methylation of L-lactate dehydrogenase from pancreatic ductal adenocarcinoma cells. <i>Electrophoresis</i> , 2012 , 33, 1850-4	3.6	11	
6	The spectra count label-free quantitation in cancer proteomics. <i>Cancer Genomics and Proteomics</i> , 2012 , 9, 135-42	3.3	14	
5	Cancer metabolism: what we can learn from proteomic analysis by mass spectrometry. <i>Cancer Genomics and Proteomics</i> , 2012 , 9, 373-81	3.3	9	
4	Mass spectrometry-based biomarker discovery. <i>Methods in Molecular Biology</i> , 2012 , 823, 251-64	1.4	4	

3	of Proteome Research, 2011 , 10, 1944-52	5.0	31	
2	Mass spectrometry analysis of the post-translational modifications of alpha-enolase from	5.6	58	

Proteomic analysis of pancreatic ductal adenocarcinoma cells reveals metabolic alterations. *Journal*

An initial characterization of the serum phosphoproteome. *Journal of Proteome Research*, **2009**, 8, 5523-3.6