Helen G Gika

List of Publications by Year in descending order

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		87888	64796
133	6,735	38	79
papers	citations	h-index	g-index
139	139	139	8047
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Global metabolic profiling procedures for urine using UPLC–MS. Nature Protocols, 2010, 5, 1005-1018.	12.0	867
2	Within-Day Reproducibility of an HPLCâ^'MS-Based Method for Metabonomic Analysis:  Application to Human Urine. Journal of Proteome Research, 2007, 6, 3291-3303.	3.7	459
3	Liquid chromatography–mass spectrometry based global metabolite profiling: A review. Analytica Chimica Acta, 2012, 711, 7-16.	5.4	452
4	Current practice of liquid chromatography–mass spectrometry in metabolomics and metabonomics. Journal of Pharmaceutical and Biomedical Analysis, 2014, 87, 12-25.	2.8	348
5	LC-MS-based methodology for global metabolite profiling in metabonomics/metabolomics. TrAC - Trends in Analytical Chemistry, 2008, 27, 251-260.	11.4	306
6	Evaluation of the repeatability of ultra-performance liquid chromatography–TOF-MS for global metabolic profiling of human urine samples. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2008, 871, 299-305.	2.3	215
7	Site and Strain-Specific Variation in Gut Microbiota Profiles and Metabolism in Experimental Mice. PLoS ONE, 2010, 5, e8584.	2.5	186
8	Hydrophilic interaction chromatography coupled to MS for metabonomic/metabolomic studies. Journal of Separation Science, 2010, 33, 716-727.	2.5	180
9	Liquid chromatography and ultra-performance liquid chromatography–mass spectrometry fingerprinting of human urine. Journal of Chromatography A, 2008, 1189, 314-322.	3.7	178
10	Mass spectrometryâ€based holistic analytical approaches for metabolite profiling in systems biology studies. Mass Spectrometry Reviews, 2011, 30, 884-906.	5.4	171
11	UPLC-MS-Based Analysis of Human Plasma for Metabonomics Using Solvent Precipitation or Solid Phase Extraction. Journal of Proteome Research, 2009, 8, 2114-2121.	3.7	159
12	Hydrophilic interaction and reversedâ€phase ultraâ€performance liquid chromatography TOFâ€MS for metabonomic analysis of Zucker rat urine. Journal of Separation Science, 2008, 31, 1598-1608.	2.5	121
13	¹ H NMR-Based Metabonomic Investigation of the Effect of Two Different Exercise Sessions on the Metabolic Fingerprint of Human Urine. Journal of Proteome Research, 2010, 9, 6405-6416.	3.7	106
14	Untargeted LC/MS-based metabolic phenotyping (metabonomics/metabolomics): The state of the art. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2019, 1117, 136-147.	2.3	106
15	Quantitative profiling of polar primary metabolites using hydrophilic interaction ultrahigh performance liquid chromatography–tandem mass spectrometry. Journal of Chromatography A, 2012, 1259, 121-127.	3.7	105
16	An overview of fecal sample preparation for global metabolic profiling. Journal of Pharmaceutical and Biomedical Analysis, 2015, 113, 137-150.	2.8	104
17	Metabolite profiling on apple volatile content based on solid phase microextraction and gas-chromatography time of flight mass spectrometry. Journal of Chromatography A, 2011, 1218, 4517-4524.	3.7	100
18	LC–MS-based holistic metabolic profiling. Problems, limitations, advantages, and future perspectives. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2014, 966, 1-6.	2.3	88

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19	Sample preparation prior to the LC–MS-based metabolomics/metabonomics of blood-derived samples. Bioanalysis, 2011, 3, 1647-1661.	1.5	82
20	Protocol for quality control in metabolic profiling of biological fluids by U(H)PLC-MS. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2016, 1008, 15-25.	2.3	78
21	Development and validation of a HILICâ€MS/MS multitargeted method for metabolomics applications. Electrophoresis, 2015, 36, 2215-2225.	2.4	77
22	LC-MS based global metabolite profiling of grapes: solvent extraction protocol optimisation. Metabolomics, 2012, 8, 175-185.	3.0	72
23	Hyphenated MS-based targeted approaches in metabolomics. Analyst, The, 2017, 142, 3079-3100.	3.5	72
24	A QC approach to the determination of day-to-day reproducibility and robustness of LC–MS methods for global metabolite profiling in metabonomics/metabolomics. Bioanalysis, 2012, 4, 2239-2247.	1.5	71
25	High temperature-ultra performance liquid chromatography–mass spectrometry for the metabonomic analysis of Zucker rat urine. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2008, 871, 279-287.	2.3	66
26	Investigation of the derivatization conditions for GC–MS metabolomics of biological samples. Bioanalysis, 2017, 9, 53-65.	1.5	65
27	Application of Ultra Performance Liquid Chromatographyâ^'Mass Spectrometry to Profiling Rat and Dog Bile. Journal of Proteome Research, 2009, 8, 2495-2500.	3.7	62
28	Sample preparation optimization in fecal metabolic profiling. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2017, 1047, 115-123.	2.3	62
29	Does the Mass Spectrometer Define the Marker? A Comparison of Global Metabolite Profiling Data Generated Simultaneously via UPLC-MS on Two Different Mass Spectrometers. Analytical Chemistry, 2010, 82, 8226-8234.	6.5	58
30	Metabolic profiling of human urine by CE-MS using a positively charged capillary coating and comparison with UPLC-MS. Molecular BioSystems, 2011, 7, 194-199.	2.9	52
31	Studying the effect of storage conditions on the metabolite content of red wine using HILIC LC–MS based metabolomics. Food Chemistry, 2016, 197, 1331-1340.	8.2	52
32	Analytical and Sample Preparation Techniques for the Determination of Food Colorants in Food Matrices. Foods, 2020, 9, 58.	4.3	52
33	Direct separation and quantitative analysis of thyroxine and triiodothyronine enantiomers in pharmaceuticals by high-performance liquid chromatography. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2004, 800, 193-201.	2.3	50
34	GC-MS analysis of organic acids in human urine in clinical settings: A study of derivatization and other analytical parameters. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2014, 964, 195-201.	2.3	49
35	Amniotic Fluid and Maternal Serum Metabolic Signatures in the Second Trimester Associated with Preterm Delivery. Journal of Proteome Research, 2017, 16, 898-910.	3.7	48
36	Methodological considerations in the development of HPLC-MS methods for the analysis of rodent plasma for metabonomic studies. Molecular BioSystems, 2009, 6, 108-120.	2.9	45

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37	Development of a validated HPLC method for the determination of iodotyrosines and iodothyronines in pharmaceuticals and biological samples using solid phase extraction. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2005, 814, 163-172.	2.3	44
38	Analysis of anaesthetics and analgesics in human urine by headspace SPME and GC. Journal of Separation Science, 2009, 32, 1018-1026.	2.5	43
39	Determination of drugs of abuse and pharmaceuticals in skeletal tissue by UHPLC–MS/MS. Forensic Science International, 2018, 290, 137-145.	2.2	40
40	Emerging Biomarkers for Prediction and Early Diagnosis of Necrotizing Enterocolitis in the Era of Metabolomics and Proteomics. Frontiers in Pediatrics, 2020, 8, 602255.	1.9	38
41	Urine metabolomics in neonates with late-onset sepsis in a case-control study. Scientific Reports, 2017, 7, 45506.	3.3	37
42	Daptomycin determination by liquid chromatography–mass spectrometry in peritoneal fluid, blood plasma, and urine of clinical patients receiving peritoneal dialysis treatment. Analytical and Bioanalytical Chemistry, 2010, 397, 2191-2197.	3.7	36
43	Profiling and biomarker identification in plasma from different Zucker rat strains via high mass accuracy multistage mass spectrometric analysis using liquid chromatography/mass spectrometry with a quadrupole ion trapâ€time of flight mass spectrometer. Rapid Communications in Mass Spectrometry, 2008, 22, 2547-2554.	1.5	35
44	Hydrophilic interaction ultra performance liquid chromatography retention prediction under gradient elution. Analytical and Bioanalytical Chemistry, 2012, 404, 701-709.	3.7	32
45	Development and validation of a fast gas chromatography mass spectrometry method for the quantification of selected non-intentionally added substances and polystyrene/polyurethane oligomers in liquid food simulants. Analytica Chimica Acta, 2020, 1130, 49-59.	5.4	32
46	A hydrophilic interaction chromatography-tandem mass spectrometry method for amino acid profiling in mussels. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2017, 1047, 197-206.	2.3	31
47	Quantification of 15 Psychotropic Drugs in Serum and Postmortem Blood Samples after a Modified Mini-QuEChERS by UHPLC–MS-MS. Journal of Analytical Toxicology, 2018, 42, 337-345.	2.8	31
48	Peak Purity Determination with a Diode Array Detector. Journal of Liquid Chromatography and Related Technologies, 2004, 27, 1083-1092.	1.0	30
49	Quality Control and Validation Issues in LC-MS Metabolomics. Methods in Molecular Biology, 2018, 1738, 15-26.	0.9	28
50	Determination of two COX-2 inhibitors in serum and synovial fluid of patients with inflammatory arthritis by ultra performance liquid chromatography–inductively coupled plasma mass spectroscopy and quadrupole time-of-flight mass spectrometry. Journal of Pharmaceutical and Biomedical Analysis, 2009, 49, 579-586.	2.8	27
51	In Vitro Evaluation of Self-Nano-Emulsifying Drug Delivery Systems (SNEDDS) Containing Room Temperature Ionic Liquids (RTILs) for the Oral Delivery of Amphotericin B. Pharmaceutics, 2020, 12, 699.	4.5	27
52	Analytical Methodologies for the Assessment of Phthalate Exposure in Humans. Critical Reviews in Analytical Chemistry, 2017, 47, 279-297.	3.5	26
53	Polystyrene Biodegradation by Tenebrio molitor Larvae: Identification of Generated Substances Using a GC-MS Untargeted Screening Method. Polymers, 2021, 13, 17.	4.5	26
54	RAPID HPLC ANALYSIS OF THYROID GLAND HORMONES TRI-IODOTHYRONINE (T3) AND THYROXINE (T4) IN HUMAN BIOLOGICAL FLUIDS AFTER SPE. Journal of Liquid Chromatography and Related Technologies, 2000, 23, 681-692.	1.0	24

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55	Sample Preparation Strategies for the Effective Quantitation of Hydrophilic Metabolites in Serum by Multi-Targeted HILIC-MS/MS. Metabolites, 2017, 7, 13.	2.9	24
56	Investigation of chronic alcohol consumption in rodents via ultra-high-performance liquid chromatography–mass spectrometry based metabolite profiling. Journal of Chromatography A, 2012, 1259, 128-137.	3.7	22
57	Computational analysis and ratiometric comparison approaches aimed to assist column selection in hydrophilic interaction liquid chromatography–tandem mass spectrometry targeted metabolomics. Journal of Chromatography A, 2015, 1406, 145-155.	3.7	22
58	Impact of Exercise and Aging on Rat Urine and Blood Metabolome. An LC-MS Based Metabolomics Longitudinal Study. Metabolites, 2017, 7, 10.	2.9	22
59	Development and validation of an ultra high performance liquid chromatography-tandem mass spectrometry method for the determination of phthalate esters in Greek grape marc spirits. Journal of Chromatography A, 2019, 1603, 165-178.	3.7	21
60	An ultra-high pressure liquid chromatography-tandem mass spectrometry method for the quantification of teicoplanin in plasma of neonates. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2017, 1047, 215-222.	2.3	20
61	A UHPLC–MS-MS Method for the Determination of 84 Drugs of Abuse and Pharmaceuticals in Blood. Journal of Analytical Toxicology, 2021, 45, 28-43.	2.8	20
62	QSRR Modeling for Metabolite Standards Analyzed by Two Different Chromatographic Columns Using Multiple Linear Regression. Metabolites, 2017, 7, 7.	2.9	19
63	Targeted profiling of hydrophilic constituents of royal jelly by hydrophilic interaction liquid chromatography–tandem mass spectrometry. Journal of Chromatography A, 2018, 1531, 53-63.	3.7	19
64	A pilot case-control study of urine metabolomics in preterm neonates with necrotizing enterocolitis. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2019, 1117, 10-21.	2.3	19
65	Retention prediction of a set of amino acids under gradient elution conditions in hydrophilic interaction liquid chromatography. Journal of Separation Science, 2012, 35, 376-383.	2.5	18
66	Global metabolic profiling for the study of alcohol-related disorders. Bioanalysis, 2014, 6, 59-77.	1.5	18
67	Impact of exercise on fecal and cecal metabolome over aging: a longitudinal study in rats. Bioanalysis, 2017, 9, 21-36.	1.5	18
68	Rat Fecal Metabolomics-Based Analysis. Methods in Molecular Biology, 2018, 1738, 149-157.	0.9	18
69	Towards the development of Self-Nano-Emulsifying Drug Delivery Systems (SNEDDS) containing trimethyl chitosan for the oral delivery of amphotericin B: In vitro assessment and cytocompatibility studies. Journal of Drug Delivery Science and Technology, 2020, 56, 101524.	3.0	18
70	Prognostic significance of metabolomic biomarkers in patients with diabetes mellitus and coronary artery disease. Cardiovascular Diabetology, 2022, 21, 70.	6.8	18
71	A targeted approach for studying the effect of sugar bee feeding on the metabolic profile of Royal Jelly. Journal of Chromatography A, 2020, 1616, 460783.	3.7	17
72	Study of Fecal and Urinary Metabolite Perturbations Induced by Chronic Ethanol Treatment in Mice by UHPLC-MS/MS Targeted Profiling. Metabolites, 2019, 9, 232.	2.9	16

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73	Development of a UHPLC-MS/MS method for the determination of 84 pharmaceuticals and drugs of abuse in human liver. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2020, 1151, 122192.	2.3	16
74	Solid phase extraction methodology for UPLCâ€MS based metabolic profiling of urine samples. Electrophoresis, 2015, 36, 2170-2178.	2.4	15
75	Correlation of the severity of coronary artery disease with patients' metabolic profile-rationale, design and baseline patient characteristics of the CorLipid trial. BMC Cardiovascular Disorders, 2021, 21, 79.	1.7	15
76	Liquid chromatography tandem mass spectrometry for the determination of nine insecticides and fungicides in human postmortem blood and urine. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2021, 1179, 122824.	2.3	15
77	Is Current Practice Adhering to Guidelines Proposed for Metabolite Identification in LC-MS Untargeted Metabolomics? A Meta-Analysis of the Literature. Journal of Proteome Research, 2022, 21, 590-598.	3.7	15
78	FoodOmicsGR_RI: A Consortium for Comprehensive Molecular Characterisation of Food Products. Metabolites, 2021, 11, 74.	2.9	14
79	Quantification of endogenous aminoacids and aminoacid derivatives in urine by hydrophilic interaction liquid chromatography tandem mass spectrometry. Journal of Chromatography A, 2021, 1642, 462005.	3.7	14
80	Headspace gas chromatography-mass spectrometry in the analysis of lavender's essential oil: Optimization by response surface methodology. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2021, 1179, 122852.	2.3	14
81	A fast SALLE GC–MS/MS multi-analyte method for the determination of 75 food packaging substances in food simulants. Food Chemistry, 2021, 361, 129998.	8.2	14
82	Liquid chromatography-mass spectrometry method for the determination of polyethylene terephthalate and polybutylene terephthalate cyclic oligomers in blood samples. Analytical and Bioanalytical Chemistry, 2022, 414, 1503-1512.	3.7	14
83	HILIC-MS/MS Multi-Targeted Method for Metabolomics Applications. Methods in Molecular Biology, 2018, 1738, 65-81.	0.9	13
84	NSAIDs Determination in Human Serum by GC-MS. Separations, 2018, 5, 37.	2.4	13
85	Urine and fecal samples targeted metabolomics of carobs treated rats. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2019, 1114-1115, 76-85.	2.3	13
86	Correlation of Serum Acylcarnitines with Clinical Presentation and Severity of Coronary Artery Disease. Biomolecules, 2022, 12, 354.	4.0	13
87	GC-MS analysis of underivatised new psychoactive substances in whole blood and urine. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2020, 1156, 122308.	2.3	12
88	Serum Ceramides as Prognostic Biomarkers of Large Thrombus Burden in Patients with STEMI: A Micro-Computed Tomography Study. Journal of Personalized Medicine, 2021, 11, 89.	2.5	12
89	Metabolic Phenotyping Study of Mouse Brains Following Acute or Chronic Exposures to Ethanol. Journal of Proteome Research, 2020, 19, 4071-4081.	3.7	11
90	Development and validation of an UHPLC-qTOF-MS method for the quantification of cyclic polyesters oligomers in pasta by applying a modified QuEChERS clean-up. Food Chemistry, 2021, 347, 129040.	8.2	11

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91	Analysis of urinary organic acids by gas chromatography tandem mass spectrometry method for metabolic profiling applications. Journal of Chromatography A, 2021, 1658, 462590.	3.7	11
92	Gut Microbiome and Degradation Product Formation during Biodegradation of Expanded Polystyrene by Mealworm Larvae under Different Feeding Strategies. Molecules, 2021, 26, 7568.	3.8	11
93	Metabolic profiling study of shikonin's cytotoxic activity in the Huh7 human hepatoma cell line. Molecular BioSystems, 2017, 13, 841-851.	2.9	10
94	Development and validation of a RPLC-MS/MS method for the quantification of ceramides in human serum. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2021, 1175, 122734.	2.3	10
95	Serum-Targeted HILIC-MS Metabolomics-Based Analysis in Infants with Ureteropelvic Junction Obstruction. Journal of Proteome Research, 2020, 19, 2294-2303.	3.7	9
96	Metabolic Profiling: Status, Challenges, and Perspective. Methods in Molecular Biology, 2018, 1738, 3-13.	0.9	8
97	Population Pharmacokinetics of Teicoplanin in Preterm and Term Neonates: Is It Time for a New Dosing Regimen?. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	8
98	Association of GRACE Risk Score with Coronary Artery Disease Complexity in Patients with Acute Coronary Syndrome. Journal of Clinical Medicine, 2021, 10, 2210.	2.4	8
99	Application of a hybrid zwitterionic hydrophilic interaction liquid chromatography column in metabolic profiling studies. Journal of Chromatography A, 2022, 1672, 463013.	3.7	8
100	Population Pharmacokinetics and Outcomes of Critically III Pediatric Patients Treated with Intravenous Colistin at Higher Than Recommended Doses. Antimicrobial Agents and Chemotherapy, 2021, 65, .	3.2	7
101	Evaluation of Cocaine Effect on Endogenous Metabolites of HepG2 Cells Using Targeted Metabolomics. Molecules, 2021, 26, 4610.	3.8	7
102	Impact of religious fasting on metabolic and hematological profile in both dyslipidemic and non-dyslipidemic fasters. European Journal of Clinical Nutrition, 2022, 76, 891-898.	2.9	7
103	Development, Validation and Application of an Ultra-High-Performance Liquid Chromatography–Tandem Mass Spectrometry (UHPLC-MS/MS) Method after QuEChERS Cleanup for Selected Dichloroanilines and Phthalates in Rice Samples. Foods, 2022, 11, 1482.	4.3	7
104	Development and validation of LC-MS/MS method for the determination of UV-filters across human skin in vitro. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2021, 1167, 122561.	2.3	6
105	Risk factors for fatal drowning in a Greek region: a retrospective case–control study. Injury Prevention, 2021, 27, injuryprev-2020-043788.	2.4	5
106	In vivo study of pro-inflammatory cytokine changes in serum and synovial fluid during treatment with celecoxib and etoricoxib and correlation with VAS pain change and synovial membrane penetration index in patients with inflammatory arthritis. Mediterranean Journal of Rheumatology, 2017, 28, 33-40.	0.8	5
107	Plasma Lipidomic and Metabolomic Profiling after Birth in Neonates Born to SARS-CoV-19 Infected and Non-Infected Mothers at Delivery: Preliminary Results. Metabolites, 2021, 11, 830.	2.9	5
108	HILIC-MS/MS Analysis of Adenosine in Patient Blood. Separations, 2021, 8, 222.	2.4	5

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109	Alprazolam and Zolpidem in Skeletal Tissue of Decomposed Body Confirms Exposure. Journal of Forensic Sciences, 2019, 64, 643-646.	1.6	4
110	Î' Simple Method for the Determination of Lacosamide in Blood by GC ―MS. Journal of Forensic Sciences, 2020, 65, 288-294.	1.6	4
111	Effect of exercise on key pharmacokinetic parameters related to metformin absorption in healthy humans: A pilot study. Scandinavian Journal of Medicine and Science in Sports, 2020, 30, 858-864.	2.9	4
112	Multitargeted hydrophilic interaction chromatography-MS/MS: limitations and perspectives. Bioanalysis, 2018, 10, 1165-1167.	1.5	3
113	GC-MS-Based Metabolic Phenotyping. , 2019, , 137-169.		3
114	A Study of Blood Fatty Acids Profile in Hyperlipidemic and Normolipidemic Subjects in Association with Common PNPLA3 and ABCB1 Polymorphisms. Metabolites, 2021, 11, 90.	2.9	3
115	Diminished Systemic Amino Acids Metabolome and Lipid Peroxidation in Ureteropelvic Junction Obstruction (UPJO) Infants Requiring Surgery. Journal of Clinical Medicine, 2021, 10, 1467.	2.4	3
116	Impact of Metabolomics Technologies on the Assessment of Peritoneal Membrane Profiles in Peritoneal Dialysis Patients: A Systematic Review. Metabolites, 2022, 12, 145.	2.9	3
117	A HILIC-MS/MS method development and validation for the quantitation of 13 acylcarnitines in human serum. Analytical and Bioanalytical Chemistry, 2022, 414, 3095-3108.	3.7	3
118	Development and Validation of a Single Step GC/MS Method for the Determination of 41 Drugs and Drugs of Abuse in Postmortem Blood. Forensic Sciences, 2022, 2, 473-491.	1.5	3
119	Liquid Chromatographic Methods Combined with Mass Spectrometry inÂMetabolomics. , 2013, , 145-161.		2
120	The Role of Mass Spectrometry in Nontargeted Metabolomics. Comprehensive Analytical Chemistry, 2014, , 213-233.	1.3	2
121	Liquid chromatographic methods combined with mass spectrometry in metabolomics. , 2020, , 149-169.		2
122	State-of-the-art in LC–MS Approaches for Probing the Polar Metabolome. New Developments in Mass Spectrometry, 2021, , 1-26.	0.2	2
123	Syncope without prodromes is associated with excessive plasma release of adenosine at the time of syncope during head-up tilt table test. International Journal of Cardiology, 2022, 363, 43-48.	1.7	2
124	Metabolic Profiling Approaches for Biomarkers of Ethanol Intake. , 2016, , 213-222.		1
125	Metabolomics: An Analytical Perspective. , 2018, , 82-82.		1
126	Development and Validation of a UHPLC-qTOF MS Method for the Determination of Sorbitol-Based Nuclear Clarifying Agents in Food Simulants after Migration from Food Contact Materials. Applied Sciences (Switzerland), 2021, 11, 3789.	2.5	1

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127	Investigation of salivary biomarkers as indicators of skeletal and dental maturity in children. Orthodontics and Craniofacial Research, 2022, , .	2.8	1
128	Efficacy and safety of Mydriatic Microdrops for Retinopathy Of Prematurity Screening (MyMiROPS): study protocol for a non-inferiority crossover randomized controlled trial. Trials, 2022, 23, 322.	1.6	1
129	Metabolic phenotyping (metabonomics/metabolomics) by liquid chromatography-mass spectrometry. , 2017, , 245-265.		0
130	Preface. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2019, 1128, 121787.	2.3	0
131	Cutting-edge analytical technologies for the comprehensive metabolic profiling of Alkanna tinctoria roots cultured in greenhouse conditions. , 2019, 85, .		O
132	A hydrophilic liquid chromatography – tandem mass spectrometry method for the determination of phenylephrine in dried blood spots from preterm infants. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2022, 1190, 123084.	2.3	0
133	Detection and determination of C12â€, C14â€, C16â€alkyldimethylamines in human blood using gas chromatography–mass spectrometry. Rapid Communications in Mass Spectrometry, 2022, 36, e9303.	1.5	0