## Jerome Zoidakis

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4577339/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Comprehensive human urine standards for comparability and standardization in clinical proteome analysis. Proteomics - Clinical Applications, 2010, 4, 464-478.	1.6	139
2	Expression and characterization of 1-aminocyclopropane-1-carboxylate deaminase from the rhizobacterium Pseudomonas putida UW4: a key enzyme in bacterial plant growth promotion. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2004, 1703, 11-19.	2.3	126
3	Profilin 1 is a Potential Biomarker for Bladder Cancer Aggressiveness. Molecular and Cellular Proteomics, 2012, 11, M111.009449.	3.8	97
4	Comparative Analysis of Label-Free and 8-Plex iTRAQ Approach for Quantitative Tissue Proteomic Analysis. PLoS ONE, 2015, 10, e0137048.	2.5	92
5	Development and Validation of Urine-based Peptide Biomarker Panels for Detecting Bladder Cancer in a Multi-center Study. Clinical Cancer Research, 2016, 22, 4077-4086.	7.0	90
6	Modular structure, local flexibility and cold-activity of a novel chitobiase from a psychrophilic antarctic bacterium. Journal of Molecular Biology, 2001, 310, 291-297.	4.2	86
7	An intrinsic role of IL-33 in Treg cell–mediated tumor immunoevasion. Nature Immunology, 2020, 21, 75-85.	14.5	82
8	Proteomics and Metabolomics for AKI Diagnosis. Seminars in Nephrology, 2018, 38, 63-87.	1.6	59
9	Plasma proteomic analysis reveals altered protein abundances in cardiovascular disease. Journal of Translational Medicine, 2018, 16, 104.	4.4	48
10	SRM/MRM targeted proteomics as a tool for biomarker validation and absolute quantification in human urine. Expert Review of Molecular Diagnostics, 2015, 15, 1441-1454.	3.1	46
11	Cervical Cancer Cell Line Secretome Highlights the Roles of Transforming Growth Factor-Beta-Induced Protein ig-h3, Peroxiredoxin-2, and NRF2 on Cervical Carcinogenesis. BioMed Research International, 2017, 2017, 1-15.	1.9	39
12	Comparison of Depletion Strategies for the Enrichment of Low-Abundance Proteins in Urine. PLoS ONE, 2015, 10, e0133773.	2.5	39
13	New Selective Peptidyl Di(chlorophenyl) Phosphonate Esters for Visualizing and Blocking Neutrophil Proteinase 3 in Human Diseases. Journal of Biological Chemistry, 2014, 289, 31777-31791.	3.4	38
14	Order of substrate binding in bacterial phenylalanine hydroxylase and its mechanistic implication for pterin-dependent oxygenases. Journal of Biological Inorganic Chemistry, 2003, 8, 121-128.	2.6	37
15	Proteomeâ€based classification of Nonmuscle Invasive Bladder Cancer. International Journal of Cancer, 2020, 146, 281-294.	5.1	35
16	Evaluation of Decellularization in Umbilical Cord Artery. Transplantation Proceedings, 2014, 46, 3232-3239.	0.6	30
17	Proteomic Analysis of Normal and Cancer Cervical Cell Lines Reveals Deregulation of Cytoskeleton-associated Proteins. Cancer Genomics and Proteomics, 2017, 14, 253-266.	2.0	30
18	Saliva Proteomics Analysis Offers Insights on Type 1 Diabetes Pathology in a Pediatric Population. Frontiers in Physiology, 2018, 9, 444.	2.8	30

JEROME ZOIDAKIS

#	Article	IF	CITATIONS
19	Urinary peptidomics analysis reveals proteases involved in diabetic nephropathy. Scientific Reports, 2017, 7, 15160.	3.3	28
20	NETs decorated with bioactive IL-33 infiltrate inflamed tissues and induce IFN-1 $\pm$ production in patients with SLE. JCI Insight, 2021, 6, .	5.0	28
21	Proteomics approaches in cervical cancer: focus on the discovery of biomarkers for diagnosis and drug treatment monitoring. Expert Review of Proteomics, 2016, 13, 731-745.	3.0	27
22	Proteomics analysis of bladder cancer invasion: Targeting EIF3D for therapeutic intervention. Oncotarget, 2017, 8, 69435-69455.	1.8	27
23	Analytical Performance of ELISA Assays in Urine: One More Bottleneck towards Biomarker Validation and Clinical Implementation. PLoS ONE, 2016, 11, e0149471.	2.5	27
24	Proteomics based identification of KDM5 histone demethylases associated with cardiovascular disease. EBioMedicine, 2019, 41, 91-104.	6.1	23
25	Microbiome in Chronic Kidney Disease (CKD): An Omics Perspective. Toxins, 2022, 14, 176.	3.4	22
26	Proteomics Analysis of Formalin Fixed Paraffin Embedded Tissues in the Investigation of Prostate Cancer. Journal of Proteome Research, 2020, 19, 2631-2642.	3.7	21
27	IMAC Fractionation in Combination with LC–MS Reveals H2B and NIF-1 Peptides As Potential Bladder Cancer Biomarkers. Journal of Proteome Research, 2013, 12, 3969-3979.	3.7	20
28	Proteomic advances in salivary diagnostics. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2020, 1868, 140494.	2.3	19
29	Silencing of Profilin-1 suppresses cell adhesion and tumor growth via predicted alterations in integrin and Ca2+ signaling in T24M-based bladder cancer models. Oncotarget, 2016, 7, 70750-70768.	1.8	19
30	Mass spectrometry-based membrane proteomics in cancer biomarker discovery. Expert Review of Molecular Diagnostics, 2014, 14, 549-563.	3.1	18
31	Cross-Talk Between Tumor Cells Undergoing Epithelial to Mesenchymal Transition and Natural Killer Cells in Tumor Microenvironment in Colorectal Cancer. Frontiers in Cell and Developmental Biology, 2021, 9, 750022.	3.7	18
32	Effect of temperature, pH, and metals on the stability and activity of phenylalanine hydroxylase from Chromobacterium violaceum. Journal of Inorganic Biochemistry, 2005, 99, 771-775.	3.5	17
33	Advances in urinary proteome analysis and applications in systems biology. Bioanalysis, 2014, 6, 2549-2569.	1.5	17
34	Proteome of olive non-glandular trichomes reveals protective protein network against (a)biotic challenge. Journal of Plant Physiology, 2018, 231, 210-218.	3.5	17
35	Multiplexed MRM-based protein quantification of putative prognostic biomarkers for chronic kidney disease progression in plasma. Scientific Reports, 2020, 10, 4815.	3.3	17
36	High Resolution Proteomic Analysis of the Cervical Cancer Cell Lines Secretome Documents Deregulation of Multiple Proteases. Cancer Genomics and Proteomics, 2017, 14, 507-521.	2.0	17

JEROME ZOIDAKIS

#	Article	IF	CITATIONS
37	Identification of novel molecular signatures of IgA nephropathy through an integrative -omics analysis. Scientific Reports, 2017, 7, 9091.	3.3	16
38	A systematic re-examination of processing of MHCI-bound antigenic peptide precursors by endoplasmic reticulum aminopeptidase 1. Journal of Biological Chemistry, 2020, 295, 7193-7210.	3.4	16
39	Applications of multiple reaction monitoring targeted proteomics assays in human plasma. Expert Review of Molecular Diagnostics, 2019, 19, 499-515.	3.1	15
40	Development and Validation of Multiple Reaction Monitoring (MRM) Assays for Clinical Applications. Methods in Molecular Biology, 2019, 1959, 205-223.	0.9	15
41	Analysis of urinary cathepsin C for diagnosing Papillon–LefÔvre syndrome. FEBS Journal, 2016, 283, 498-509.	4.7	14
42	Insights into Biomechanical and Proteomic Characteristics of Small Diameter Vascular Grafts Utilizing the Human Umbilical Artery. Biomedicines, 2020, 8, 280.	3.2	13
43	Protein Interactome of Muscle Invasive Bladder Cancer. PLoS ONE, 2015, 10, e0116404.	2.5	12
44	Integrative analysis of extracellular and intracellular bladder cancer cell line proteome with transcriptome: improving coverage and validity of –omics findings. Scientific Reports, 2016, 6, 25619.	3.3	12
45	Short Term Results of Fibrin Gel Obtained from Cord Blood Units: A Preliminary in Vitro Study. Bioengineering, 2019, 6, 66.	3.5	12
46	Evaluation of Peripheral Blood and Cord Blood Platelet Lysates in Isolation and Expansion of Multipotent Mesenchymal Stromal Cells. Bioengineering, 2018, 5, 19.	3.5	11
47	The family of 14â€3â€3 proteins and specifically 14â€3â€3σ are upâ€regulated during the development of renal pathologies. Journal of Cellular and Molecular Medicine, 2018, 22, 4139-4149.	3.6	10
48	Novel structural approaches concerning HPV proteins: Insight into targeted therapies for cervical cancer (Review). Oncology Reports, 2018, 39, 1547-1554.	2.6	10
49	Implications of the mitochondrial interactome of mammalian thioredoxin 2 for normal cellular function and disease. Free Radical Biology and Medicine, 2019, 137, 59-73.	2.9	10
50	Urinary Proteomics in Predicting Heart Transplantation Outcomes (uPROPHET)—Rationale and database description. PLoS ONE, 2017, 12, e0184443.	2.5	9
51	Chloride Intracellular Channel 4 Overexpression in the Proximal Tubules of Kidneys from the Spontaneously Hypertensive Rat: Insight from Proteomic Analysis. Nephron, 2018, 138, 60-70.	1.8	8
52	Tissue proteomics studies in the investigation of prostate cancer. Expert Review of Proteomics, 2018, 15, 593-611.	3.0	8
53	Downregulation of Salivary Proteins, Protective against Dental Caries, in Type 1 Diabetes. Proteomes, 2021, 9, 33.	3.5	8
54	PeptiCKDdb—peptide- and protein-centric database for the investigation of genesis and progression of chronic kidney disease. Database: the Journal of Biological Databases and Curation, 2016, 2016, baw128.	3.0	7

JEROME ZOIDAKIS

#	Article	IF	CITATIONS
55	Diagnostic and Prognostic Performance of Secreted Protein Acidic and Rich in Cysteine (SPARC) Assay for Detecting Primary and Recurrent Urinary Bladder Cancer. Proteomics - Clinical Applications, 2019, 13, 1800148.	1.6	7
56	Proteome analysis of leaf, stem and callus in Viscum album and identification of lectins and viscotoxins with bioactive properties. Plant Cell, Tissue and Organ Culture, 2020, 141, 167-178.	2.3	7
57	Role of the second coordination sphere residue tyrosine 179 in substrate affinity and catalytic activity of phenylalanine hydroxylase. Journal of Biological Inorganic Chemistry, 2004, 9, 289-296.	2.6	6
58	Membrane proteomics of cervical cancer cell lines reveal insights on the process of cervical carcinogenesis. International Journal of Oncology, 2018, 53, 2111-2122.	3.3	6
59	Protein biomarkers for cardiorenal syndrome. Expert Review of Proteomics, 2019, 16, 325-336.	3.0	6
60	Effect of Heme Oxygenase-1 Deficiency on Glomerular Proteomics. American Journal of Nephrology, 2016, 43, 441-450.	3.1	5
61	RGS14414-Mediated Activation of the 14-3-3ζ in Rodent Perirhinal Cortex Induces Dendritic Arborization, an Increase in Spine Number, Long-Lasting Memory Enhancement, and the Prevention of Memory Deficits. Cerebral Cortex, 2022, 32, 1894-1910.	2.9	5
62	BcCluster: A Bladder Cancer Database at the Molecular Level. Bladder Cancer, 2016, 2, 65-76.	0.4	4
63	Systematic review on recent potential biomarkers of chronic obstructive pulmonary disease. Expert Review of Molecular Diagnostics, 2019, 19, 37-45.	3.1	4
64	Proteomic Analysis of Mouse Kidney Tissue Associates Peroxisomal Dysfunction with Early Diabetic Kidney Disease. Biomedicines, 2022, 10, 216.	3.2	4
65	The ERAP1 active site cannot productively access the N-terminus of antigenic peptide precursors stably bound onto MHC class I. Scientific Reports, 2021, 11, 16475.	3.3	3
66	Gene Expression Monotonicity across Bladder Cancer Stages Informs on the Molecular Pathogenesis and Identifies a Prognostic Eight-Gene Signature. Cancers, 2022, 14, 2542.	3.7	3
67	Application of Preparative Electrophoresis for Clinical Proteomics in Urine: Is it Feasible?. Journal of Medical Biochemistry, 2009, 28, 268-273.	1.7	2
68	Simple and Efficient Stratification of Invasive Bladder Cancer Patients. EBioMedicine, 2016, 12, 6-7.	6.1	2
69	High resolution analysis of the intracellular proteome of cervical cancer cell lines unveils novel regulators of cervical carcinogenesis. Oncology Reports, 2019, 42, 1441-1450.	2.6	2
70	Plasma Proteomics in Healthy Subjects with Differences in Tissue Glucocorticoid Sensitivity Identifies A Novel Proteomic Signature. Biomedicines, 2022, 10, 184.	3.2	1
71	FP308URINARY PROTEOMICS TO DECIPHER MOLECULAR PATHOPHYSIOLOGY OF CKD PROGRESSION. Nephrology Dialysis Transplantation, 2015, 30, iii170-iii170.	0.7	0