## Blendi Ura

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8459079/publications.pdf Version: 2024-02-01



RIENDILLOA

#	Article	IF	CITATIONS
1	Gel-Based Proteomic Identification of Suprabasin as a Potential New Candidate Biomarker in Endometrial Cancer. International Journal of Molecular Sciences, 2022, 23, 2076.	4.1	7
2	Two Dimensional-Difference in Gel Electrophoresis (2D-DIGE) Proteomic Approach for the Identification of Biomarkers in Endometrial Cancer Serum. Cancers, 2021, 13, 3639.	3.7	13
3	Proteomic Study Identifies Glycolytic and Inflammation Pathways Involved in Recurrent Otitis Media. International Journal of Molecular Sciences, 2020, 21, 9291.	4.1	3
4	Phosphoproteins Involved in the Inhibition of Apoptosis and in Cell Survival in the Leiomyoma. Journal of Clinical Medicine, 2019, 8, 691.	2.4	14
5	Leiomyoma phosphoproteins involved in inhibition of oxidative stress and synthesis of reactive oxygen species. International Journal of Molecular Medicine, 2019, 44, 2329-2335.	4.0	1
6	In vitro treatment of congenital disorder of glycosylation type Ia using PLGA nanoparticles loaded with GDP‑Man. International Journal of Molecular Medicine, 2019, 44, 262-272.	4.0	4
7	Interstitial Fluid in Gynecologic Tumors and Its Possible Application in the Clinical Practice. International Journal of Molecular Sciences, 2018, 19, 4018.	4.1	8
8	Dysregulated chaperones associated with cell proliferation and negative apoptosis regulation in the uterine leiomyoma. Oncology Letters, 2018, 15, 8005-8010.	1.8	2
9	Identification of proteins with different abundance associated with cell migration and proliferation in leiomyoma interstitial fluid by proteomics. Oncology Letters, 2017, 13, 3912-3920.	1.8	9
10	A proteomic approach for the identification of biomarkers in endometrial cancer uterine aspirate. Oncotarget, 2017, 8, 109536-109545.	1.8	19
11	Serological Proteome Analysis for Identification of Potential Antigen in Atopic Dermatitis. Pediatrics & Health Research, 2016, 01, .	0.0	Ο
12	A Proteomic Approach for the Identification of Up-Regulated Proteins Involved in the Metabolic Process of the Leiomyoma. International Journal of Molecular Sciences, 2016, 17, 540.	4.1	13
13	Abnormal expression of leiomyoma cytoskeletal proteins involved in cell migration. Oncology Reports, 2016, 35, 3094-3100.	2.6	11
14	Changes in Protein Expression in Two Cholangiocarcinoma Cell Lines Undergoing Formation of Multicellular Tumor Spheroids In Vitro. PLoS ONE, 2015, 10, e0118906.	2.5	16
15	Two-dimensional gel electrophoresis analysis of the leiomyoma interstitial fluid reveals altered protein expression with a possible involvement in pathogenesis. Oncology Reports, 2015, 33, 2219-2226.	2.6	13
16	Comparative analysis of the seminal plasma proteomes of oligoasthenozoospermic and normozoospermic men. Reproductive BioMedicine Online, 2015, 30, 522-531.	2.4	33
17	Association between up-regulated expression proteins and circulating steroidal hormones in leiomyoma. Medical Hypotheses, 2015, 85, 515.	1.5	3
18	Potential role of circulating microRNAs as early markers of preeclampsia. Taiwanese Journal of Obstetrics and Gynecology, 2014, 53, 232-234.	1.3	86

#	Article	IF	CITATIONS
19	The side chain of glutamine 13 is the acyl-donor amino acid modified by type 2 transglutaminase in subunit T of the native rabbit skeletal muscle troponin complex. Amino Acids, 2013, 44, 227-234.	2.7	1
20	Comparative proteomic analysis of ductal breast carcinoma demonstrates an altered expression of chaperonins and cytoskeletal proteins. Molecular Medicine Reports, 2013, 7, 1700-1704.	2.4	9