

Chia-Wei Hsu

List of Publications by Year in descending order

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26
papers

1,434
citations

394421

19
h-index

552781

26
g-index

26
all docs

26
docs citations

26
times ranked

2724
citing authors

#	ARTICLE	IF	CITATIONS
1	Novel plasma metabolite markers of attention-deficit/hyperactivity disorder identified using high-performance chemical isotope labelling-based liquid chromatography-mass spectrometry. <i>World Journal of Biological Psychiatry</i> , 2021, 22, 139-148.	2.6	8
2	Association of exacerbation phenotype with the sputum microbiome in chronic obstructive pulmonary disease patients during the clinically stable state. <i>Journal of Translational Medicine</i> , 2021, 19, 121.	4.4	31
3	Reducing Lung ATP Levels and Alleviating Asthmatic Airway Inflammation through Adeno-Associated Viral Vector-Mediated CD39 Expression. <i>Biomedicines</i> , 2021, 9, 656.	3.2	4
4	Metabolomic profiling of parapneumonic effusion reveals a regulatory role of dipeptides in interleukin-8 production in neutrophil-like cells. <i>Analytica Chimica Acta</i> , 2020, 1128, 238-250.	5.4	2
5	Proteomic Profiling of Paired Interstitial Fluids Reveals Dysregulated Pathways and Salivary NID1 as a Biomarker of Oral Cavity Squamous Cell Carcinoma*[S]. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 1939-1949.	3.8	22
6	Identification of Salivary Biomarkers for Oral Cancer Detection with Untargeted and Targeted Quantitative Proteomics Approaches. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 1796-1806.	3.8	61
7	Integrated analyses utilizing metabolomics and transcriptomics reveal perturbation of the polyamine pathway in oral cavity squamous cell carcinoma. <i>Analytica Chimica Acta</i> , 2019, 1050, 113-122.	5.4	34
8	Metabolic reprogramming of hydrogenosomal amino acids in <i>Trichomonas vaginalis</i> under glucose restriction. <i>Journal of Microbiology, Immunology and Infection</i> , 2019, 52, 630-637.	3.1	11
9	Bretschneider solution-induced alterations in the urine metabolome in cardiac surgery patients. <i>Scientific Reports</i> , 2018, 8, 17774.	3.3	7
10	Oral Microbiota Community Dynamics Associated With Oral Squamous Cell Carcinoma Staging. <i>Frontiers in Microbiology</i> , 2018, 9, 862.	3.5	211
11	APOBEC3A is an oral cancer prognostic biomarker in Taiwanese carriers of an APOBEC deletion polymorphism. <i>Nature Communications</i> , 2017, 8, 465.	12.8	89
12	Proteomics Analysis Reveals Involvement of Krt17 in Areca Nut-Induced Oral Carcinogenesis. <i>Journal of Proteome Research</i> , 2016, 15, 2981-2997.	3.7	30
13	Identification and Characterization of Potential Biomarkers by Quantitative Tissue Proteomics of Primary Lung Adenocarcinoma. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 2396-2410.	3.8	65
14	Saliva proteome profiling reveals potential salivary biomarkers for detection of oral cavity squamous cell carcinoma. <i>Proteomics</i> , 2015, 15, 3394-3404.	2.2	86
15	In-depth Proteomic Analysis of Six Types of Exudative Pleural Effusions for Nonsmall Cell Lung Cancer Biomarker Discovery. <i>Molecular and Cellular Proteomics</i> , 2015, 14, 917-932.	3.8	32
16	Proteomics Analysis of EV71-Infected Cells Reveals the Involvement of Host Protein NEDD4L in EV71 Replication. <i>Journal of Proteome Research</i> , 2015, 14, 1818-1830.	3.7	24
17	Proteomic analyses of genes regulated by heterogeneous nuclear ribonucleoproteins <sc>A</sc>/<sc>B</sc> in <sc>J</sc>/<sc>urkat cells. <i>Proteomics</i> , 2014, 14, 1357-1366.	2.2	12
18	Secretome Profiling of Primary Cells Reveals That THBS2 Is a Salivary Biomarker of Oral Cavity Squamous Cell Carcinoma. <i>Journal of Proteome Research</i> , 2014, 13, 4796-4807.	3.7	50

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19	Low-molecular-mass secretome profiling identifies C α C motif chemokine 5 as a potential plasma biomarker and therapeutic target for nasopharyngeal carcinoma. <i>Journal of Proteomics</i> , 2013, 94, 186-201.	2.4	29
20	Proteome-wide Dysregulation by Glucose-6-phosphate Dehydrogenase (G6PD) Reveals a Novel Protective Role for G6PD in Aflatoxin B ₁ -Mediated Cytotoxicity. <i>Journal of Proteome Research</i> , 2013, 12, 3434-3448.	3.7	21
21	Interactome-wide Analysis Identifies End-binding Protein 1 as a Crucial Component for the Speck-like Particle Formation of Activated Absence in Melanoma 2 (AIM2) Inflammasomes. <i>Molecular and Cellular Proteomics</i> , 2012, 11, 1230-1244.	3.8	24
22	Comparative and Targeted Proteomic Analyses of Urinary Microparticles from Bladder Cancer and Hernia Patients. <i>Journal of Proteome Research</i> , 2012, 11, 5611-5629.	3.7	185
23	Identification of Guanylate-Binding Protein 1 as a Potential Oral Cancer Marker Involved in Cell Invasion Using Omics-Based Analysis. <i>Journal of Proteome Research</i> , 2011, 10, 3778-3788.	3.7	68
24	Quantitative plasma proteome analysis reveals aberrant level of blood coagulation-related proteins in nasopharyngeal carcinoma. <i>Journal of Proteomics</i> , 2011, 74, 744-757.	2.4	14
25	Candidate Serological Biomarkers for Cancer Identified from the Secretomes of 23 Cancer Cell Lines and the Human Protein Atlas. <i>Molecular and Cellular Proteomics</i> , 2010, 9, 1100-1117.	3.8	177
26	Discovery of Novel Bladder Cancer Biomarkers by Comparative Urine Proteomics Using iTRAQ Technology. <i>Journal of Proteome Research</i> , 2010, 9, 5803-5815.	3.7	137