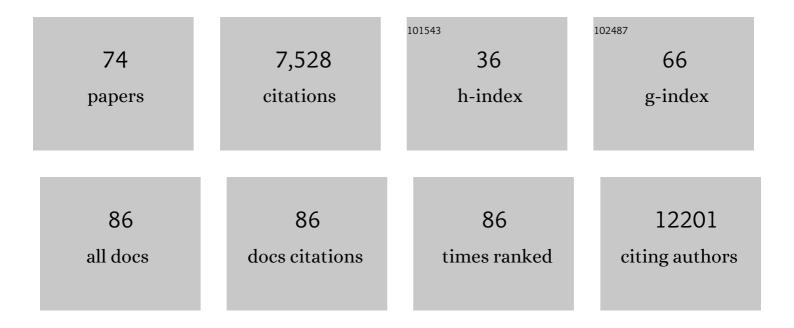
List of Publications by Year in descending order

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ΠΑΥΙΟ Ι ΡΑCΗΑΡΙΝΙ

#	Article	IF	CITATIONS
1	A Mitochondrial Protein Compendium Elucidates Complex I Disease Biology. Cell, 2008, 134, 112-123.	28.9	1,766
2	Upstream open reading frames cause widespread reduction of protein expression and are polymorphic among humans. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 7507-7512.	7.1	741
3	Calorie Restriction and SIRT3 Trigger Global Reprogramming of the Mitochondrial Protein Acetylome. Molecular Cell, 2013, 49, 186-199.	9.7	584
4	A Single Kinase Generates the Majority of the Secreted Phosphoproteome. Cell, 2015, 161, 1619-1632.	28.9	264
5	Mitochondrial Protein Interaction Mapping Identifies Regulators of Respiratory Chain Function. Molecular Cell, 2016, 63, 621-632.	9.7	241
6	Biochemistry of Mitochondrial Coenzyme Q Biosynthesis. Trends in Biochemical Sciences, 2017, 42, 824-843.	7.5	239
7	Mitochondrial modulation: reversible phosphorylation takes center stage?. Trends in Biochemical Sciences, 2006, 31, 26-34.	7.5	211
8	Mutation of C20orf7 Disrupts Complex I Assembly and Causes Lethal Neonatal Mitochondrial Disease. American Journal of Human Genetics, 2008, 83, 468-478.	6.2	175
9	Neutron-encoded mass signatures for multiplexed proteome quantification. Nature Methods, 2013, 10, 332-334.	19.0	165
10	Maximal Oxidative Capacity during Exercise Is Associated with Skeletal Muscle Fuel Selection and Dynamic Changes in Mitochondrial Protein Acetylation. Cell Metabolism, 2015, 21, 468-478.	16.2	165
11	Hallmarks of a new era in mitochondrial biochemistry. Genes and Development, 2013, 27, 2615-2627.	5.9	146
12	A Quantitative Map of the Liver Mitochondrial Phosphoproteome Reveals Posttranslational Control of Ketogenesis. Cell Metabolism, 2012, 16, 672-683.	16.2	141
13	Involvement of a Mitochondrial Phosphatase in the Regulation of ATP Production and Insulin Secretion in Pancreatic β Cells. Molecular Cell, 2005, 19, 197-207.	9.7	138
14	Mitochondrial protein hyperacetylation in the failing heart. JCl Insight, 2016, 1, .	5.0	133
15	Mitochondrial protein functions elucidated by multi-omic mass spectrometry profiling. Nature Biotechnology, 2016, 34, 1191-1197.	17.5	122
16	Mitochondrial COQ9 is a lipid-binding protein that associates with COQ7 to enable coenzyme Q biosynthesis. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E4697-705.	7.1	113
17	Quantification of Mitochondrial Acetylation Dynamics Highlights Prominent Sites of Metabolic Regulation. Journal of Biological Chemistry, 2013, 288, 26209-26219.	3.4	105
18	Mitochondrial ADCK3 Employs an Atypical Protein Kinase-like Fold to Enable Coenzyme Q Biosynthesis. Molecular Cell, 2015, 57, 83-94.	9.7	104

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19	Cerebellar Ataxia and Coenzyme Q Deficiency through Loss of Unorthodox Kinase Activity. Molecular Cell, 2016, 63, 608-620.	9.7	101
20	Classification of T-cell activation via autofluorescence lifetime imaging. Nature Biomedical Engineering, 2021, 5, 77-88.	22.5	92
21	Two-stage metabolic remodelling in macrophages in response to lipopolysaccharide and interferon-Î ³ stimulation. Nature Metabolism, 2019, 1, 731-742.	11.9	90
22	Multi-omics Reveal Specific Targets of the RNA-Binding Protein Puf3p and Its Orchestration of Mitochondrial Biogenesis. Cell Systems, 2018, 6, 125-135.e6.	6.2	80
23	A PTEN-like Phosphatase with a Novel Substrate Specificity. Journal of Biological Chemistry, 2004, 279, 38590-38596.	3.4	70
24	Coenzyme Q biosynthetic proteins assemble in a substrate-dependent manner into domains at ER–mitochondria contacts. Journal of Cell Biology, 2019, 218, 1353-1369.	5.2	69
25	Complementary RNA and Protein Profiling Identifies Iron as a Key Regulator of Mitochondrial Biogenesis. Cell Reports, 2013, 3, 237-245.	6.4	67
26	Conserved Lipid and Small-Molecule Modulation of COQ8 Reveals Regulation of the Ancient Kinase-like UbiB Family. Cell Chemical Biology, 2018, 25, 154-165.e11.	5.2	63
27	Analysis of the Acidic Proteome with Negative Electron-Transfer Dissociation Mass Spectrometry. Analytical Chemistry, 2012, 84, 2875-2882.	6.5	57
28	Amine-reactive Neutron-encoded Labels for Highly Plexed Proteomic Quantitation. Molecular and Cellular Proteomics, 2013, 12, 3360-3369.	3.8	57
29	Systems Analyses Reveal Physiological Roles and Genetic Regulators of Liver Lipid Species. Cell Systems, 2018, 6, 722-733.e6.	6.2	54
30	Pharmacological Targeting of the Mitochondrial Phosphatase PTPMT1. Journal of Pharmacology and Experimental Therapeutics, 2010, 333, 584-592.	2.5	53
31	Genetic Regulation of Plasma Lipid Species and Their Association with Metabolic Phenotypes. Cell Systems, 2018, 6, 709-721.e6.	6.2	52
32	Clinico enetic, Imaging and Molecular Delineation of <scp> <i>COQ8A</i> </scp> â€Ataxia: A Multicenter Study of 59 Patients. Annals of Neurology, 2020, 88, 251-263.	5.3	52
33	Mitochondrial DNA Variant in COX1 Subunit Significantly Alters Energy Metabolism of Geographically Divergent Wild Isolates in Caenorhabditis elegans. Journal of Molecular Biology, 2014, 426, 2199-2216.	4.2	49
34	Defining mitochondrial protein functions through deep multiomic profiling. Nature, 2022, 606, 382-388.	27.8	49
35	Quantitative shotgun proteome analysis by direct infusion. Nature Methods, 2020, 17, 1222-1228.	19.0	48
36	Inhibitors of bacterial tubulin target bacterial membranes <i>in vivo</i> . MedChemComm, 2013, 4, 112-119.	3.4	45

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37	Neutron-Encoded Mass Signatures for Quantitative Top-Down Proteomics. Analytical Chemistry, 2014, 86, 2314-2319.	6.5	45
38	Ptc7p Dephosphorylates Select Mitochondrial Proteins to Enhance Metabolic Function. Cell Reports, 2017, 18, 307-313.	6.4	45
39	Multi-omic Mitoprotease Profiling Defines a Role for Oct1p in Coenzyme Q Production. Molecular Cell, 2017, 68, 970-977.e11.	9.7	45
40	Pptc7 is an essential phosphatase for promoting mammalian mitochondrial metabolism and biogenesis. Nature Communications, 2019, 10, 3197.	12.8	45
41	Multiplexed Quantification for Data-Independent Acquisition. Analytical Chemistry, 2015, 87, 2570-2575.	6.5	43
42	Intelligent Data Acquisition Blends Targeted and Discovery Methods. Journal of Proteome Research, 2014, 13, 2152-2161.	3.7	39
43	Pathogenic Bi-allelic Mutations in NDUFAF8 Cause Leigh Syndrome with an Isolated Complex I Deficiency. American Journal of Human Genetics, 2020, 106, 92-101.	6.2	39
44	An Isoprene Lipid-Binding Protein Promotes Eukaryotic Coenzyme Q Biosynthesis. Molecular Cell, 2019, 73, 763-774.e10.	9.7	37
45	Mass spectrometry proteomics reveals a function for mammalian CALCOCO1 in MTOR-regulated selective autophagy. Autophagy, 2020, 16, 2219-2237.	9.1	37
46	Erythropoietin signaling regulates heme biosynthesis. ELife, 2017, 6, .	6.0	36
47	Obesity-dependent CDK1 signaling stimulates mitochondrial respiration at complex I in pancreatic β-cells. Journal of Biological Chemistry, 2019, 294, 4656-4666.	3.4	35
48	A PTEN-related 5-Phosphatidylinositol Phosphatase Localized in the Golgi. Journal of Biological Chemistry, 2003, 278, 39866-39873.	3.4	32
49	Solution NMR Structure of Yeast Succinate Dehydrogenase Flavinylation Factor Sdh5 Reveals a Putative Sdh1 Binding Site. Biochemistry, 2012, 51, 8475-8477.	2.5	29
50	Iron Deprivation Induces Transcriptional Regulation of Mitochondrial Biogenesis. Journal of Biological Chemistry, 2016, 291, 20827-20837.	3.4	28
51	DNA Polymerase Î, Increases Mutational Rates in Mitochondrial DNA. ACS Chemical Biology, 2018, 13, 900-908.	3.4	26
52	UbiB proteins regulate cellular CoQ distribution in Saccharomyces cerevisiae. Nature Communications, 2021, 12, 4769.	12.8	26
53	Integrative proteomics and biochemical analyses define Ptc6p as the Saccharomyces cerevisiae pyruvate dehydrogenase phosphatase. Journal of Biological Chemistry, 2017, 292, 11751-11759.	3.4	25
54	Dual Specificity Phosphatases 18 and 21 Target to Opposing Sides of the Mitochondrial Inner Membrane. Journal of Biological Chemistry, 2008, 283, 15440-15450.	3.4	24

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55	Identification and Quantification of Murine Mitochondrial Proteoforms Using an Integrated Top-Down and Intact-Mass Strategy. Journal of Proteome Research, 2018, 17, 3526-3536.	3.7	23
56	The extensive and functionally uncharacterized mitochondrial phosphoproteome. Journal of Biological Chemistry, 2021, 297, 100880.	3.4	23
57	Multi-Omic Single-Shot Technology for Integrated Proteome and Lipidome Analysis. Analytical Chemistry, 2021, 93, 4217-4222.	6.5	22
58	Argonaut: A Web Platform for Collaborative Multi-omic Data Visualization and Exploration. Patterns, 2020, 1, 100122.	5.9	18
59	A Gly-Zipper Motif Mediates Homodimerization of the Transmembrane Domain of the Mitochondrial Kinase ADCK3. Journal of the American Chemical Society, 2014, 136, 14068-14077.	13.7	17
60	Systems Biochemistry Approaches to Defining Mitochondrial Protein Function. Cell Metabolism, 2020, 31, 669-678.	16.2	16
61	ADCK3/COQ8A: the choice target of the UbiB protein kinase-like family. Nature Reviews Drug Discovery, 2019, 18, 815-815.	46.4	15
62	Automated Gas-Phase Purification for Accurate, Multiplexed Quantification on a Stand-Alone Ion-Trap Mass Spectrometer. Analytical Chemistry, 2013, 85, 2079-2086.	6.5	13
63	Defining intermediates and redundancies in coenzyme Q precursor biosynthesis. Journal of Biological Chemistry, 2021, 296, 100643.	3.4	12
64	Loss of C2orf69 defines a fatal autoinflammatory syndrome in humans and zebrafish that evokes a glycogen-storage-associated mitochondriopathy. American Journal of Human Genetics, 2021, 108, 1301-1317.	6.2	11
65	Prolyl endopeptidase-like is a (thio)esterase involved in mitochondrial respiratory chain function. IScience, 2021, 24, 103460.	4.1	8
66	Protein Tyrosine Phosphatases. , 2004, , 536-542.		3
67	Transomics: Mitochondrial Systems Analyses Get Supercomplex. Cell Metabolism, 2016, 24, 13-14.	16.2	1
68	Clinico-Genetic, Imaging and Molecular Delineation of COQ8A-Ataxia: A Multicenter Study of 59 Patients. , 2020, 88, 251.		1
69	A path to the powerhouse: systemsâ€ŧoâ€structure approaches for studying mitochondrial proteins. Protein Science, 2018, 27, 1518-1525.	7.6	0
70	Exploring the Role of an Atypical Kinase in Ubiquinone Biosynthesis. FASEB Journal, 2012, 26, 565.12.	0.5	0
71	Ironâ€dependent regulation of mitochondrial form and function. FASEB Journal, 2013, 27, lb65.	0.5	0
72	Mitochondrial protein functions revealed by global proteinâ€lipidâ€metabolite profiles. FASEB Journal, 2016, 30, 1100.4.	0.5	0

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73	COQ9 Membrane Association and Its Role in Coenzyme Q Biosynthesis. FASEB Journal, 2018, 32, 815.8.	0.5	0
74	Design and Evaluation of Cleavable CoQâ€ŧriphenylphosphonium Analogs. FASEB Journal, 2022, 36, .	0.5	0