Rebeca Kawahara

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2534464/publications.pdf

Version: 2024-02-01

36 papers 1,080 citations

430754 18 h-index 434063 31 g-index

41 all docs

41 docs citations

times ranked

41

1858 citing authors

#	Article	IF	CITATIONS
1	$\langle i > N < i > -$ acetyl- $\hat{\Gamma}^2$ -D-hexosaminidases mediate the generation of paucimannosidic proteins via a putative noncanonical truncation pathway in human neutrophils. Glycobiology, 2022, 32, 218-229.	1.3	15
2	The Complexity and Dynamics of the Tissue Glycoproteome Associated With Prostate Cancer Progression. Molecular and Cellular Proteomics, 2021, 20, 100026.	2.5	39
3	Hyper-truncated Asn355- and Asn391-glycans modulate the activity of neutrophil granule myeloperoxidase. Journal of Biological Chemistry, 2021, 296, 100144.	1.6	31
4	Serum N-Glycomics Stratifies Bacteremic Patients Infected with Different Pathogens. Journal of Clinical Medicine, 2021, 10, 516.	1.0	12
5	Comparative analysis of the protein profile from biofortified cultivars of quality protein maize and conventional maize by gel-based and gel-free proteomic approaches. LWT - Food Science and Technology, 2021, 138, 110683.	2.5	3
6	Mapping the SARS-CoV-2 spike glycoprotein-derived peptidome presented by HLA class II on dendritic cells. Cell Reports, 2021, 35, 109179.	2.9	63
7	Towards structure-focused glycoproteomics. Biochemical Society Transactions, 2021, 49, 161-186.	1.6	60
8	Trends in oligomannosylation and $\hat{l}\pm 1,2$ -mannosidase expression in human cancers. Oncotarget, 2021, 12, 2188-2205.	0.8	17
9	Community evaluation of glycoproteomics informatics solutions reveals high-performance search strategies for serum glycopeptide analysis. Nature Methods, 2021, 18, 1304-1316.	9.0	74
10	ADAM17 cytoplasmic domain modulates Thioredoxin-1 conformation and activity. Redox Biology, 2020, 37, 101735.	3.9	6
11	The intracellular bacterium Rickettsia rickettsii exerts an inhibitory effect on the apoptosis of tick cells. Parasites and Vectors, 2020, 13, 603.	1.0	11
12	High-resolution longitudinal N- and O-glycoprofiling of human monocyte-to-macrophage transition. Glycobiology, 2020, 30, 679-694.	1.3	26
13	Protein Paucimannosylation Is an Enriched <i>N</i> â€Glycosylation Signature of Human Cancers. Proteomics, 2019, 19, e1900010.	1.3	52
14	Tissue Proteome Signatures Associated with Five Grades of Prostate Cancer and Benign Prostatic Hyperplasia. Proteomics, 2019, 19, e1900174.	1.3	27
15	Zika Virus Impairs Neurogenesis and Synaptogenesis Pathways in Human Neural Stem Cells and Neurons. Frontiers in Cellular Neuroscience, 2019, 13, 64.	1.8	65
16	Integrated Proteomics Reveals Apoptosis-related Mechanisms Associated with Placental Malaria*. Molecular and Cellular Proteomics, 2019, 18, 182-199.	2.5	15
17	Thioredoxin-1 Negatively Modulates ADAM17 Activity Through Direct Binding and Indirect Reductive Activity. Antioxidants and Redox Signaling, 2018, 29, 717-734.	2.5	9
18	Development of a Trypanosoma cruzi strain typing assay using MS2 peptide spectral libraries (Tc-STAMS2). PLoS Neglected Tropical Diseases, 2018, 12, e0006351.	1.3	12

#	Article	IF	Citations
19	Distinct urinary glycoprotein signatures in prostate cancer patients. Oncotarget, 2018, 9, 33077-33097.	0.8	33
20	Protein glycosylation in Trypanosoma cruzi and mass spectrometry-based strategies for glycan and glycoprotein characterization. , 2018, , .		0
21	Comprehensive glycoprofiling of the epimastigote and trypomastigote stages of Trypanosoma cruzi. Journal of Proteomics, 2017, 151, 182-192.	1.2	52
22	Novel DNA coding regions and protein arginylation reveal unexplored T. cruzi proteome and PTMs. International Journal of Mass Spectrometry, 2017, 418, 51-66.	0.7	4
23	MP87-03 URINARY MMP-9 AS CANDIDATE FOR A NON-INVASIVE PROSTATE CANCER BIOMARKER REVEALED BY QUANTITATIVE PROTEOMICS ANALYSIS. Journal of Urology, 2017, 197, .	0.2	0
24	Mass spectrometry-based proteomics revealed Glypican-1 as a novel ADAM17 substrate. Journal of Proteomics, 2017, 151, 53-65.	1.2	23
25	Analysis of the Salivary Gland Transcriptome of Unfed and Partially Fed Amblyomma sculptum Ticks and Descriptive Proteome of the Saliva. Frontiers in Cellular and Infection Microbiology, 2017, 7, 476.	1.8	79
26	Quantitative proteomic analysis of amastigotes from Leishmania (L.) amazonensis LV79 and PH8 strains reveals molecular traits associated with the virulence phenotype. PLoS Neglected Tropical Diseases, 2017, 11, e0006090.	1.3	22
27	A targeted proteomic strategy for the measurement of oral cancer candidate biomarkers in human saliva. Proteomics, 2016, 16, 159-173.	1.3	66
28	EEF1D modulates proliferation and epithelial–mesenchymal transition in oral squamous cell carcinoma. Clinical Science, 2016, 130, 785-799.	1.8	33
29	Site-specific characterization of N-linked glycosylation in human urinary glycoproteins and endogenous glycopeptides. Glycoconjugate Journal, 2016, 33, 937-951.	1.4	15
30	Integrative analysis to select cancer candidate biomarkers to targeted validation. Oncotarget, 2015, 6, 43635-43652.	0.8	18
31	Agrin and Perlecan Mediate Tumorigenic Processes in Oral Squamous Cell Carcinoma. PLoS ONE, 2014, 9, e115004.	1.1	44
32	ADAM17 mediates OSCC development in an orthotopic murine model. Molecular Cancer, 2014, 13, 24.	7.9	16
33	Deciphering the Role of the ADAM17-Dependent Secretome in Cell Signaling. Journal of Proteome Research, 2014, 13, 2080-2093.	1.8	38
34	Integrated Proteomics Identified Up-Regulated Focal Adhesion-Mediated Proteins in Human Squamous Cell Carcinoma in an Orthotopic Murine Model. PLoS ONE, 2014, 9, e98208.	1.1	10
35	Melanogenesis stimulation in B16-F10 melanoma cells induces cell cycle alterations, increased ROS levels and a differential expression of proteins as revealed by proteomic analysis. Experimental Cell Research, 2012, 318, 1913-1925.	1.2	41
36	Novel Processed Form of Syndecan-1 Shed from SCC-9 Cells Plays a Role in Cell Migration. PLoS ONE, 2012, 7, e43521.	1,1	37