## **Richard D Beger**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9278829/publications.pdf Version: 2024-02-01

		117453	114278
107	4,528	34	63
papers	citations	h-index	g-index
112	112	112	6596
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Metabolomics enables precision medicine: "A White Paper, Community Perspective― Metabolomics, 2016, 12, 149.	1.4	434
2	CERAPP: Collaborative Estrogen Receptor Activity Prediction Project. Environmental Health Perspectives, 2016, 124, 1023-1033.	2.8	264
3	Summary recommendations for standardization and reporting of metabolic analyses. Nature Biotechnology, 2005, 23, 833-838.	9.4	261
4	Isolation of human intestinal bacteria metabolizing the natural isoflavone glycosides daidzin and genistin. Archives of Microbiology, 2000, 174, 422-428.	1.0	238
5	A Review of Applications of Metabolomics in Cancer. Metabolites, 2013, 3, 552-574.	1.3	217
6	Metabolomics approaches for discovering biomarkers of drug-induced hepatotoxicity and nephrotoxicity. Toxicology and Applied Pharmacology, 2010, 243, 154-166.	1.3	202
7	Use cases, best practice and reporting standards for metabolomics in regulatory toxicology. Nature Communications, 2019, 10, 3041.	5.8	131
8	Metabonomics evaluation of urine from rats given acute and chronic doses of acetaminophen using NMR and UPLC/MS. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2008, 871, 328-340.	1.2	115
9	Neuroprotective effect of the chemical chaperone, trehalose in a chronic MPTP-induced Parkinson's disease mouse model. NeuroToxicology, 2014, 44, 250-262.	1.4	103
10	Towards quality assurance and quality control in untargeted metabolomics studies. Metabolomics, 2019, 15, 4.	1.4	101
11	Metabonomics of acute kidney injury in children after cardiac surgery. Pediatric Nephrology, 2008, 23, 977-984.	0.9	89
12	Serum metabolomic profiles from patients with acute kidney injury: A pilot study. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2012, 893-894, 107-113.	1.2	81
13	Metabonomic models of human pancreatic cancer using 1D proton NMR spectra of lipids in plasma. Metabolomics, 2006, 2, 125-134.	1.4	80
14	Potential of extracellular microRNAs as biomarkers of acetaminophen toxicity in children. Toxicology and Applied Pharmacology, 2015, 284, 180-187.	1.3	73
15	Translational biomarkers of acetaminophen-induced acute liver injury. Archives of Toxicology, 2015, 89, 1497-1522.	1.9	72
16	Monitoring the health to disease continuum with global metabolic profiling and systems biology. Pharmacogenomics, 2006, 7, 1077-1086.	0.6	67
17	Targeted liquid chromatography–mass spectrometry analysis of serum acylcarnitines in acetaminophen toxicity in children. Biomarkers in Medicine, 2014, 8, 147-159.	0.6	62
18	Dissemination and analysis of the quality assurance (QA) and quality control (QC) practices of LC–MS based untargeted metabolomics practitioners. Metabolomics, 2020, 16, 113.	1.4	56

#	Article	IF	CITATIONS
19	Current Concepts in Pharmacometabolomics, Biomarker Discovery, and Precision Medicine. Metabolites, 2020, 10, 129.	1.3	56
20	Protein phi and psi dihedral restraints determined from multidimensional hypersurface correlations of backbone chemical shifts and their use in the determination of protein tertiary structures. Journal of Biomolecular NMR, 1997, 10, 129-142.	1.6	54
21	The Liver Toxicity Biomarker Study: Phase I Design and Preliminary Results. Toxicologic Pathology, 2009, 37, 52-64.	0.9	53
22	Quality assurance and quality control processes: summary of a metabolomics community questionnaire. Metabolomics, 2017, 13, 1.	1.4	53
23	Metabonomics evaluations of age-related changes in the urinary compositions of male Sprague Dawley rats and effects of data normalization methods on statistical and quantitative analysis. BMC Bioinformatics, 2007, 8, S3.	1.2	50
24	An Integrated Study of Acute Effects of Valproic Acid in the Liver Using Metabonomics, Proteomics, and Transcriptomics Platforms. OMICS A Journal of Integrative Biology, 2006, 10, 1-14.	1.0	49
25	Metabolomics as an Extension of Proteomic Analysis: Study of Acute Kidney Injury. Seminars in Nephrology, 2007, 27, 609-620.	0.6	47
26	Progress towards an OECD reporting framework for transcriptomics and metabolomics in regulatory toxicology. Regulatory Toxicology and Pharmacology, 2021, 125, 105020.	1.3	46
27	Reference materials for MS-based untargeted metabolomics and lipidomics: a review by the metabolomics quality assurance and quality control consortium (mQACC). Metabolomics, 2022, 18, 24.	1.4	43
28	Metabolomics evaluation of the effects of green tea extract on acetaminophen-induced hepatotoxicity in mice. Food and Chemical Toxicology, 2013, 62, 707-721.	1.8	42
29	Multiple microRNAs function as self-protective modules in acetaminophen-induced hepatotoxicity in humans. Archives of Toxicology, 2018, 92, 845-858.	1.9	42
30	The Utility of a Rodent Model in Detecting Pediatric Drug-Induced Nephrotoxicity. Toxicological Sciences, 2007, 99, 637-648.	1.4	41
31	Studies of Acetaminophen and Metabolites in Urine and Their Correlations with Toxicity Using Metabolomics. Drug Metabolism Letters, 2009, 3, 130-136.	0.5	41
32	Models of Polychlorinated Dibenzodioxins, Dibenzofurans, and Biphenyls Binding Affinity to the Aryl Hydrocarbon Receptor Developed Using13C NMR Data. Journal of Chemical Information and Computer Sciences, 2001, 41, 1322-1329.	2.8	39
33	Evaluations of the trans-sulfuration pathway in multiple liver toxicity studies. Toxicology and Applied Pharmacology, 2009, 235, 25-32.	1.3	39
34	13C NMR Quantitative Spectrometric Data-Activity Relationship (QSDAR) Models of Steroids Binding the Aromatase Enzyme. Journal of Chemical Information and Computer Sciences, 2001, 41, 1360-1366.	2.8	38
35	Early metabolomics changes in heart and plasma during chronic doxorubicin treatment in B6C3F <sub>1</sub> mice. Journal of Applied Toxicology, 2016, 36, 1486-1495.	1.4	37
36	Identification of metabolite profiles of the catechol-O-methyl transferase inhibitor tolcapone in rat urine using LC/MS-based metabonomics analysis. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2009, 877, 2557-2565.	1.2	36

#	Article	IF	CITATIONS
37	Single valproic acid treatment inhibits glycogen and RNA ribose turnover while disrupting glucose-derived cholesterol synthesis in liver as revealed by the [U-13C6]-d-glucose tracer in mice. Metabolomics, 2009, 5, 336-345.	1.4	34
38	Assessment of usnic acid toxicity in rat primary hepatocytes using 13C isotopomer distribution analysis of lactate, glutamate and glucose. Food and Chemical Toxicology, 2011, 49, 2968-2974.	1.8	32
39	Metabolomic biomarkers: their role in the critical path. Drug Discovery Today: Technologies, 2007, 4, 13-16.	4.0	30
40	Developing 13C NMR quantitative spectrometric data-activity relationship (QSDAR) models of steroid binding globulin. , 2001, 15, 659-669.		29
41	Immune response proteins as predictive biomarkers of doxorubicin-induced cardiotoxicity in breast cancer patients. Experimental Biology and Medicine, 2018, 243, 248-255.	1.1	29
42	Differential gene expression in mouse liver associated with the hepatoprotective effect of clofibrateâ~†. Toxicology and Applied Pharmacology, 2007, 222, 169-179.	1.3	27
43	Evaluating effects of penicillin treatment on the metabolome of rats. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2013, 932, 134-143.	1.2	26
44	Metabolic fate of fructose in human adipocytes: a targeted 13C tracer fate association study. Metabolomics, 2015, 11, 529-544.	1.4	26
45	Proteomic analysis of acetaminophen-induced hepatotoxicity and identification of heme oxygenase 1 as a potential plasma biomarker of liver injury. Proteomics - Clinical Applications, 2017, 11, 1600123.	0.8	26
46	Interest is high in improving quality control for clinical metabolomics: setting the path forward for community harmonization of quality control standards. Metabolomics, 2019, 15, 1.	1.4	26
47	The Role of Metabolic Biomarkers in Drug Toxicity Studies. Toxicology Mechanisms and Methods, 2008, 18, 301-311.	1.3	25
48	Study of valproic acid-induced endogenous and exogenous metabolite alterations using LC–MS-based metabolomics. Bioanalysis, 2010, 2, 207-216.	0.6	25
49	Evaluation of metabolism of azo dyes and their effects on <i>Staphylococcus aureus</i> metabolome. Journal of Industrial Microbiology and Biotechnology, 2017, 44, 1471-1481.	1.4	23
50	BUILDING AN ORGAN-SPECIFIC CARCINOGENIC DATABASE FOR SAR ANALYSES. Journal of Toxicology and Environmental Health - Part A: Current Issues, 2004, 67, 1363-1389.	1,1	22
51	Ageâ€related differences in susceptibility to toxic effects of valproic acid in rats. Journal of Applied Toxicology, 2008, 28, 628-637.	1.4	22
52	Metabolomics data and the biomarker qualification process. Metabolomics, 2012, 8, 2-7.	1.4	22
53	Metabolomics as a tool for personalizing medicine: 2012 update. Personalized Medicine, 2013, 10, 149-161.	0.8	22
54	Emerging technologies and their impact on regulatory science. Experimental Biology and Medicine, 2022, 247, 1-75.	1.1	22

4

#	Article	IF	CITATIONS
55	Dose-response analysis of epigenetic, metabolic, and apical endpoints after short-term exposure to experimental hepatotoxicants. Food and Chemical Toxicology, 2017, 109, 690-702.	1.8	21
56	Comparative Structural Connectivity Spectra Analysis (CoSCoSA) Models of Steroid Binding to the Corticosteroid Binding Globulin. Journal of Chemical Information and Computer Sciences, 2002, 42, 1123-1131.	2.8	20
57	NMR-based metabonomic evaluation of livers from rats chronically treated with tamoxifen, mestranol, and phenobarbital. Metabolomics, 2005, 1, 87-94.	1.4	20
58	Metabolomic analysis of urine from rats chronically dosed with acrylamide using NMR and LC/MS. Metabolomics, 2010, 6, 550-563.	1.4	20
59	The Liver Toxicity Biomarker Study Phase I: Markers for the Effects of Tolcapone or Entacapone. Toxicologic Pathology, 2012, 40, 951-964.	0.9	20
60	Computational identification of a phospholipidosis toxicophore using 13 C and 15 N NMR-distance based fingerprints. Bioorganic and Medicinal Chemistry, 2014, 22, 6706-6714.	1.4	20
61	Aptamer-Based Proteomics Identifies Mortality-Associated Serum Biomarkers in Dialysis-Dependent AKI Patients. Kidney International Reports, 2018, 3, 1202-1213.	0.4	20
62	An Integrated Analysis of Metabolites, Peptides, and Inflammation Biomarkers for Assessment of Preanalytical Variability of Human Plasma. Journal of Proteome Research, 2019, 18, 2411-2421.	1.8	20
63	Hepatic Transcript Profiles of Cytochrome P450 Genes Predict Sex Differences in Drug Metabolism. Drug Metabolism and Disposition, 2020, 48, 447-458.	1.7	20
64	Metabolomics evaluation of hydroxyproline as a potential marker of melamine and cyanuric acid nephrotoxicity in male and female Fischer F344 rats. Food and Chemical Toxicology, 2012, 50, 3978-3983.	1.8	19
65	Microbiota of MR1 deficient mice confer resistance against Clostridium difficile infection. PLoS ONE, 2019, 14, e0223025.	1.1	19
66	Evaluation of NMR spectral data of urine in conjunction with measured clinical chemistry and histopathology parameters to assess the effects of liver and kidney toxicants. Metabolomics, 2007, 3, 87-100.	1.4	18
67	Fructose Alters Intermediary Metabolism of Glucose in Human Adipocytes and Diverts Glucose to Serine Oxidation in the One–Carbon Cycle Energy Producing Pathway. Metabolites, 2015, 5, 364-385.	1.3	17
68	Stability of the Human Plasma Proteome to Pre-analytical Variability as Assessed by an Aptamer-Based Approach. Journal of Proteome Research, 2019, 18, 3661-3670.	1.8	17
69	Discovery of Novel Proteomic Biomarkers for the Prediction of Kidney Recovery from Dialysis-Dependent AKI Patients. Kidney360, 2021, 2, 1716-1727.	0.9	16
70	Producing13C NMR, Infrared Absorption, and Electron Ionization Mass Spectrometric Data Models of the Monodechlorination of Chlorobenzenes, Chlorophenols, and Chloroanilines. Journal of Chemical Information and Computer Sciences, 2000, 40, 1449-1455.	2.8	15
71	Partial least square and kâ€nearest neighbor algorithms for improved 3D quantitative spectral data–activity relationship consensus modeling of acute toxicity. Environmental Toxicology and Chemistry, 2014, 33, 1271-1282.	2.2	15
72	Combining NMR spectral and structural data to form models of polychlorinated dibenzodioxins, dibenzofurans, and biphenyls binding to the AhR. Journal of Computer-Aided Molecular Design, 2002, 16, 727-740.	1.3	14

#	Article	IF	CITATIONS
73	Metabolomics Analysis of Urine Samples from Children after Acetaminophen Overdose. Metabolites, 2017, 7, 46.	1.3	14
74	The New Data Quality Task Group (DQTG): ensuring high quality data today and in the future. Metabolomics, 2014, 10, 539-540.	1.4	13
75	Circulating mitochondrial biomarkers for drug-induced liver injury. Biomarkers in Medicine, 2015, 9, 1215-1223.	0.6	13
76	Metabolomics Test Materials for Quality Control: A Study of a Urine Materials Suite. Metabolites, 2019, 9, 270.	1.3	13
77	Metabolomics as a Truly Translational Tool for Precision Medicine. International Journal of Toxicology, 2021, 40, 413-426.	0.6	13
78	Improving proton MR spectroscopy of brain tissue for noninvasive diagnostics. Journal of Magnetic Resonance Imaging, 2010, 32, 818-829.	1.9	12
79	Metabolomics evaluation of the impact of smokeless tobacco exposure on the oral bacterium Capnocytophaga sputigena. Toxicology in Vitro, 2016, 36, 133-141.	1.1	12
80	Testicular function in cultured postnatal mouse testis fragments is similar to that of animals during the first wave of spermatogenesis. Birth Defects Research, 2019, 111, 270-280.	0.8	12
81	Discriminant Function Analyses of Liver-Specific Carcinogens. Journal of Chemical Information and Computer Sciences, 2004, 44, 1107-1110.	2.8	11
82	Complementary PLS and KNN algorithms for improved 3D-QSDAR consensus modeling of AhR binding. Journal of Cheminformatics, 2013, 5, 47.	2.8	11
83	3D-SDAR modeling of hERG potassium channel affinity: A case study in model design and toxicophore identification. Journal of Molecular Graphics and Modelling, 2017, 72, 246-255.	1.3	10
84	Comparative structural connectivity spectra analysis (CoSCoSA) models of steroids binding to the aromatase enzyme. Journal of Molecular Recognition, 2002, 15, 154-162.	1.1	9
85	The use of carbon thirteen nuclear magnetic resonance spectra to predict dioxin and furan binding affinities to the aryl hydrocarbon receptor. Environmental Toxicology and Chemistry, 2003, 22, 501-509.	2.2	9
86	Computational modeling of biologically active molecules using NMR spectra. Drug Discovery Today, 2006, 11, 429-435.	3.2	9
87	Discovery of early urinary biomarkers in preclinical study of gentamicin-induced kidney injury and recovery in rats. Metabolomics, 2012, 8, 1181-1193.	1.4	9
88	Models of Steroid Binding Based on the Minimum Deviation of Structurally Assigned 13C NMR Spectra Analysis (MiDSASA). Journal of Chemical Information and Computer Sciences, 2004, 44, 1489-1496.	2.8	8
89	Metabolomics: a tool for personalizingÂmedicine?. Personalized Medicine, 2008, 5, 495-504.	0.8	8
90	Targeted metabolomic profiling indicates structure-based perturbations in serum phospholipids in children with acetaminophen overdose. Toxicology Reports, 2016, 3, 747-755.	1.6	8

#	Article	IF	CITATIONS
91	Serum metabolite profiles predict outcomes in critically ill patients receiving renal replacement therapy. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2021, 1187, 123024.	1.2	8
92	Pharmacometabolomics in drug safety and drug-exposome interactions. Metabolomics, 2016, 12, 1.	1.4	7
93	Comparison of Effects of Diet on Mammary Cancer: Efficacy of Various Preventive Agents and Metabolomic Changes of Different Diets and Agents. Cancer Prevention Research, 2018, 11, 831-840.	0.7	7
94	Bile Acid Profile and its Changes in Response to Cefoperazone Treatment in MR1 Deficient Mice. Metabolites, 2020, 10, 127.	1.3	7
95	<sup>13</sup> C NMR–Distance Matrix Descriptors: Optimal Abstract 3D Space Granularity for Predicting Estrogen Binding. Journal of Chemical Information and Modeling, 2012, 52, 1854-1864.	2.5	6
96	Computational identification of structural factors affecting the mutagenic potential of aromatic amines: study design and experimental validation. Archives of Toxicology, 2018, 92, 2369-2384.	1.9	6
97	Pharmacometabolomic Pathway Response of Effective Anticancer Agents on Different Diets in Rats with Induced Mammary Tumors. Metabolites, 2019, 9, 149.	1.3	6
98	Comprehensive analysis of alterations in lipid and bile acid metabolism by carbon tetrachloride using integrated transcriptomics and metabolomics. Metabolomics, 2014, 10, 1293-1304.	1.4	5
99	Rigorous 3-dimensional spectral data activity relationship approach modeling strategy for ToxCast estrogen receptor data classification, validation, and feature extraction. Environmental Toxicology and Chemistry, 2017, 36, 823-830.	2.2	4
100	Distinct lipid signatures are identified in the plasma of rats with chronic inflammation induced by estradiol benzoate and sex hormones. Metabolomics, 2020, 16, 95.	1.4	4
101	Quantitative structure–toxicity relationships in translational toxicology. Current Opinion in Toxicology, 2020, 23-24, 46-49.	2.6	4
102	Metabolomicsâ€based pathway changes in testis fragments treated with ethinylestradiol in vitro. Birth Defects Research, 2019, 111, 1643-1654.	0.8	2
103	Determination of structural factors affecting binding to mu, kappa and delta opioid receptors. Archives of Toxicology, 2020, 94, 1215-1227.	1.9	2
104	Targeted 13C-Labeled Tracer Fate Associations for Drug Efficacy Testing in Cancer. , 2015, , 349-372.		2
105	Evaluating Cefoperazone-Induced Gut Metabolic Functional Changes in MR1-Deficient Mice. Metabolites, 2022, 12, 380.	1.3	1
106	Identification of structural factors that affect binding to cannabinoid receptor type 1. Journal of Molecular Structure, 2022, 1249, 131589.	1.8	0
107	An Aptamerâ€Based Approach to Assess the Human Plasma Proteome for Preâ€Analytical Variability. FASEB Journal, 2018, 32, 802.5.	0.2	0