Hector C Keun

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/1129235/hector-c-keun-publications-by-year.pdf

Version: 2024-04-09

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

128 8,537 49 91 h-index g-index citations papers 5.56 136 9,795 7.1 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
128	Indisulam targets RNA splicing and metabolism to serve as a therapeutic strategy for high-risk neuroblastoma <i>Nature Communications</i> , 2022 , 13, 1380	17.4	1
127	Metabonomics study of the effects of single copy mutant KRAS in the presence or absence of WT allele using human HCT116 isogenic cell lines. <i>Metabolomics</i> , 2021 , 17, 104	4.7	1
126	Advancing tools for human early lifecourse exposome research and translation (ATHLETE): Project overview <i>Environmental Epidemiology</i> , 2021 , 5, e166	0.2	2
125	Systems level profiling of chemotherapy-induced stress resolution in cancer cells reveals druggable trade-offs. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	4
124	Metabolic signatures of greater body size and their associations with risk of colorectal and endometrial cancers in the European Prospective Investigation into Cancer and Nutrition. <i>BMC Medicine</i> , 2021 , 19, 101	11.4	6
123	Phase I expansion study of the first-in-class monocarboxylate transporter 1 (MCT1) inhibitor AZD3965 in patients with diffuse large B-cell lymphoma (DLBCL) and Burkitt lymphoma (BL) <i>Journal of Clinical Oncology</i> , 2021 , 39, 3115-3115	2.2	4
122	Enhanced triacylglycerol catabolism by carboxylesterase 1 promotes aggressive colorectal carcinoma. <i>Journal of Clinical Investigation</i> , 2021 , 131,	15.9	5
121	Variability of multi-omics profiles in a population-based child cohort. <i>BMC Medicine</i> , 2021 , 19, 166	11.4	7
120	Urinary metabolite quantitative trait loci in children and their interaction with dietary factors. <i>Human Molecular Genetics</i> , 2021 , 29, 3830-3844	5.6	1
119	Prospective analysis of circulating metabolites and endometrial cancer risk. <i>Gynecologic Oncology</i> , 2021 , 162, 475-481	4.9	4
118	The early-life exposome and epigenetic age acceleration in children. <i>Environment International</i> , 2021 , 155, 106683	12.9	5
117	Lactic acidosis induces resistance to the pan-Akt inhibitor uprosertib in colon cancer cells. <i>British Journal of Cancer</i> , 2020 , 122, 1298-1308	8.7	9
116	Tracing Nutrient Flux Following Monocarboxylate Transporter-1 Inhibition with AZD3965. <i>Cancers</i> , 2020 , 12,	6.6	2
115	A case of malignant hyperlactaemic acidosis appearing upon treatment with the mono-carboxylase transporter 1 inhibitor AZD3965. <i>British Journal of Cancer</i> , 2020 , 122, 1141-1145	8.7	5
114	Metabolic characterization of colorectal cancer cells harbouring different KRAS mutations in codon 12, 13, 61 and 146 using human SW48 isogenic cell lines. <i>Metabolomics</i> , 2020 , 16, 51	4.7	7
113	Prenatal Exposure to Perfluoroalkyl Substances Associated With Increased Susceptibility to Liver Injury in Children. <i>Hepatology</i> , 2020 , 72, 1758-1770	11.2	27
112	EHydroxybutyrate Oxidation Promotes the Accumulation of Immunometabolites in Activated Microglia Cells. <i>Metabolites</i> , 2020 , 10,	5.6	2

(2018-2020)

111	In utero and childhood exposure to tobacco smoke and multi-layer molecular signatures in children. <i>BMC Medicine</i> , 2020 , 18, 243	11.4	6
110	RNA-binding motif protein 39 (RBM39): An emerging cancer target. <i>British Journal of Pharmacology</i> , 2020 ,	8.6	2
109	Opportunities at the Interface of Network Science and Metabolic Modeling. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020 , 8, 591049	5.8	5
108	The coordinated action of VCP/p97 and GCN2 regulates cancer cell metabolism and proteostasis during nutrient limitation. <i>Oncogene</i> , 2019 , 38, 3216-3231	9.2	23
107	Astrocyte adenosine deaminase loss increases motor neuron toxicity in amyotrophic lateral sclerosis. <i>Brain</i> , 2019 , 142, 586-605	11.2	44
106	Intracellular Elicits the Production of Host Very Long-Chain Saturated Fatty Acids with Antimicrobial Activity. <i>Metabolites</i> , 2019 , 9,	5.6	8
105	The impact of p53 on aristolochic acid I-induced nephrotoxicity and DNA damage in vivo and in vitro. <i>Archives of Toxicology</i> , 2019 , 93, 3345-3366	5.8	11
104	The 14q32 maternally imprinted locus is a major source of longitudinally stable circulating microRNAs as measured by small RNA sequencing. <i>Scientific Reports</i> , 2019 , 9, 15787	4.9	5
103	Predictive modelling using pathway scores: robustness and significance of pathway collections. <i>BMC Bioinformatics</i> , 2019 , 20, 543	3.6	8
102	Integrated Systems Level Examination of Proteasome Inhibitor Stress Recovery in Myeloma Cells Reveals Druggable Vulnerabilities Linked to Multiple Metabolic Processes. <i>Blood</i> , 2019 , 134, 1818-1818	3 2.2	
101	Spinal Cord Metabolic Signatures in Models of Fast- and Slow-Progressing SOD1 Amyotrophic Lateral Sclerosis. <i>Frontiers in Neuroscience</i> , 2019 , 13, 1276	5.1	9
100	The Impact of p53 on Aristolochic Acid I-Induced Gene Expression In Vivo. <i>International Journal of Molecular Sciences</i> , 2019 , 20,	6.3	3
99	Acetaminophen cytotoxicity in HepG2 cells is associated with a decoupling of glycolysis from the TCA cycle, loss of NADPH production, and suppression of anabolism. <i>Archives of Toxicology</i> , 2019 , 93, 341-353	5.8	14
98	Pre-diagnostic blood immune markers, incidence and progression of B-cell lymphoma and multiple myeloma: Univariate and functionally informed multivariate analyses. <i>International Journal of Cancer</i> , 2018 , 143, 1335-1347	7.5	10
97	A model-based assay design to reproduce in vivo patterns of acute drug-induced toxicity. <i>Archives of Toxicology</i> , 2018 , 92, 553-555	5.8	17
96	Determinants of the urinary and serum metabolome in children from six European populations. <i>BMC Medicine</i> , 2018 , 16, 202	11.4	56
95	Persistence of Epigenomic Effects After Recovery From Repeated Treatment With Two Nephrocarcinogens. <i>Frontiers in Genetics</i> , 2018 , 9, 558	4.5	4
94	Urine Metabolic Signatures of Multiple Environmental Pollutants in Pregnant Women: An Exposome Approach. <i>Environmental Science & Exposome Approach</i> . 2018, 52, 13469-13480	10.3	32

93	In-utero and childhood chemical exposome in six European mother-child cohorts. <i>Environment International</i> , 2018 , 121, 751-763	12.9	79
92	Intracellular Staphylococcus aureus Modulates Host Central Carbon Metabolism To Activate Autophagy. <i>MSphere</i> , 2018 , 3,	5	37
91	Human Early Life Exposome (HELIX) study: a European population-based exposome cohort. <i>BMJ Open</i> , 2018 , 8, e021311	3	88
90	Omics-based responses induced by bosentan in human hepatoma HepaRG cell cultures. <i>Archives of Toxicology</i> , 2018 , 92, 1939-1952	5.8	21
89	Blood-based omic profiling supports female susceptibility to tobacco smoke-induced cardiovascular diseases. <i>Scientific Reports</i> , 2017 , 7, 42870	4.9	19
88	CYP3A7*1C allele is associated with reduced levels of 2-hydroxylation pathway oestrogen metabolites. <i>British Journal of Cancer</i> , 2017 , 116, 382-388	8.7	8
87	Inhibition of monocarboxyate transporter 1 by AZD3965 as a novel therapeutic approach for diffuse large B-cell lymphoma and Burkitt lymphoma. <i>Haematologica</i> , 2017 , 102, 1247-1257	6.6	58
86	Metabolomic characterisation of the effects of oncogenic PIK3CA transformation in a breast epithelial cell line. <i>Scientific Reports</i> , 2017 , 7, 46079	4.9	14
85	Metabolite signatures of doxorubicin induced toxicity in human induced pluripotent stem cell-derived cardiomyocytes. <i>Amino Acids</i> , 2017 , 49, 1955-1963	3.5	14
84	Assessment of metabolic phenotypic variability in children's urine using H NMR spectroscopy. <i>Scientific Reports</i> , 2017 , 7, 46082	4.9	23
83	Interlaboratory Reproducibility of a Targeted Metabolomics Platform for Analysis of Human Serum and Plasma. <i>Analytical Chemistry</i> , 2017 , 89, 656-665	7.8	131
82	Altered Metabolic Profiles Associate with Toxicity in SOD1 Astrocyte-Neuron Co-Cultures. <i>Scientific Reports</i> , 2017 , 7, 50	4.9	10
81	Lipid degradation promotes prostate cancer cell survival. <i>Oncotarget</i> , 2017 , 8, 38264-38275	3.3	41
80	T cell inhibitory mechanisms in a model of aggressive Non-Hodgkin's Lymphoma. <i>OncoImmunology</i> , 2017 , 7, e1365997	7.2	1
79	Metabolomic Analysis Reveals Increased Aerobic Glycolysis and Amino Acid Deficit in a Cellular Model of Amyotrophic Lateral Sclerosis. <i>Molecular Neurobiology</i> , 2016 , 53, 2222-40	6.2	44
78	p53 Loss in MYC-Driven Neuroblastoma Leads to Metabolic Adaptations Supporting Radioresistance. <i>Cancer Research</i> , 2016 , 76, 3025-35	10.1	25
77	Suppression of MTHFD2 in MCF-7 Breast Cancer Cells Increases Glycolysis, Dependency on Exogenous Glycine, and Sensitivity to Folate Depletion. <i>Journal of Proteome Research</i> , 2016 , 15, 2618-25	5.6	23
76	Identification of genomic biomarkers for anthracycline-induced cardiotoxicity in human iPSC-derived cardiomyocytes: an in vitro repeated exposure toxicity approach for safety assessment. <i>Archives of Toxicology</i> , 2016 , 90, 2763-2777	5.8	62

(2013-2016)

75	A Systems Oncology Approach Identifies NT5E as a Key Metabolic Regulator in Tumor Cells and Modulator of Platinum Sensitivity. <i>Journal of Proteome Research</i> , 2016 , 15, 280-90	5.6	17
74	Metabonomic analysis of ovarian tumour cyst fluid by proton nuclear magnetic resonance spectroscopy. <i>Oncotarget</i> , 2016 , 7, 7216-26	3.3	25
73	Over-representation of correlation analysis (ORCA): a method for identifying associations between variable sets. <i>Bioinformatics</i> , 2015 , 31, 102-8	7.2	2
72	Influence of glutathione-S-transferase (GST) inhibition on lung epithelial cell injury: role of oxidative stress and metabolism. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2015 , 308, L1274-85	5.8	17
71	convISA: A simple, convoluted method for isotopomer spectral analysis of fatty acids and cholesterol. <i>Metabolic Engineering</i> , 2015 , 32, 125-132	9.7	14
70	Metabolic profiling in human exposome studies. <i>Mutagenesis</i> , 2015 , 30, 755-62	2.8	33
69	Plasma metabolomic profiles of breast cancer patients after short-term limonene intervention. <i>Cancer Prevention Research</i> , 2015 , 8, 86-93	3.2	27
68	Diurnal rhythms in the human urine metabolome during sleep and total sleep deprivation. <i>Scientific Reports</i> , 2015 , 5, 14843	4.9	88
67	diXa: a data infrastructure for chemical safety assessment. <i>Bioinformatics</i> , 2015 , 31, 1505-7	7.2	27
66	Alterations of choline phospholipid metabolism in endometrial cancer are caused by choline kinase alpha overexpression and a hyperactivated deacylation pathway. <i>Cancer Research</i> , 2014 , 74, 6867-77	10.1	61
65	Metabolomic characterization of nipple aspirate fluid by (1)H NMR spectroscopy and GC-MS. <i>Journal of Proteome Research</i> , 2014 , 13, 883-9	5.6	15
64	Metabolomic studies of patient material by high-resolution magic angle spinning nuclear magnetic resonance spectroscopy. <i>Methods in Enzymology</i> , 2014 , 543, 297-313	1.7	5
63	Metabonomic analysis of water extracts from different angelica roots by IH-nuclear magnetic resonance spectroscopy. <i>Molecules</i> , 2014 , 19, 3460-70	4.8	12
62	The human early-life exposome (HELIX): project rationale and design. <i>Environmental Health Perspectives</i> , 2014 , 122, 535-44	8.4	219
61	Serum metabolomic pertubations among workers exposed to 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD). <i>Environmental and Molecular Mutagenesis</i> , 2013 , 54, 558-65	3.2	20
60	Evaluation of 1H NMR metabolic profiling using biofluid mixture design. <i>Analytical Chemistry</i> , 2013 , 85, 6674-81	7.8	4
59	1H HR-MAS NMR spectroscopy of tumor-induced local metabolic "field-effects" enables colorectal cancer staging and prognostication. <i>Journal of Proteome Research</i> , 2013 , 12, 959-68	5.6	92
58	Performance in omics analyses of blood samples in long-term storage: opportunities for the exploitation of existing biobanks in environmental health research. <i>Environmental Health Perspectives</i> , 2013 , 121, 480-7	8.4	111

57	Metabolomics in toxicology and preclinical research. <i>ALTEX: Alternatives To Animal Experimentation</i> , 2013 , 30, 209-25	4.3	135
56	Metabolic characterization of Leishmania major infection in activated and nonactivated macrophages. <i>Journal of Proteome Research</i> , 2012 , 11, 4211-22	5.6	24
55	Intra- and interlaboratory reproducibility of ultra performance liquid chromatography-time-of-flight mass spectrometry for urinary metabolic profiling. <i>Analytical Chemistry</i> , 2012 , 84, 2424-32	7.8	36
54	Metabolic profiling detects early effects of environmental and lifestyle exposure to cadmium in a human population. <i>BMC Medicine</i> , 2012 , 10, 61	11.4	98
53	Chiral metabonomics: 1H NMR-based enantiospecific differentiation of metabolites in human urine via direct cosolvation with Eyclodextrin. <i>Analytical Chemistry</i> , 2012 , 84, 2868-74	7.8	31
52	Biofluid metabonomics using (1)H NMR spectroscopy: the road to biomarker discovery in gastroenterology and hepatology. <i>Expert Review of Gastroenterology and Hepatology</i> , 2012 , 6, 239-51	4.2	22
51	Choline-releasing glycerophosphodiesterase EDI3 drives tumor cell migration and metastasis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 8155-60	11.5	69
50	Reply to Moestue et al.: Untangling the contribution of choline metabolism to the metastatic process: Fig. 1 <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, E2507-E2507	11.5	1
49	Metabolic signatures of malignant progression in prostate epithelial cells. <i>International Journal of Biochemistry and Cell Biology</i> , 2011 , 43, 1002-9	5.6	40
48	Nuclear magnetic resonance (NMR)-based metabolomics. <i>Methods in Molecular Biology</i> , 2011 , 708, 321	-3 1. 4	30
47	Evaluation of urinary ribonucleoside profiling for clinical biomarker discovery using constant neutral loss scanning liquid chromatography/tandem mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2011 , 25, 2071-82	2.2	13
46	Meeting-in-the-middle using metabolic profiling - a strategy for the identification of intermediate biomarkers in cohort studies. <i>Biomarkers</i> , 2011 , 16, 83-8	2.6	88
45	Pharmacometabonomic profiling as a predictor of toxicity in patients with inoperable colorectal cancer treated with capecitabine. <i>Clinical Cancer Research</i> , 2011 , 17, 3019-28	12.9	90
44	Consensus-phenotype integration of transcriptomic and metabolomic data implies a role for metabolism in the chemosensitivity of tumour cells. <i>PLoS Computational Biology</i> , 2011 , 7, e1001113	5	65
43	High-resolution magic-angle-spinning NMR spectroscopy for metabolic profiling of intact tissues. <i>Nature Protocols</i> , 2010 , 5, 1019-32	18.8	309
42	Metabolic profiling detects field effects in nondysplastic tissue from esophageal cancer patients. <i>Cancer Research</i> , 2010 , 70, 9129-36	10.1	38
41	1H NMR spectroscopy-based interventional metabolic phenotyping: a cohort study of rheumatoid arthritis patients. <i>Journal of Proteome Research</i> , 2010 , 9, 4545-53	5.6	75
40	Effect of the histone deacetylase inhibitor trichostatin a on the metabolome of cultured primary hepatocytes. <i>Journal of Proteome Research</i> , 2010 , 9, 413-9	5.6	12

(2008-2010)

39	Ultra performance liquid chromatography-mass spectrometry profiling of bile acid metabolites in biofluids: application to experimental toxicology studies. <i>Analytical Chemistry</i> , 2010 , 82, 5282-9	7.8	79
38	NMR-based metabolic profiling identifies biomarkers of liver regeneration following partial hepatectomy in the rat. <i>Journal of Proteome Research</i> , 2010 , 9, 59-69	5.6	68
37	Plasma metabolic profiling reveals age-dependency of systemic effects of green tea polyphenols in mice with and without prostate cancer. <i>Molecular BioSystems</i> , 2010 , 6, 1911-6		3
36	Metabolic Profiling for Biomarker Discovery 2010 , 47-74		
35	Circulating sphingosine-1-phosphate inversely correlates with chemotherapy-induced weight gain during early breast cancer. <i>Breast Cancer Research and Treatment</i> , 2010 , 124, 543-9	4.4	6
34	Genetic algorithms for simultaneous variable and sample selection in metabonomics. <i>Bioinformatics</i> , 2009 , 25, 112-8	7.2	50
33	Serum molecular signatures of weight change during early breast cancer chemotherapy. <i>Clinical Cancer Research</i> , 2009 , 15, 6716-23	12.9	50
32	Citrate transport and metabolism in mammalian cells: prostate epithelial cells and prostate cancer. <i>BioEssays</i> , 2009 , 31, 10-20	4.1	98
31	Cluster analysis statistical spectroscopy using nuclear magnetic resonance generated metabolic data sets from perturbed biological systems. <i>Analytical Chemistry</i> , 2009 , 81, 6581-9	7.8	30
30	Pivotal role for two electron reduction in 2,3-dimethoxy-1,4-naphthoquinone and 2-methyl-1,4-naphthoquinone metabolism and kinetics in vivo that prevents liver redox stress. <i>Chemical Research in Toxicology</i> , 2009 , 22, 717-25	4	17
29	Mechanistic aspects and novel biomarkers of responder and non-responder phenotypes in galactosamine-induced hepatitis. <i>Journal of Proteome Research</i> , 2009 , 8, 5175-87	5.6	35
28	Detection of metabolic alterations in non-tumor gastrointestinal tissue of the Apc(Min/+) mouse by (1)H MAS NMR spectroscopy. <i>Journal of Proteome Research</i> , 2009 , 8, 1423-30	5.6	31
27	Metabolic profiling of human colorectal cancer using high-resolution magic angle spinning nuclear magnetic resonance (HR-MAS NMR) spectroscopy and gas chromatography mass spectrometry (GC/MS). <i>Journal of Proteome Research</i> , 2009 , 8, 352-61	5.6	377
26	Temporal metabonomic modeling of l-arginine-induced exocrine pancreatitis. <i>Journal of Proteome Research</i> , 2008 , 7, 4435-45	5.6	49
25	The carcinoGENOMICS project: critical selection of model compounds for the development of omics-based in vitro carcinogenicity screening assays. <i>Mutation Research - Reviews in Mutation Research</i> , 2008 , 659, 202-10	7	50
24	Heteronuclear 19F-1H statistical total correlation spectroscopy as a tool in drug metabolism: study of flucloxacillin biotransformation. <i>Analytical Chemistry</i> , 2008 , 80, 1073-9	7.8	50
23	Metabolic profiling of transgenic adenocarcinoma of mouse prostate (TRAMP) tissue by 1H-NMR analysis: evidence for unusual phospholipid metabolism. <i>Prostate</i> , 2008 , 68, 1035-47	4.2	31
22	Robust algorithms for automated chemical shift calibration of 1D 1H NMR spectra of blood serum. <i>Analytical Chemistry</i> , 2008 , 80, 7158-62	7.8	52

21	Direct quantitative trait locus mapping of mammalian metabolic phenotypes in diabetic and normoglycemic rat models. <i>Nature Genetics</i> , 2007 , 39, 666-72	36.3	132
20	Metabolic profiling, metabolomic and metabonomic procedures for NMR spectroscopy of urine, plasma, serum and tissue extracts. <i>Nature Protocols</i> , 2007 , 2, 2692-703	18.8	1536
19	Standard reporting requirements for biological samples in metabolomics experiments: mammalian/in vivo experiments. <i>Metabolomics</i> , 2007 , 3, 179-188	4.7	63
18	Application of metabonomics in drug development. <i>Pharmacogenomics</i> , 2007 , 8, 731-41	2.6	53
17	Prediction and classification of drug toxicity using probabilistic modeling of temporal metabolic data: the consortium on metabonomic toxicology screening approach. <i>Journal of Proteome Research</i> , 2007 , 6, 4407-22	5.6	146
16	Metabolic profile biomarkers of metal contamination in a sentinel terrestrial species are applicable across multiple sites. <i>Environmental Science & Environmental Science & E</i>	10.3	93
15	Metabonomic modeling of drug toxicity 2006 , 109, 92-106		128
14	Impact of analytical bias in metabonomic studies of human blood serum and plasma. <i>Analytical Chemistry</i> , 2006 , 78, 4307-18	7.8	200
13	Metabolomic evaluation of rat liver and testis to characterize the toxicity of triazole fungicides. <i>Metabolomics</i> , 2006 , 2, 63-73	4.7	34
12	Summary recommendations for standardization and reporting of metabolic analyses. <i>Nature Biotechnology</i> , 2005 , 23, 833-8	44.5	233
11	Comparative metabonomics of differential hydrazine toxicity in the rat and mouse. <i>Toxicology and Applied Pharmacology</i> , 2005 , 204, 135-51	4.6	107
10	The Consortium for Metabonomic Toxicology (COMET): aims, activities and achievements. <i>Pharmacogenomics</i> , 2005 , 6, 691-9	2.6	255
9	Geometric trajectory analysis of metabolic responses to toxicity can define treatment specific profiles. <i>Chemical Research in Toxicology</i> , 2004 , 17, 579-87	4	131
8	Spectral editing and pattern recognition methods applied to high-resolution magic-angle spinning 1H nuclear magnetic resonance spectroscopy of liver tissues. <i>Analytical Biochemistry</i> , 2003 , 323, 26-32	3.1	134
7	Contemporary issues in toxicology the role of metabonomics in toxicology and its evaluation by the COMET project. <i>Toxicology and Applied Pharmacology</i> , 2003 , 187, 137-46	4.6	342
6	NMR-based metabonomic toxicity classification: hierarchical cluster analysis and k-nearest-neighbour approaches. <i>Analytica Chimica Acta</i> , 2003 , 490, 3-15	6.6	127
5	Improved analysis of multivariate data by variable stability scaling: application to NMR-based metabolic profiling. <i>Analytica Chimica Acta</i> , 2003 , 490, 265-276	6.6	149
4	Toxicity classification from metabonomic data using a density superposition approach: © LOUDS I <i>Analytica Chimica Acta</i> , 2003 , 490, 109-122	6.6	69

LIST OF PUBLICATIONS

3	Analytical reproducibility in (1)H NMR-based metabonomic urinalysis. <i>Chemical Research in Toxicology</i> , 2002 , 15, 1380-6	4	239
2	Cryogenic probe 13C NMR spectroscopy of urine for metabonomic studies. <i>Analytical Chemistry</i> , 2002 , 74, 4588-93	7.8	174
1	Metabolic profiling of rodent biological fluids via 1H NMR spectroscopy using a 1 mm microlitre probe. <i>Analyst, The</i> , 2002 , 127, 582-4	5	44