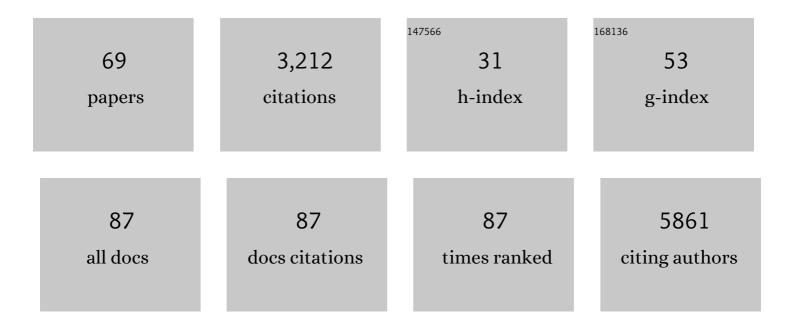
David Meierhofer

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2476499/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Variants in mitochondrial amidoxime reducing component 1 and hydroxysteroid 17â€beta dehydrogenase 13 reduce severity of nonalcoholic fatty liver disease in children and suppress fibrotic pathways through distinct mechanisms. Hepatology Communications, 2022, 6, 1934-1948.	2.0	18
2	Sleep neuron depolarization promotes protective gene expression changes and FOXO activation. Current Biology, 2022, 32, 2248-2262.e9.	1.8	5
3	Alternative splicing of BUD13 determines the severity of a developmental disorder with lipodystrophy and progeroid features. Genetics in Medicine, 2022, 24, 1927-1940.	1.1	2
4	Functional Consequences of Metabolic Zonation in Murine Livers: Insights for an Old Story. Hepatology, 2021, 73, 795-810.	3.6	35
5	Metabolic heterogeneity of human hepatocellular carcinoma: implications for personalized pharmacological treatment. FEBS Journal, 2021, 288, 2332-2346.	2.2	12
6	Regulation of the cytochrome P450 epoxyeicosanoid pathway is associated with distinct histologic features in pediatric non-alcoholic fatty liver disease. Prostaglandins Leukotrienes and Essential Fatty Acids, 2021, 164, 102229.	1.0	6
7	Viability Assessment in Liver Transplantation—What Is the Impact of Dynamic Organ Preservation?. Biomedicines, 2021, 9, 161.	1.4	47
8	Defective metabolic programming impairs early neuronal morphogenesis in neural cultures and an organoid model of Leigh syndrome. Nature Communications, 2021, 12, 1929.	5.8	55
9	Dietary-challenged mice with Alzheimer-like pathology show increased energy expenditure and reduced adipocyte hypertrophy and steatosis. Aging, 2021, 13, 10891-10919.	1.4	2
10	Dnmt1 has de novo activity targeted to transposable elements. Nature Structural and Molecular Biology, 2021, 28, 594-603.	3.6	83
11	Deletion of mTOR in liver epithelial cells enhances hepatic metastasis of colon cancer. Journal of Pathology, 2021, 255, 270-284.	2.1	6
12	A BRD4-mediated elongation control point primes transcribing RNA polymerase II for 3′-processing and termination. Molecular Cell, 2021, 81, 3589-3603.e13.	4.5	31
13	Cell autonomous requirement of neurofibromin (Nf1) for postnatal muscle hypertrophic growth and metabolic homeostasis. Journal of Cachexia, Sarcopenia and Muscle, 2020, 11, 1758-1778.	2.9	8
14	Systematic Surveys of Iron Homeostasis Mechanisms Reveal Ferritin Superfamily and Nucleotide Surveillance Regulation to be Modified by PINK1 Absence. Cells, 2020, 9, 2229.	1.8	9
15	Decreased Mitochondrial DNA Content Drives OXPHOS Dysregulation in Chromophobe Renal Cell Carcinoma. Cancer Research, 2020, 80, 3830-3840.	0.4	9
16	Endocytosis-Mediated Replenishment of Amino Acids Favors Cancer Cell Proliferation and Survival in Chromophobe Renal Cell Carcinoma. Cancer Research, 2020, 80, 5491-5501.	0.4	11
17	Rapid and Culture Free Identification of Francisella in Hare Carcasses by High-Resolution Tandem Mass Spectrometry Proteotyping. Frontiers in Microbiology, 2020, 11, 636.	1.5	8
18	Hypothermic oxygenated perfusion protects from mitochondrial injury before liver transplantation. EBioMedicine, 2020, 60, 103014.	2.7	111

DAVID MEIERHOFER

#	Article	IF	CITATIONS
19	SON and SRRM2 are essential for nuclear speckle formation. ELife, 2020, 9, .	2.8	122
20	Are Hydroethidine-Based Probes Reliable for Reactive Oxygen Species Detection?. Antioxidants and Redox Signaling, 2019, 31, 359-367.	2.5	27
21	PS-168-Novel real time prediction of liver graft function during hypothermic oxygenated machine perfusion prior to liver transplantation. Journal of Hepatology, 2019, 70, e104-e105.	1.8	2
22	Generation of an Atxn2-CAG100 knock-in mouse reveals N-acetylaspartate production deficit due to early Nat8l dysregulation. Neurobiology of Disease, 2019, 132, 104559.	2.1	24
23	Glutathione Metabolism in Renal Cell Carcinoma Progression and Implications for Therapies. International Journal of Molecular Sciences, 2019, 20, 3672.	1.8	61
24	Brain maturation is associated with increasing tissue stiffness and decreasing tissue fluidity. Acta Biomaterialia, 2019, 99, 433-442.	4.1	55
25	Papillary Renal Cell Carcinomas Rewire Glutathione Metabolism and Are Deficient in Both Anabolic Glucose Synthesis and Oxidative Phosphorylation. Cancers, 2019, 11, 1298.	1.7	15
26	Acylcarnitine profiling by low-resolution LC-MS. PLoS ONE, 2019, 14, e0221342.	1.1	16
27	Mutations in NDUFS1 Cause Metabolic Reprogramming and Disruption of the Electron Transfer. Cells, 2019, 8, 1149.	1.8	30
28	Characterization of Lipid and Lipid Droplet Metabolism in Human HCC. Cells, 2019, 8, 512.	1.8	60
29	Mutual Zonated Interactions of Wnt and Hh Signaling Are Orchestrating the Metabolism of the Adult Liver in Mice and Human. Cell Reports, 2019, 29, 4553-4567.e7.	2.9	15
30	In Human and Mouse Spino-Cerebellar Tissue, Ataxin-2 Expansion Affects Ceramide-Sphingomyelin Metabolism. International Journal of Molecular Sciences, 2019, 20, 5854.	1.8	19
31	Novel Real-time Prediction of Liver Graft Function During Hypothermic Oxygenated Machine Perfusion Before Liver Transplantation. Annals of Surgery, 2019, 270, 783-790.	2.1	146
32	Genetic determinants of steatosis and fibrosis progression in paediatric nonâ€ e lcoholic fatty liver disease. Liver International, 2019, 39, 540-556.	1.9	54
33	HEPATOKIN1 is a biochemistry-based model of liver metabolism for applications in medicine and pharmacology. Nature Communications, 2018, 9, 2386.	5.8	44
34	Human iPSC-Derived Neural Progenitors Are an Effective Drug Discovery Model for Neurological mtDNA Disorders. Cell Stem Cell, 2017, 20, 659-674.e9.	5.2	126
35	Quantitative Global Proteomics of Yeast PBP1 Deletion Mutants and Their Stress Responses Identifies Glucose Metabolism, Mitochondrial, and Stress Granule Changes. Journal of Proteome Research, 2017, 16, 504-515.	1.8	22
36	Defining Human Tyrosine Kinase Phosphorylation Networks Using Yeast as an In Vivo Model Substrate. Cell Systems, 2017, 5, 128-139.e4.	2.9	20

DAVID MEIERHOFER

#	Article	IF	CITATIONS
37	Efficient Prevention of Neurodegenerative Diseases by Depletion of Starvation Response Factor Ataxin-2. Trends in Neurosciences, 2017, 40, 507-516.	4.2	51
38	\ddot{I} O Cells Feature De-Ubiquitination of SLC Transporters and Increased Levels and Fluxes of Amino Acids. International Journal of Molecular Sciences, 2017, 18, 879.	1.8	5
39	Progression of pathology in PINK1-deficient mouse brain from splicing via ubiquitination, ER stress, and mitophagy changes to neuroinflammation. Journal of Neuroinflammation, 2017, 14, 154.	3.1	63
40	Renal oncocytoma characterized by the defective complex I of the respiratory chain boosts the synthesis of the ROS scavenger glutathione. Oncotarget, 2017, 8, 105882-105904.	0.8	32
41	PHF13 is a molecular reader and transcriptional co-regulator of H3K4me2/3. ELife, 2016, 5, .	2.8	22
42	Advantages and Pitfalls of Mass Spectrometry Based Metabolome Profiling in Systems Biology. International Journal of Molecular Sciences, 2016, 17, 632.	1.8	129
43	Serial interactome capture of the human cell nucleus. Nature Communications, 2016, 7, 11212.	5.8	122
44	Ataxin-2 (Atxn2)-Knock-Out Mice Show Branched Chain Amino Acids and Fatty Acids Pathway Alterations. Molecular and Cellular Proteomics, 2016, 15, 1728-1739.	2.5	70
45	An Impaired Respiratory Electron Chain Triggers Down-regulation of the Energy Metabolism and De-ubiquitination of Solute Carrier Amino Acid Transporters. Molecular and Cellular Proteomics, 2016, 15, 1526-1538.	2.5	19
46	Identification and characterization of DNA sequences that prevent glucocorticoid receptor binding to nearby response elements. Nucleic Acids Research, 2016, 44, 6142-6156.	6.5	10
47	Mutations in Subunits of the Activating Signal Cointegrator 1 Complex Are Associated with Prenatal Spinal Muscular Atrophy and Congenital Bone Fractures. American Journal of Human Genetics, 2016, 98, 473-489.	2.6	56
48	Phosphorylation of the chromatin remodeling factor DPF3a induces cardiac hypertrophy through releasing HEY repressors from DNA. Nucleic Acids Research, 2016, 44, 2538-2553.	6.5	30
49	The long non-coding RNA PARROT is an upstream regulator of c-Myc and affects proliferation and translation. Oncotarget, 2016, 7, 33934-33947.	0.8	6
50	GORAB Missense Mutations Disrupt RAB6 and ARF5 Binding and Golgi Targeting. Journal of Investigative Dermatology, 2015, 135, 2368-2376.	0.3	28
51	Bioenergetic cues shift FXR splicing towards FXRα2 to modulate hepatic lipolysis and fatty acid metabolism. Molecular Metabolism, 2015, 4, 891-902.	3.0	33
52	Recurrent De Novo Mutations Affecting Residue Arg138 of Pyrroline-5-Carboxylate Synthase Cause a Progeroid Form of Autosomal-Dominant Cutis Laxa. American Journal of Human Genetics, 2015, 97, 483-492.	2.6	70
53	Metabolome and Proteome Profiling of Complex I Deficiency Induced by Rotenone. Journal of Proteome Research, 2015, 14, 224-235.	1.8	71
54	Integrative Analysis of Transcriptomics, Proteomics, and Metabolomics Data of White Adipose and Liver Tissue of High-Fat Diet and Rosiglitazone-Treated Insulin-Resistant Mice Identified Pathway Alterations and Molecular Hubs. Journal of Proteome Research, 2014, 13, 5592-5602.	1.8	51

DAVID MEIERHOFER

#	Article	IF	CITATIONS
55	Transcriptomics assisted proteomic analysis of <i>Nicotiana occidentalis</i> infected by <i>Candidatus</i> Phytoplasma mali strain AT. Proteomics, 2014, 14, 1882-1889.	1.3	39
56	A Y2H-seq approach defines the human protein methyltransferase interactome. Nature Methods, 2013, 10, 339-342.	9.0	99
57	Protein Sets Define Disease States and Predict In Vivo Effects of Drug Treatment. Molecular and Cellular Proteomics, 2013, 12, 1965-1979.	2.5	29
58	Comprehensive proteomic datasets for studying adipocyte–macrophage cell–cell communication. Proteomics, 2013, 13, 3424-3428.	1.3	7
59	Lipoic Acid Synthetase Deficiency Causes Neonatal-Onset Epilepsy, Defective Mitochondrial Energy Metabolism, and Glycine Elevation. American Journal of Human Genetics, 2011, 89, 792-797.	2.6	104
60	Global Analysis of Ubiquitination. Neuromethods, 2011, , 197-209.	0.2	0
61	Quantitative Analysis of global Ubiquitination in HeLa Cells by Mass Spectrometry. Journal of Proteome Research, 2008, 7, 4566-4576.	1.8	182
62	Tandem Affinity Purification Combined with Mass Spectrometry to Identify Components of Protein Complexes. Methods in Molecular Biology, 2008, 439, 309-326.	0.4	33
63	Loss of Complex I due to Mitochondrial DNA Mutations in Renal Oncocytoma. Clinical Cancer Research, 2008, 14, 2270-2275.	3.2	154
64	Platelet transfusion can mimic somatic mtDNA mutations. Leukemia, 2006, 20, 362-363.	3.3	11
65	Multiplex primer extension analysis for rapid detection of major European mitochondrial haplogroups. Electrophoresis, 2006, 27, 3864-3868.	1.3	21
66	Mitochondrial DNA mutations in renal cell carcinomas revealed no general impact on energy metabolism. British Journal of Cancer, 2006, 94, 268-274.	2.9	58
67	Rapid screening of the entire mitochondrial DNA for low-level heteroplasmic mutations. Mitochondrion, 2005, 5, 282-296.	1.6	43
68	Decrease of mitochondrial DNA content and energy metabolism in renal cell carcinoma. Carcinogenesis, 2004, 25, 1005-1010.	1.3	144
69	Severe depletion of mitochondrial DNA in spinal muscular atrophy. Acta Neuropathologica, 2003, 105, 245-251.	3.9	72