

Indira Pla

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

331
citations

1040056

9
h-index

888059

17
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28
all docs

28
docs citations

28
times ranked

613
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteomic Alterations in Follicular Fluid of Human Small Antral Follicles Collected from Polycystic Ovariesâ€”A Pilot Study. <i>Life</i> , 2022, 12, 391.	2.4	0
2	Novel protein markers of androgen activity in humans: proteomic study of plasma from young chemically castrated men. <i>ELife</i> , 2022, 11, .	6.0	3
3	The human melanoma proteome atlasâ€”Defining the molecular pathology. <i>Clinical and Translational Medicine</i> , 2021, 11, e473.	4.0	14
4	The Human Melanoma Proteome Atlasâ€”Complementing the melanoma transcriptome. <i>Clinical and Translational Medicine</i> , 2021, 11, e451.	4.0	20
5	A biobanking turningâ€”point in the use of formalinâ€”fixed, paraffin tumor blocks to unveil kinase signaling in melanoma. <i>Clinical and Translational Medicine</i> , 2021, 11, e466.	4.0	6
6	Proteome of fluid from human ovarian small antral follicles reveals insights in folliculogenesis and oocyte maturation. <i>Human Reproduction</i> , 2021, 36, 756-770.	0.9	17
7	Short-Term Effect of Induced Alterations in Testosterone Levels on Fasting Plasma Amino Acid Levels in Healthy Young Men. <i>Life</i> , 2021, 11, 1276.	2.4	2
8	Mapping the Melanoma Plasma Proteome (MPP) Using Single-Shot Proteomics Interfaced with the WiMT Database. <i>Cancers</i> , 2021, 13, 6224.	3.7	4
9	Novel functional proteins coded by the human genome discovered in metastases of melanoma patients. <i>Cell Biology and Toxicology</i> , 2020, 36, 261-272.	5.3	9
10	Short-term effect of pharmacologically induced alterations in testosterone levels on common blood biomarkers in a controlled healthy human model. <i>Scandinavian Journal of Clinical and Laboratory Investigation</i> , 2020, 80, 25-31.	1.2	7
11	Mass spectrometry-based analysis of formalin-fixed, paraffin-embedded distal cholangiocarcinoma identifies stromal thrombospondin-2 as a potential prognostic marker. <i>Journal of Translational Medicine</i> , 2020, 18, 343.	4.4	10
12	Proteomic analysis enables distinction of earlyâ€”versus advancedâ€”stage lung adenocarcinomas. <i>Clinical and Translational Medicine</i> , 2020, 10, e106.	4.0	7
13	Protein Expression in Metastatic Melanoma and the Link to Disease Presentation in a Range of Tumor Phenotypes. <i>Cancers</i> , 2020, 12, 767.	3.7	2
14	A pilot proteomic study reveals different protein profiles related to testosterone and gonadotropin changes in a short-term controlled healthy human cohort. <i>Journal of Proteomics</i> , 2020, 220, 103768.	2.4	4
15	Alpha-1-acid glycoprotein 1 is upregulated in pancreatic ductal adenocarcinoma and confers a poor prognosis. <i>Translational Research</i> , 2019, 212, 67-79.	5.0	11
16	Progressive changes in human follicular fluid composition over the course of ovulation: quantitative proteomic analyses. <i>Molecular and Cellular Endocrinology</i> , 2019, 495, 110522.	3.2	29
17	Proteomic signatures of brain regions affected by tau pathology in early and late stages of Alzheimer's disease. <i>Neurobiology of Disease</i> , 2019, 130, 104509.	4.4	46
18	Quantitative proteomics identifies brain acid soluble protein 1 (BASP1) as a prognostic biomarker candidate in pancreatic cancer tissue. <i>EBioMedicine</i> , 2019, 43, 282-294.	6.1	43

#	ARTICLE	IF	CITATIONS
19	Clinical protein science in translational medicine targeting malignant melanoma. <i>Cell Biology and Toxicology</i> , 2019, 35, 293-332.	5.3	33
20	Improved survival prognostication of node-positive malignant melanoma patients utilizing shotgun proteomics guided by histopathological characterization and genomic data. <i>Scientific Reports</i> , 2019, 9, 5154.	3.3	12
21	The Hidden Story of Heterogeneous B-raf V600E Mutation Quantitative Protein Expression in Metastatic Melanoma—Association with Clinical Outcome and Tumor Phenotypes. <i>Cancers</i> , 2019, 11, 1981.	3.7	16
22	Quantitative Assessment of Urea In-Solution Lys-C/Trypsin Digestions Reveals Superior Performance at Room Temperature over Traditional Proteolysis at 37 Å°C. <i>Journal of Proteome Research</i> , 2018, 17, 2556-2561.	3.7	32
23	Mass spectrometry evaluation of a neuroblastoma SH-SY5Y cell culture protocol. <i>Analytical Biochemistry</i> , 2018, 559, 51-54.	2.4	2