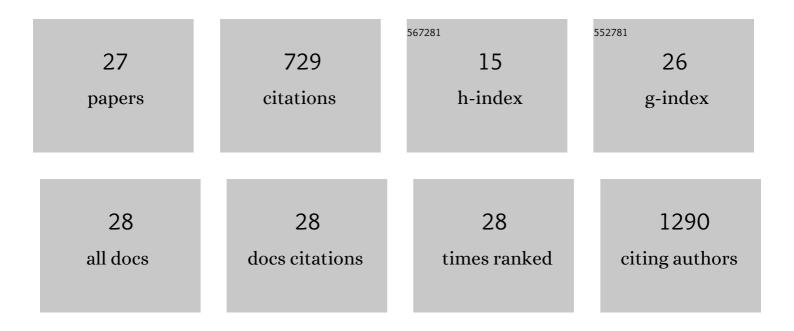
## Renu Pandey

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8087125/publications.pdf Version: 2024-02-01



Ρενιί Ρλνσεν

#	Article	IF	CITATIONS
1	Inhibition of mitochondrial complex I reverses NOTCH1-driven metabolic reprogramming in T-cell acute lymphoblastic leukemia. Nature Communications, 2022, 13, 2801.	12.8	25
2	Novel Strategy for Untargeted Chiral Metabolomics using Liquid Chromatography-High Resolution Tandem Mass Spectrometry. Analytical Chemistry, 2021, 93, 5805-5814.	6.5	17
3	Stable Isotope Dilution LC-HRMS Assay To Determine Free SN-38, Total SN-38, and SN-38G in a Tumor Xenograft Model after Intravenous Administration of Antibody–Drug Conjugate (Sacituzumab) Tj ETQq1 1 0	.784&\$4 rg	BT <b>\$</b> Overloc®
4	The RNA-binding protein SERBP1 functions as a novel oncogenic factor in glioblastoma by bridging cancer metabolism and epigenetic regulation. Genome Biology, 2020, 21, 195.	8.8	55
5	Identification of a synergistic combination of dimethylaminoparthenolide and shikonin alters metabolism and inhibits proliferation of pediatric precursorâ€B cell acute lymphoblastic leukemia. Molecular Carcinogenesis, 2020, 59, 399-411.	2.7	19
6	Enzyme-mediated depletion of serum <scp>l</scp> -Met abrogates prostate cancer growth via multiple mechanisms without evidence of systemic toxicity. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 13000-13011.	7.1	27
7	Glutaminase Inhibition Overcomes Acquired Resistance to Mitochondrial Complex I in NOTCH1-Driven T-Cell Acute Lymphoblastic Leukemias (T-ALL) Via Block of Glutamine Driven Reductive Metabolism. Blood, 2019, 134, 806-806.	1.4	1
8	Highly sensitive and selective determination of redox states of coenzymes Q9 and Q10 in mice tissues: Application of orbitrap mass spectrometry. Analytica Chimica Acta, 2018, 1011, 68-76.	5.4	23
9	Mitochondrial Complex I Inhibitor Iacs-010759 Reverses the NOTCH1-Driven Metabolic Reprogramming in T-ALL Via Blockade of Oxidative Phosphorylation: Synergy with Chemotherapy and Clutaminase Inhibition. Blood, 2018, 132, 4020-4020.	1.4	7
10	Metabolomic signature of brain cancer. Molecular Carcinogenesis, 2017, 56, 2355-2371.	2.7	86
11	Bioguided chemical characterization of the antiproliferative fraction of edible pseudo bulbs of Malaxis acuminata D. Don by HPLC-ESI-QTOF-MS. Medicinal Chemistry Research, 2017, 26, 3307-3314.	2.4	8
12	Major bioactive phenolics in Bergenia species from the Indian Himalayan region: Method development, validation and quantitative estimation using UHPLC-QqQLIT-MS/MS. PLoS ONE, 2017, 12, e0180950.	2.5	16
13	Rapid screening and quantitative determination of bioactive compounds from fruit extracts of Myristica species and their in vitro antiproliferative activity. Food Chemistry, 2016, 211, 483-493.	8.2	26
14	Rapid quantitative analysis of multi-components in Andrographis paniculata using UPLC-QqQLIT-MS/MS: Application to soil sodicity and organic farming. Industrial Crops and Products, 2016, 83, 423-430.	5.2	9
15	A rapid and highly sensitive method for simultaneous determination of bioactive constituents in leaf extracts of six Ocimum species using ultra high performance liquid chromatography-hybrid linear ion trap triple quadrupole mass spectrometry. Analytical Methods, 2016, 8, 333-341.	2.7	9
16	HPLC–QTOF–MS/MS-based rapid screening of phenolics and triterpenic acids in leaf extracts of <i>Ocimum</i> species and their interspecies variation. Journal of Liquid Chromatography and Related Technologies, 2016, 39, 225-238.	1.0	49
17	Quality control assessment of polyherbal formulation based on a quantitative determination multimarker approach by ultra high performance liquid chromatography with tandem mass spectrometry using polarity switching combined with multivariate analysis. Journal of Separation Science, 2015, 38, 3183-3191.	2.5	8
18	Science, 2013, 58, 5185-5191. Simultaneous quantitative determination of multiple bioactive markers in <i>Ocimum sanctum</i> obtained from different locations and its marketed herbal formulations using UPLCâ€ESIâ€MS/MS combined with principal component analysis. Phytochemical Analysis, 2015, 26, 383-394.	2.4	27

Renu Pandey

#	Article	IF	CITATIONS
19	Ultra high performance liquid chromatography tandem mass spectrometry method for the simultaneous determination of multiple bioactive constituents in fruit extracts of <i>Myristica fragrans</i> and its marketed polyherbal formulations using a polarity switching technique. Journal of Separation Science, 2015, 38, 1277-1285.	2.5	16
20	Simultaneous determination of multi-class bioactive constituents for quality assessment of Garcinia species using UHPLC–QqQ LIT –MS/MS. Industrial Crops and Products, 2015, 77, 861-872.	5.2	21
21	Quantification of multianalyte by UPLC–QqQLIT–MS/MS and in-vitro anti-proliferative screening in Cassia species. Industrial Crops and Products, 2015, 76, 1133-1141.	5.2	16
22	Quantitative determination of chemical constituents of Piper spp. using UPLC–ESI–MS/MS. Industrial Crops and Products, 2015, 76, 967-976.	5.2	13
23	Development and validation of an ultra high performance liquid chromatography electrospray ionization tandem mass spectrometry method for the simultaneous determination of selected flavonoids in <i>Ginkgo biloba</i> . Journal of Separation Science, 2014, 37, 3610-3618.	2.5	30
24	A rapid analytical method for characterization and simultaneous quantitative determination of phytoconstituents in Piper betle landraces using UPLC-ESI-MS/MS. Analytical Methods, 2014, 6, 7349.	2.7	11
25	A strategy to access fused triazoloquinoline and related nucleoside analogues. Tetrahedron, 2013, 69, 8547-8558.	1.9	20
26	Characteristic differences in metabolite profile in male and female plants of dioecious Piper betle L Journal of Biosciences, 2012, 37, 1061-1066.	1.1	18
27	Optimized metabolite extraction from blood serum for 1H nuclear magnetic resonance spectroscopy. Analytical Biochemistry, 2008, 377, 16-23.	2.4	164