## Peter J Oefner

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3113331/publications.pdf

Version: 2024-02-01

220 papers

25,056 citations

9254 74 h-index 152 g-index

224 all docs

224 docs citations

times ranked

224

29183 citing authors

#	Article	IF	CITATIONS
1	Familial Hemiplegic Migraine and Episodic Ataxia Type-2 Are Caused by Mutations in the Ca2+ Channel Gene CACNL1A4. Cell, 1996, 87, 543-552.	13.5	2,287
2	LDHA-Associated Lactic Acid Production Blunts Tumor Immunosurveillance by T and NK Cells. Cell Metabolism, 2016, 24, 657-671.	7.2	1,126
3	CD133+ and CD133â° Glioblastoma-Derived Cancer Stem Cells Show Differential Growth Characteristics and Molecular Profiles. Cancer Research, 2007, 67, 4010-4015.	0.4	1,027
4	Y chromosome sequence variation and the history of human populations. Nature Genetics, 2000, 26, 358-361.	9.4	935
5	The Genetic Legacy of Paleolithic Homo sapiens sapiens in Extant Europeans: A Y Chromosome Perspective. Science, 2000, 290, 1155-1159.	6.0	783
6	Denaturing high-performance liquid chromatography: A review. Human Mutation, 2001, 17, 439-474.	1.1	674
7	Detection of Numerous Y Chromosome Biallelic Polymorphisms by Denaturing High-Performance Liquid Chromatography. Genome Research, 1997, 7, 996-1005.	2.4	617
8	Deficiency in glutamine but not glucose induces MYC-dependent apoptosis in human cells. Journal of Cell Biology, 2007, 178, 93-105.	2.3	599
9	Dissecting the architecture of a quantitative trait locus in yeast. Nature, 2002, 416, 326-330.	13.7	524
10	Systematic screen for human disease genes in yeast. Nature Genetics, 2002, 31, 400-404.	9.4	503
11	Metagenomic Analysis of the Stool Microbiome in Patients Receiving Allogeneic Stem Cell Transplantation: Loss of Diversity Is Associated with Use of Systemic Antibiotics and More Pronounced in Gastrointestinal Graft-versus-Host Disease. Biology of Blood and Marrow Transplantation, 2014, 20, 640-645.	2.0	444
12	The extent of linkage disequilibrium in Arabidopsis thaliana. Nature Genetics, 2002, 30, 190-193.	9.4	425
13	The Role of Selection in the Evolution of Human Mitochondrial Genomes. Genetics, 2006, 172, 373-387.	1.2	395
14	Y-Chromosome Evidence for a Northward Migration of Modern Humans into Eastern Asia during the Last Ice Age. American Journal of Human Genetics, 1999, 65, 1718-1724.	2.6	394
15	Origin, Diffusion, and Differentiation of Y-Chromosome Haplogroups E and J: Inferences on the Neolithization of Europe and Later Migratory Events in the Mediterranean Area. American Journal of Human Genetics, 2004, 74, 1023-1034.	2.6	345
16	Blind Analysis of Denaturing High-Performance Liquid Chromatography as a Tool for Mutation Detection. Genomics, 1998, 52, 44-49.	1.3	334
17	Lactic Acid and Acidification Inhibit TNF Secretion and Glycolysis of Human Monocytes. Journal of Immunology, 2010, 184, 1200-1209.	0.4	325
18	A Back Migration from Asia to Sub-Saharan Africa Is Supported by High-Resolution Analysis of Human Y-Chromosome Haplotypes. American Journal of Human Genetics, 2002, 70, 1197-1214.	2.6	318

#	Article	IF	Citations
19	Excavating Y-chromosome haplotype strata in Anatolia. Human Genetics, 2004, 114, 127-148.	1.8	318
20	African Origin of Modern Humans in East Asia: A Tale of 12,000 Y Chromosomes. Science, 2001, 292, 1151-1153.	6.0	310
21	GLUT1 Expression Is Increased in Hepatocellular Carcinoma and Promotes Tumorigenesis. American Journal of Pathology, 2009, 174, 1544-1552.	1.9	283
22	Genome-wide mapping with biallelic markers in Arabidopsis thaliana. Nature Genetics, 1999, 23, 203-207.	9.4	260
23	Phylogeography of Y-Chromosome Haplogroup I Reveals Distinct Domains of Prehistoric Gene Flow in Europe. American Journal of Human Genetics, 2004, 75, 128-137.	2.6	256
24	Srebp-controlled glucose metabolism is essential for NK cell functional responses. Nature Immunology, 2017, 18, 1197-1206.	7.0	249
25	Genome sequencing and comparative analysis of <i>Saccharomyces cerevisiae</i> strain YJM789. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 12825-12830.	3.3	240
26	Transcriptional Profiles of CD133+ and CD133â^' Glioblastoma-Derived Cancer Stem Cell Lines Suggest Different Cells of Origin. Cancer Research, 2010, 70, 2030-2040.	0.4	237
27	High-Resolution Analysis of Human Y-Chromosome Variation Shows a Sharp Discontinuity and Limited Gene Flow between Northwestern Africa and the Iberian Peninsula. American Journal of Human Genetics, 2001, 68, 1019-1029.	2.6	234
28	A Simple Procedure for the Analysis of Single Nucleotide Polymorphisms Facilitates Map-Based Cloning in Arabidopsis. Plant Physiology, 2000, 124, 1483-1492.	2.3	227
29	Melanesian and Asian Origins of Polynesians: mtDNA and Y Chromosome Gradients Across the Pacific. Molecular Biology and Evolution, 2006, 23, 2234-2244.	<b>3.</b> 5	216
30	Denaturing High-Performance Liquid Chromatography Detects Reliably BRCA1 and BRCA2 Mutations. Genomics, 1999, 62, 369-376.	1.3	214
31	Transformation of follicular lymphoma to diffuse large-cell lymphoma: Alternative patterns with increased or decreased expression of c-myc and its regulated genes. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 8886-8891.	3.3	204
32	Revealing the prehistoric settlement of Australia by Y chromosome and mtDNA analysis. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 8726-8730.	3.3	204
33	Lactate promotes glioma migration by TGF-β2–dependent regulation of matrix metalloproteinase-2. Neuro-Oncology, 2009, 11, 368-380.	0.6	204
34	Metabolite extraction from adherently growing mammalian cells for metabolomics studies: optimization of harvesting and extraction protocols. Analytical and Bioanalytical Chemistry, 2011, 399, 1127-1139.	1.9	200
35	Origins and Divergence of the Roma (Gypsies). American Journal of Human Genetics, 2001, 69, 1314-1331.	2.6	188
36	Microbiota Disruption Induced by Early Use of Broad-Spectrum Antibiotics Is an Independent Risk Factor of Outcome after Allogeneic Stem Cell Transplantation. Biology of Blood and Marrow Transplantation, 2017, 23, 845-852.	2.0	183

#	Article	IF	CITATIONS
37	Integrative Analysis of the Mitochondrial Proteome in Yeast. PLoS Biology, 2004, 2, e160.	2.6	181
38	Delaying aging and the aging-associated decline in protein homeostasis by inhibition of tryptophan degradation. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 14912-14917.	3.3	180
39	Advances in amino acid analysis. Analytical and Bioanalytical Chemistry, 2009, 393, 445-452.	1.9	168
40	Third-party fecal microbiota transplantation following allo-HCT reconstitutes microbiome diversity. Blood Advances, 2018, 2, 745-753.	2.5	167
41	Rare, Evolutionarily Unlikely Missense Substitutions in ATM Confer Increased Risk of Breast Cancer. American Journal of Human Genetics, 2009, 85, 427-446.	2.6	165
42	Reduced Y-Chromosome, but Not Mitochondrial DNA, Diversity in Human Populations from West New Guinea. American Journal of Human Genetics, 2003, 72, 281-302.	2.6	160
43	Double genetic disruption of lactate dehydrogenases A and B is required to ablate the "Warburg effect―restricting tumor growth to oxidative metabolism. Journal of Biological Chemistry, 2018, 293, 15947-15961.	1.6	160
44	A Metabolome-Wide Association Study of Kidney Function and Disease in the General Population. Journal of the American Society of Nephrology: JASN, 2016, 27, 1175-1188.	3.0	159
45	Automated GC–MS analysis of free amino acids in biological fluids. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2008, 870, 222-232.	1.2	158
46	Urinary amino acid analysis: A comparison of iTRAQ®–LC–MS/MS, GC–MS, and amino acid analyzer. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2009, 877, 1838-1846.	1.2	150
47	Task-specific expression of the foraging gene in harvester ants. Molecular Ecology, 2005, 14, 813-818.	2.0	147
48	Sequential Elimination of Major-Effect Contributors Identifies Additional Quantitative Trait Loci Conditioning High-Temperature Growth in Yeast. Genetics, 2008, 180, 1661-1670.	1.2	145
49	Y-chromosomal evidence of a pastoralist migration through Tanzania to southern Africa. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 10693-10698.	3.3	133
50	Premature Termination Mutations in FBN1: Distinct Effects on Differential Allelic Expression and on Protein and Clinical Phenotypes. American Journal of Human Genetics, 2002, 71, 223-237.	2.6	131
51	Quantitative profiling of tryptophan metabolites in serum, urine, and cell culture supernatants by liquid chromatography–tandem mass spectrometry. Analytical and Bioanalytical Chemistry, 2011, 401, 3249-3261.	1.9	130
52	The German Chronic Kidney Disease (GCKD) study: design and methods. Nephrology Dialysis Transplantation, 2012, 27, 1454-1460.	0.4	127
53	Urinary Metabolite Quantification Employing 2D NMR Spectroscopy. Analytical Chemistry, 2008, 80, 9288-9297.	3.2	123
54	NMR Metabolomic Analysis of Dairy Cows Reveals Milk Glycerophosphocholine to Phosphocholine Ratio as Prognostic Biomarker for Risk of Ketosis. Journal of Proteome Research, 2012, 11, 1373-1381.	1.8	122

#	Article	IF	Citations
55	Rapid and Accurate Sizing of DNA Fragments by Ion-Pair Chromatography on Alkylated Nonporous Poly(styrene-divinylbenzene) Particles. Analytical Chemistry, 1995, 67, 578-585.	3.2	118
56	Contributions of ATM mutations to familial breast and ovarian cancer. Cancer Research, 2003, 63, 3325-33.	0.4	113
57	Comprehensive two-dimensional gas chromatography in metabolomics. Analytical and Bioanalytical Chemistry, 2012, 402, 1993-2013.	1.9	104
58	Capillary zone electrophoresis and micellar electrokinetic chromatography of 4-aminobenzonitrile carbohydrate derivatives. Electrophoresis, 1994, 15, 941-952.	1.3	101
59	Genotyping single nucleotide polymorphisms by primer extension and high performance liquid chromatography. Human Genetics, 1999, 104, 89-93.	1.8	101
60	Genetic studies of urinary metabolites illuminate mechanisms of detoxification and excretion in humans. Nature Genetics, 2020, 52, 167-176.	9.4	101
61	Denaturing HPLC-Identified Novel FBN1 Mutations, Polymorphisms, and Sequence Variants in Marfan Syndrome and Related Connective Tissue Disorders. Genetic Testing and Molecular Biomarkers, 1997, 1, 237-242.	1.7	99
62	A Novel Y-Chromosome Variant Puts an Upper Limit on the Timing of First Entry into the Americas. American Journal of Human Genetics, 2003, 73, 700-705.	2.6	99
63	Mistargeting of Peroxisomal EHHADH and Inherited Renal Fanconi's Syndrome. New England Journal of Medicine, 2014, 370, 129-138.	13.9	99
64	High-resolution liquid chromatography of DNA fragments on non-porous poly(styrene-divinylbenzene) particles. Nucleic Acids Research, 1993, 21, 1061-1066.	6.5	97
65	Global sequence diversity of BRCA2: analysis of 71 breast cancer families and 95 control individuals of worldwide populations [published erratum appears in Hum Mol Genet 1999 Apr;8(4):717-9]. Human Molecular Genetics, 1999, 8, 413-423.	1.4	97
66	D-2-hydroxyglutarate interferes with HIF-1 $\hat{l}$ ± stability skewing T-cell metabolism towards oxidative phosphorylation and impairing Th17 polarization. OncoImmunology, 2018, 7, e1445454.	2.1	97
67	Global Analysis of ATM Polymorphism Reveals Significant Functional Constraint. American Journal of Human Genetics, 2001, 69, 396-412.	2.6	93
68	Capillary electrophoresis of carbohydrates. Glycobiology, 1994, 4, 397-412.	1.3	87
69	New Aspects of an Old Drug – Diclofenac Targets MYC and Glucose Metabolism in Tumor Cells. PLoS ONE, 2013, 8, e66987.	1.1	86
70	High-resolution capillary electrophoretic analysis of DNA in free solution. Electrophoresis, 1992, 13, 18-31.	1.3	85
71	Validation of microarray-based resequencing of 93 worldwide mitochondrial genomes. Human Mutation, 2009, 30, 115-122.	1.1	83
72	Polyol Pathway Links Glucose Metabolism to the Aggressiveness of Cancer Cells. Cancer Research, 2018, 78, 1604-1618.	0.4	83

#	Article	IF	CITATIONS
73	Capillary zone electrophoresis ofp-aminobenzoic acid derivatives of aldoses, ketoses and uronic acids. Electrophoresis, 1993, 14, 1004-1010.	1.3	81
74	BRCA1-related breast cancer in Austrian breast and ovarian cancer families: SpecificBRCA1 mutations and pathological characteristics., 1998, 77, 354-360.		81
75	Monolithic capillary columns for liquid chromatography–electrospray ionization mass spectrometry in proteomic and genomic research. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2002, 782, 111-125.	1.2	79
76	Diclofenac inhibits lactate formation and efficiently counteracts local immune suppression in a murine glioma model. International Journal of Cancer, 2013, 132, 843-853.	2.3	77
77	Reduced genetic structure of the Iberian peninsula revealed by Y-chromosome analysis: implications for population demography. European Journal of Human Genetics, 2004, 12, 855-863.	1.4	76
78	Frequentist Estimation of Coalescence Times From Nucleotide Sequence Data Using a Tree-Based Partition. Genetics, 2002, 161, 447-459.	1.2	76
79	Metabolic plasticity of human T cells: Preserved cytokine production under glucose deprivation or mitochondrial restriction, but 2â€deoxyâ€glucose affects effector functions. European Journal of Immunology, 2015, 45, 2504-2516.	1.6	75
80	Proteome analysis of mitochondrial outer membrane from Neurospora crassa. Proteomics, 2006, 6, 72-80.	1.3	74
81	Conservation of the class I beta-tubulin gene in human populations and lack of mutations in lung cancers and paclitaxel-resistant ovarian cancers. Molecular Cancer Therapeutics, 2002, 1, 215-25.	1.9	72
82	Maori origins, Y-chromosome haplotypes and implications for human history in the Pacific. Human Mutation, 2001, 17, 271-280.	1.1	70
83	Direct and tumor microenvironment mediated influences of 5′â€deoxyâ€5′â€(methylthio)adenosine on tum progression of malignant melanoma. Journal of Cellular Biochemistry, 2009, 106, 210-219.	ior 1.2	70
84	High-Accuracy DNA Sequence Variation Screening by DHPLC. BioTechniques, 2000, 29, 1084-1092.	0.8	69
85	Capillary electrophoretic analysis of flavonoids. Electrophoresis, 1992, 13, 35-38.	1.3	68
86	Identification by Denaturing High-Performance Liquid Chromatography of Numerous Polymorphisms in a Candidate Region for Multiple Sclerosis Susceptibility. Genomics, 1999, 56, 247-253.	1.3	66
87	Reconstruction of patrilineages and matrilineages of Samaritans and other Israeli populations from Y-Chromosome and mitochondrial DNA sequence Variation. Human Mutation, 2004, 24, 248-260.	1.1	66
88	First-generation SNP/InDel markers tagging loci for pathogen resistance in the potato genome. Plant Biotechnology Journal, 2003, 1, 399-410.	4.1	63
89	Development of a quantitative, validated Capillary electrophoresisâ€time of flight – mass spectrometry method with integrated highâ€confidence analyte identification for metabolomics. Electrophoresis, 2008, 29, 2203-2214.	1.3	63
90	Genotyping of SNPs in a Polyploid Genome by Pyrosequencingâ,,¢. BioTechniques, 2002, 32, 592-603.	0.8	62

#	Article	IF	CITATIONS
91	An Automated Hydrodynamic Process for Controlled, Unbiased DNA Shearing. Genome Research, 1998, 8, 848-855.	2.4	61
92	Cloning of the Arabidopsis RSF1 Gene by Using a Mapping Strategy Based on High-Density DNA Arrays and Denaturing High-Performance Liquid Chromatography. Plant Cell, 2000, 12, 2485-2498.	3.1	61
93	Detection of autosomal dominant polycystic kidney disease by NMR spectroscopic fingerprinting of urine. Kidney International, 2011, 79, 1244-1253.	2.6	59
94	Extracellular Citrate Affects Critical Elements of Cancer Cell Metabolism and Supports Cancer Development <i>In Vivo</i> . Cancer Research, 2018, 78, 2513-2523.	0.4	59
95	High-performance liquid chromatographic separation of detritylated oligonucleotides on highly cross-linked poly-(styrene-divinylbenzene) particles. Journal of Chromatography A, 1992, 599, 113-118.	1.8	58
96	Integrative Normalization and Comparative Analysis for Metabolic Fingerprinting by Comprehensive Two-Dimensional Gas Chromatographyâ°'Time-of-Flight Mass Spectrometry. Analytical Chemistry, 2009, 81, 5731-5739.	3.2	56
97	Comparative study of capillary zone electrphoresis and high-performance liquid chromatography in the analysis of oligonucleotides and DNA. Journal of Chromatography A, 1992, 625, 331-340.	1.8	55
98	Down-Regulation of Methylthioadenosine Phosphorylase (MTAP) Induces Progression of Hepatocellular Carcinoma via Accumulation of 5′-Deoxy-5′-Methylthioadenosine (MTA). American Journal of Pathology, 2011, 178, 1145-1152.	1.9	54
99	Rapid Quantification of Gene Expression by Competitive RT-PCR and Ion-Pair Reversed-Phase HPLC. BioTechniques, 1996, 20, 250-257.	0.8	53
100	Capillary electrophoresis and column chromatography in biomedical chiral amino acid analysis. Analytical and Bioanalytical Chemistry, 2009, 394, 695-706.	1.9	53
101	Comparison of derivatization and chromatographic methods for GC–MS analysis of amino acid enantiomers in physiological samples. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2010, 878, 1103-1112.	1.2	53
102	Improved enantiomer resolution and quantification of free d-amino acids in serum and urine by comprehensive two-dimensional gas chromatography–time-of-flight mass spectrometry. Journal of Chromatography A, 2011, 1218, 4537-4544.	1.8	53
103	Glycine Amidinotransferase (GATM), Renal Fanconi Syndrome, and Kidney Failure. Journal of the American Society of Nephrology: JASN, 2018, 29, 1849-1858.	3.0	53
104	Mitochondrial DNA copy number is associated with mortality and infections in a large cohort of patients with chronic kidney disease. Kidney International, 2019, 96, 480-488.	2.6	53
105	Mutation detection by capillary denaturing high-performance liquid chromatography using monolithic columns. Journal of Proteomics, 2001, 47, 5-19.	2.4	52
106	Reduced Expression of Fibroblast Growth Factor Receptor 2IIIb in Hepatocellular Carcinoma Induces a More Aggressive Growth. American Journal of Pathology, 2010, 176, 1433-1442.	1.9	52
107	Tryptophan catabolism is associated with acute GVHD after human allogeneic stem cell transplantation and indicates activation of indoleamine 2,3-dioxygenase. Blood, 2011, 118, 6971-6974.	0.6	52
108	Mutational analyses of BRCA1 and BRCA2 in Ashkenazi and non-Ashkenazi Jewish women with familial breast and ovarian cancer. Human Mutation, 2000, 16, 491-501.	1.1	50

#	Article	IF	Citations
109	Hypoxiaâ€inducible protein 2 Hig2/Hilpda mediates neutral lipid accumulation in macrophages and contributes to atherosclerosis in apolipoprotein E–deficient mice. FASEB Journal, 2017, 31, 4971-4984.	0.2	50
110	Suppressive effects of tumor cell-derived $5\hat{a}\in^2$ -deoxy- $5\hat{a}\in^2$ -methylthioadenosine on human T cells. Oncolmmunology, 2016, 5, e1184802.	2.1	48
111	Regulation and function of the atypical cadherin FAT1 in hepatocellular carcinoma. Carcinogenesis, 2014, 35, 1407-1415.	1.3	46
112	A decade of high-resolution liquid chromatography of nucleic acids on styrene–divinylbenzene copolymers. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2002, 782, 27-55.	1.2	45
113	TwoATM variants and breast cancer risk. Human Mutation, 2005, 25, 594-595.	1.1	44
114	Deficient Tryptophan Catabolism along the Kynurenine Pathway Reveals That the Epididymis Is in a Unique Tolerogenic State. Journal of Biological Chemistry, 2011, 286, 8030-8042.	1.6	44
115	Modeling the temporal interplay of molecular signaling and gene expression by using dynamic nested effects models. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 6447-6452.	3.3	43
116	Comparison of serum versus plasma collection in gas chromatography – Mass spectrometryâ€based metabolomics. Electrophoresis, 2010, 31, 2365-2373.	1.3	43
117	Performance Evaluation of Gas Chromatography–Atmospheric Pressure Chemical Ionization–Time-of-Flight Mass Spectrometry for Metabolic Fingerprinting and Profiling. Analytical Chemistry, 2011, 83, 7514-7522.	3.2	43
118	Quantification of intermediates of the methionine and polyamine metabolism by liquid chromatography–tandem mass spectrometry in cultured tumor cells and liver biopsies. Journal of Chromatography A, 2010, 1217, 3282-3288.	1.8	39
119	Metformin inhibits proliferation and migration of glioblastoma cells independently of TGF-Î <sup>2</sup> 2. Cell Cycle, 2016, 15, 1755-1766.	1.3	39
120	Genome-Wide Association Studies of Metabolites in Patients with CKD Identify Multiple Loci and Illuminate Tubular Transport Mechanisms. Journal of the American Society of Nephrology: JASN, 2018, 29, 1513-1524.	3.0	39
121	Isotachophoretic analysis of flavonoids and phenolcarboxylic acids of relevance to phytopharmaceutical industry. Journal of Chromatography A, 1991, 559, 499-504.	1.8	35
122	High-performance liquid chromatographic determination of free polyamines in human seminal plasma. Clinica Chimica Acta, 1992, 205, 11-18.	0.5	35
123	Collagen XVI Induces Expression of MMP9 via Modulation of AP-1 Transcription Factors and Facilitates Invasion of Oral Squamous Cell Carcinoma. PLoS ONE, 2014, 9, e86777.	1.1	35
124	Experimental test of a method for determining causal connectivities of species in reactions. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 1494-1498.	3.3	34
125	Performance Evaluation of Algorithms for the Classification of Metabolic <sup>1</sup> H NMR Fingerprints. Journal of Proteome Research, 2012, 11, 6242-6251.	1.8	33
126	Data Normalization of <sup>1</sup> H NMR Metabolite Fingerprinting Data Sets in the Presence of Unbalanced Metabolite Regulation. Journal of Proteome Research, 2015, 14, 3217-3228.	1.8	32

#	Article	IF	Citations
127	Quantitative analysis of $5\hat{a}\in^2$ -deoxy- $5\hat{a}\in^2$ -methylthioadenosine in melanoma cells by liquid chromatography-stable isotope ratio tandem mass spectrometry. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2008, 876, 123-128.	1.2	30
128	Distinct metabolic differences between various human cancer and primary cells. Electrophoresis, 2013, 34, 2836-2847.	1.3	29
129	Surface-charge reversed capillary zone electrophoresis of inorganic and organic anions. Electrophoresis, 1995, 16, 46-56.	1.3	28
130	Capillary electrophoretic determination of the component monosaccharides in hemicelluloses. Fresenius' Journal of Analytical Chemistry, 1994, 348, 825-831.	1.5	26
131	Early changes in the liverâ€soluble proteome from mice fed a nonalcoholic steatohepatitis inducing diet. Proteomics, 2012, 12, 1437-1451.	1.3	26
132	Visceral adipose tissue but not subcutaneous adipose tissue is associated with urine and serum metabolites. PLoS ONE, 2017, 12, e0175133.	1.1	26
133	Systematic Evaluation of Non-Uniform Sampling Parameters in the Targeted Analysis of Urine Metabolites by 1H,1H 2D NMR Spectroscopy. Scientific Reports, 2018, 8, 4249.	1.6	26
134	Applicability of tandem mass spectrometry to the automated comparative sequencing of long-chain oligonucleotides. Journal of the American Society for Mass Spectrometry, 2004, 15, 510-522.	1.2	25
135	Optimized suppression of adducts in polymerase chain reaction products for semi-quantitative SNP genotyping by liquid chromatography-mass spectrometry. Journal of the American Society for Mass Spectrometry, 2004, 15, 1897-1906.	1.2	25
136	High CD206 levels in Hodgkin lymphomaâ€educated macrophages are linked to matrixâ€remodeling and lymphoma dissemination. Molecular Oncology, 2020, 14, 571-589.	2.1	25
137	Capillary Array High-Performance Liquid Chromatography of Nucleic Acids and Proteins. Analytical Chemistry, 2002, 74, 4688-4693.	3.2	24
138	Mutation scanning by ion-pair reversed-phase high-performance liquid chromatography-electrospray ionization mass spectrometry (ICEMS). Human Mutation, 2003, 21, 86-95.	1.1	24
139	Comparison of two algorithmic data processing strategies for metabolic fingerprinting by comprehensive two-dimensional gas chromatography–time-of-flight mass spectrometry. Journal of Chromatography A, 2011, 1218, 7031-8.	1.8	24
140	Correlations between Milk and Plasma Levels of Amino and Carboxylic Acids in Dairy Cows. Journal of Proteome Research, 2013, 12, 5223-5232.	1.8	24
141	Continuous Water Infusion Enhances Atmospheric Pressure Chemical Ionization of Methyl Chloroformate Derivatives in Gas Chromatography Coupled to Time-of-Flight Mass Spectrometry-Based Metabolomics. Analytical Chemistry, 2014, 86, 9186-9195.	3.2	24
142	Quantification of Metabolites by NMR Spectroscopy in the Presence of Protein. Journal of Proteome Research, 2017, 16, 1784-1796.	1.8	24
143	Temperature-Modulated Array High-Performance Liquid Chromatography. Genome Research, 2001, 11, 1944-1951.	2.4	23
144	LEF1 supports metastatic brain colonization by regulating glutathione metabolism and increasing ROS resistance in breast cancer. International Journal of Cancer, 2020, 146, 3170-3183.	2.3	23

#	Article	IF	CITATIONS
145	Distinct von Hippel-Lindau gene and hypoxia-regulated alterations in gene and protein expression patterns of renal cell carcinoma and their effects on metabolism. Oncotarget, 2015, 6, 11395-11406.	0.8	23
146	Polymorphisms within the <i> APOBR &lt; /i &gt; gene are highly associated with milk levels of prognostic ketosis biomarkers in dairy cows. Physiological Genomics, 2015, 47, 129-137.</i>	1.0	22
147	Causal Modeling of Cancer-Stromal Communication Identifies PAPPA as a Novel Stroma-Secreted Factor Activating NFκB Signaling in Hepatocellular Carcinoma. PLoS Computational Biology, 2015, 11, e1004293.	1.5	22
148	Comprehensive Metaboproteomics of Burkitt's and Diffuse Large B-Cell Lymphoma Cell Lines and Primary Tumor Tissues Reveals Distinct Differences in Pyruvate Content and Metabolism. Journal of Proteome Research, 2017, 16, 1105-1120.	1.8	22
149	Incidence of Arterial Hypotension in Patients Receiving Peroral or Continuous Intra-arterial Nimodipine After Aneurysmal or Perimesencephalic Subarachnoid Hemorrhage. Neurocritical Care, 2019, 31, 32-39.	1.2	22
150	Evaluation of dilution and normalization strategies to correct for urinary output in HPLC-HRTOFMS metabolomics. Analytical and Bioanalytical Chemistry, 2016, 408, 8483-8493.	1.9	21
151	Biological and clinical significance of tryptophan-catabolizing enzymes in cutaneous T-cell lymphomas. Oncolmmunology, 2017, 6, e1273310.	2.1	21
152	The association between acute graft-versus-host disease and antimicrobial peptide expression in the gastrointestinal tract after allogeneic stem cell transplantation. PLoS ONE, 2017, 12, e0185265.	1.1	21
153	Younger birth cohort correlates with higher breast and ovarian cancer risk in EuropeanBRCA1 mutation carriers. Human Mutation, 2005, 26, 583-589.	1.1	20
154	Topical Diclofenac Reprograms Metabolism and Immune Cell Infiltration in Actinic Keratosis. Frontiers in Oncology, 2019, 9, 605.	1.3	20
155	Kynurenine induces T cell fat catabolism and has limited suppressive effects in vivo. EBioMedicine, 2021, 74, 103734.	2.7	20
156	MetaboQuant: a tool combining individual peak calibration and outlier detection for accurate metabolite quantification in 1D $<$ sup>1 $<$ sup>H and $<$ sup>1 $<$ sup>H- $<$ sup>13 $<$ sup>C HSQC NMR spectra. BioTechniques, 2013, 54, 251-256.	0.8	19
157	Quantification of alternatively spliced RUSH mRNA isoforms by QRT-PCR and IP-RP-HPLC analysis: a new approach to measuring regulated splicing efficiency. Gene, 1997, 198, 1-4.	1.0	18
158	Conservation of theRB1gene in human and primates. Human Mutation, 2005, 25, 396-409.	1.1	18
159	Identification of Plasma Metabolites Prognostic of Acute Kidney Injury after Cardiac Surgery with Cardiopulmonary Bypass. Journal of Proteome Research, 2015, 14, 2897-2905.	1.8	18
160	De novo polyamine synthesis supports metabolic and functional responses in activated murine NK cells. European Journal of Immunology, 2021, 51, 91-102.	1.6	18
161	Multiplex Analysis of Single-Nucleotide Extension Products on a 16-Capillary, Denaturing, High-Performance Liquid Chromatography Array. Genomics, 2002, 79, 793-798.	1.3	17
162	Characterization of the breast cancer associated ATM7271T>G (V2424G) mutation by gene expression profiling. Genes Chromosomes and Cancer, 2006, 45, 1169-1181.	1.5	17

#	Article	IF	CITATIONS
163	Enhanced metabolite profiling using a redesigned atmospheric pressure chemical ionization source for gas chromatography coupled to high-resolution time-of-flight mass spectrometry. Analytical and Bioanalytical Chemistry, 2015, 407, 6669-6680.	1.9	17
164	Quantification and 13C-Tracer analysis of total reduced glutathione by HPLC-QTOFMS/MS. Analytica Chimica Acta, 2019, 1080, 127-137.	2.6	17
165	Polymerase chain reaction fidelity and denaturing high-performance liquid chromatography. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2002, 782, 105-110.	1.2	16
166	Identifying new candidate genes for hereditary facial paresis on chromosome 3q21–q22 by RNA in situ hybridization in mouse. Genomics, 2005, 86, 55-67.	1.3	16
167	Current Experimental, Bioinformatic and Statistical Methods used in NMR Based Metabolomics. Current Metabolomics, 2013, 1, 253-268.	0.5	16
168	Results from the German Chronic Kidney Disease (GCKD) study support association of relative telomere length with mortality in a large cohort of patients with moderate chronic kidney disease. Kidney International, 2020, 98, 488-497.	2.6	16
169	Isotachophoretic separation of organic acids in biological fluids. Journal of Chromatography A, 1990, 516, 251-262.	1.8	15
170	Nâ€cadherin promoter polymorphisms and risk of osteoarthritis. FASEB Journal, 2014, 28, 683-691.	0.2	15
171	Scale-Invariant Biomarker Discovery in Urine and Plasma Metabolite Fingerprints. Journal of Proteome Research, 2017, 16, 3596-3605.	1.8	15
172	Assessment of urinary 3-indoxyl sulfate as a marker for gut microbiota diversity and abundance of <i>Clostridiales</i> . Gut Microbes, 2019, 10, 133-141.	4.3	15
173	A Novel Metabolic Signature To Predict the Requirement of Dialysis or Renal Transplantation in Patients with Chronic Kidney Disease. Journal of Proteome Research, 2019, 18, 1796-1805.	1.8	15
174	Sequence variation and the biological function of genes: methodological and biological considerations. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2002, 782, 3-25.	1.2	14
175	Fisher's theorems for multivariable, time- and space-dependent systems, with applications in population genetics and chemical kinetics. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 9848-9853.	3.3	14
176	Central European BRCA2 mutation carriers: Birth cohort status correlates with onset of breast cancer. Maturitas, 2014, 77, 68-72.	1.0	14
177	Shear Force Processing of Lipoaspirates for Stem Cell Enrichment Does Not Affect Secretome of Human Cells Detected by Mass Spectrometry In Vitro. Plastic and Reconstructive Surgery, 2020, 146, 749e-758e.	0.7	14
178	Potential biomarkers to predict outcome of faecal microbiota transfer for recurrent Clostridioides difficile infection. Digestive and Liver Disease, 2019, 51, 944-951.	0.4	13
179	LDHB Overexpression Can Partially Overcome T Cell Inhibition by Lactic Acid. International Journal of Molecular Sciences, 2022, 23, 5970.	1.8	13
180	Combining genome sequences and new technologies for dissecting the genetics of complex phenotypes. Trends in Plant Science, 2000, 5, 397-401.	4.3	12

#	Article	IF	CITATIONS
181	FEM1A is a candidate gene for polycystic ovary syndrome. Gynecological Endocrinology, 2005, 21, 330-335.	0.7	11
182	Analysis of FAS (CD95) Gene Mutations in Higher-Grade Transformation of Follicle Center Lymphoma. Leukemia and Lymphoma, 2003, 44, 1317-1323.	0.6	10
183	Mutation of the ATM Gene is Not Involved in the Pathogenesis of Either Follicle Center Lymphoma or its Transformation to Higher-grade Lymphoma. Leukemia and Lymphoma, 2002, 43, 1079-1085.	0.6	9
184	Kinetic laws, phase–phase expansions, renormalization group, and INR calibration. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 6465-6470.	3.3	9
185	Self-Reported Medication Use and Urinary Drug Metabolites in the German Chronic Kidney Disease (GCKD) Study. Journal of the American Society of Nephrology: JASN, 2021, 32, 2315-2329.	3.0	9
186	An R-Package for the Deconvolution and Integration of 1D NMR Data: MetaboDecon1D. Metabolites, 2021, 11, 452.	1.3	9
187	Cytokine-specific autoantibodies shape the gut microbiome in autoimmune polyendocrine syndrome type 1. Journal of Allergy and Clinical Immunology, 2021, 148, 876-888.	1.5	9
188	Population genetic implications from DNA polymorphism in random human genomic sequences. Human Mutation, 2002, 20, 209-217.	1.1	8
189	A serum microRNA sequence reveals fragile X protein pathology in amyotrophic lateral sclerosis. Brain, 2021, 144, 1214-1229.	3.7	8
190	Presenilin $1/\hat{A}$ -secretase modulates P-cadherin processing and influences cell adhesion in oral squamous cell carcinoma cell lines. Carcinogenesis, 2013, 34, 2622-2628.	1.3	7
191	Selenophosphate synthetase in the male accessory glands of an insect without selenoproteins. Journal of Insect Physiology, 2014, 71, 46-51.	0.9	7
192	Expression and Function of Methylthioadenosine Phosphorylase in Chronic Liver Disease. PLoS ONE, 2013, 8, e80703.	1.1	7
193	Random Activation Energy Model and Disordered Kinetics, from Static to Dynamic Disorder. Journal of Physical Chemistry B, 2005, 109, 21241-21257.	1.2	6
194	Hyphenated mass spectrometry in the analysis of the central carbon metabolism. Analytical and Bioanalytical Chemistry, 2008, 391, 895-898.	1.9	6
195	Variation This work was financially supported by the Austrian Science Fund (P-14133-PHY) and the National Institutes of Health (HG01932). The suggestion of Dr. Jeff van Ness from the Keck Graduate Institute of Applied Life Sciences in Claremont, CA, USA, to utilize butyldimethylamine for IP-RP-HPLC separations is gratefully acknowledged Angewandte Chemie - International Edition, 2001, 40.	7.2	6
196	3828 3830 Impact of lifestyle factors on preneoplastic changes in prophylactic oophorectomies of BRCA mutation carriers. European Journal of Cancer Prevention, 2012, 21, 199-204.	0.6	6
197	Incremental parameter evaluation from incomplete data with application to the population pharmacology of anticoagulants. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 4627-4632.	3.3	5
198	Robust Metabolite Quantification from J-Compensated 2D 1H-13C-HSQC Experiments. Metabolites, 2020, 10, 449.	1.3	5

#	Article	IF	Citations
199	Loss-Function Learning for Digital Tissue Deconvolution. Journal of Computational Biology, 2020, 27, 342-355.	0.8	5
200	Characterization of the Methylthioadenosine Phosphorylase Polymorphism rs7023954 - Incidence and Effects on Enzymatic Function in Malignant Melanoma. PLoS ONE, 2016, 11, e0160348.	1.1	5
201	Impact of separation capacity on the isotachophoretic analysis of organic acids in human seminal plasma and prostatic fluid. Electrophoresis, 1992, 13, 122-127.	1.3	4
202	Genotyping African haplotypes in ATM using a co-spotted single-base extension assay. Human Mutation, 2003, 22, 214-221.	1.1	4
203	Allelic loss analysis by denaturing high-performance liquid chromatography and electrospray ionization mass spectrometry. Human Mutation, 2007, 28, 303-311.	1.1	4
204	Associations between urinary 3-indoxyl sulfate, a gut microbiome-derived biomarker, and patient outcomes after intensive care unit admission. Journal of Critical Care, 2021, 63, 15-21.	1.0	4
205	DTD: An R Package for Digital Tissue Deconvolution. Journal of Computational Biology, 2020, 27, 386-389.	0.8	4
206	Acidic Microenvironments Found in Cutaneous Leishmania Lesions Curtail NO-Dependent Antiparasitic Macrophage Activity. Frontiers in Immunology, 2022, 13, 789366.	2.2	4
207	An isotachophoretic analysis of the interaction of bilirubin and biliverdin with bovine serum albumin. Electrophoresis, 1985, 6, 538-544.	1.3	3
208	Isotachophoretic Analysis of Organic Acids in Human Cerebrospinal Fluid and Serum. Annals of the New York Academy of Sciences, 1988, 529, 193-198.	1.8	3
209	Serotonin and tryptophan metabolites, autoantibodies and gut microbiome in APECED. Endocrine Connections, 2019, 8, 69-77.	0.8	3
210	Validation Study for Non-Invasive Prediction of IDH Mutation Status in Patients with Glioma Using In Vivo 1H-Magnetic Resonance Spectroscopy and Machine Learning. Cancers, 2022, 14, 2762.	1.7	3
211	The Microbiome and Allogeneic Stem Cell Transplantation. Current Stem Cell Reports, 2015, 1, 53-59.	0.7	2
212	Inducing anti-tumor cytokines and an immune response in melanoma by inhibition of MIA using the peptide AR71. European Journal of Dermatology, 2013, 23, 820-825.	0.3	2
213	Transition Event Statistics in Genetics and Disordered Kinetics. Theoretical Approaches for Extracting Rate Distributions from Experimental Dataâ€. Journal of Physical Chemistry B, 2006, 110, 18945-18952.	1.2	1
214	Editorial: Electrophoresis 13/2005. Electrophoresis, 2005, 26, 2493-2493.	1.3	0
215	Editorial: Electrophoresis 14/2005. Electrophoresis, 2005, 26, 2685-2685.	1.3	0
216	Bioanalysis: Developments and Trends. Electrophoresis, 2006, 27, 2527-2528.	1.3	0

## PETER J OEFNER

#	Article	IF	CITATIONS
217	Bioanalysis. Electrophoresis, 2007, 28, 1849-1850.	1.3	0
218	BIOANALYSIS. Electrophoresis, 2008, 29, 2447-2448.	1.3	0
219	Genetics and the History of The Samaritans: Y-Chromosomal Microsatellites and Genetic Affinity between Samaritans and Cohanim. Human Biology, 2013, 85, 825.	0.4	0
220	Principles of Systems Biology, No. 31. Cell Systems, 2018, 7, 133-135.	2.9	0