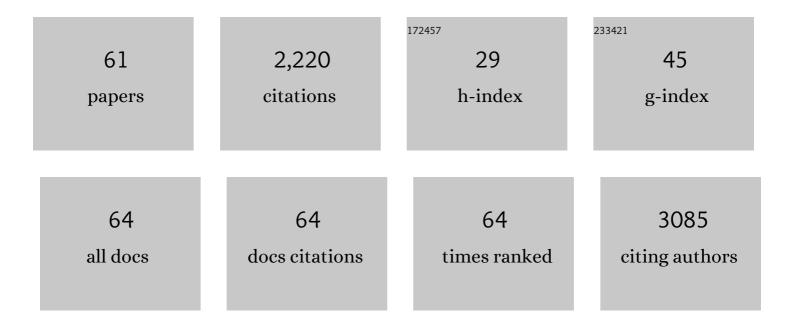
## Margherita Ruoppolo

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Clinical validation of cutoff target ranges in newborn screening of metabolic disorders by tandem mass spectrometry: A worldwide collaborative project. Genetics in Medicine, 2011, 13, 230-254.	2.4	308
2	Enhanced interpretation of newborn screening results without analyte cutoff values. Genetics in Medicine, 2012, 14, 648-655.	2.4	117
3	Dysregulation of lipid metabolism and pathological inflammation in patients with COVID-19. Scientific Reports, 2021, 11, 2941.	3.3	102
4	Pro-sequence assisted folding and disulfide bond formation of human nerve growth factorâ€â€Dedicated to Rita Levi-Montalcini.22Edited by R. Huber. Journal of Molecular Biology, 2001, 305, 523-533.	4.2	90
5	Targeted metabolomics in the expanded newborn screening for inborn errors of metabolism. Molecular BioSystems, 2015, 11, 1525-1535.	2.9	73
6	"Classical organic acidurias― diagnosis and pathogenesis. Clinical and Experimental Medicine, 2017, 17, 305-323.	3.6	69
7	Proteomics Identification of Acyl-acceptor and Acyl-donor Substrates for Transglutaminase in a Human Intestinal Epithelial Cell Line. Journal of Biological Chemistry, 2003, 278, 31766-31773.	3.4	62
8	The Serum Metabolome of Moderate and Severe COVID-19 Patients Reflects Possible Liver Alterations Involving Carbon and Nitrogen Metabolism. International Journal of Molecular Sciences, 2021, 22, 9548.	4.1	56
9	Folding and Oxidation of the Antibody Domain CH3. Journal of Molecular Biology, 2002, 319, 1267-1277.	4.2	53
10	Maternal vitamin B12 deficiency detected in expanded newborn screening. Clinical Biochemistry, 2014, 47, 312-317.	1.9	53
11	Protein crossâ€ŧalk in <scp>CD</scp> 133+ colon cancer cells indicates activation of the <scp>W</scp> nt pathway and upregulation of <scp>SR</scp> p20 that is potentially involved in tumorigenicity. Proteomics, 2012, 12, 2045-2059.	2.2	52
12	Protein–protein interaction networks as a new perspective to evaluate distinct functional roles of voltage-dependent anion channel isoforms. Molecular BioSystems, 2017, 13, 2466-2476.	2.9	50
13	Analysis of the Ribosomal Protein S19 Interactome. Molecular and Cellular Proteomics, 2007, 6, 382-393.	3.8	49
14	COVIDomics: The Proteomic and Metabolomic Signatures of COVID-19. International Journal of Molecular Sciences, 2022, 23, 2414.	4.1	49
15	Identification and Characterization of Structural Domains of Human ERp57. Journal of Biological Chemistry, 2004, 279, 13607-13615.	3.4	47
16	Human Serum Albumin Is an Essential Component of the Host Defense Mechanism Against Clostridium difficile Intoxication. Journal of Infectious Diseases, 2018, 218, 1424-1435.	4.0	45
17	Female and male human babies have distinct blood metabolomic patterns. Molecular BioSystems, 2015, 11, 2483-2492.	2.9	40
18	Targeted metabolomic profiling in rat tissues reveals sex differences. Scientific Reports, 2018, 8, 4663.	3.3	40

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19	Sex differences in the human metabolome. Biology of Sex Differences, 2022, 13, .	4.1	38
20	Proteomics Reveals that Methylmalonyl-CoA Mutase Modulates Cell Architecture and Increases Susceptibility to Stress. International Journal of Molecular Sciences, 2020, 21, 4998.	4.1	36
21	The proteome of cblC defect: in vivo elucidation of altered cellular pathways in humans. Journal of Inherited Metabolic Disease, 2015, 38, 969-979.	3.6	34
22	Influence of Sex on Urinary Organic Acids: A Cross-Sectional Study in Children. International Journal of Molecular Sciences, 2020, 21, 582.	4.1	33
23	Serum metabolomic profiles suggest influence of sex and oral contraceptive use. American Journal of Translational Research (discontinued), 2014, 6, 614-24.	0.0	33
24	Mass Spectrometry-Based Metabolomic and Proteomic Strategies in Organic Acidemias. BioMed Research International, 2016, 2016, 1-13.	1.9	32
25	Label-Free Quantitative Proteomics in a Methylmalonyl-CoA Mutase-Silenced Neuroblastoma Cell Line. International Journal of Molecular Sciences, 2018, 19, 3580.	4.1	32
26	Targeted Metabolomic Analysis of a Mucopolysaccharidosis IIIB Mouse Model Reveals an Imbalance of Branched-Chain Amino Acid and Fatty Acid Metabolism. International Journal of Molecular Sciences, 2020, 21, 4211.	4.1	32
27	Characterization of low-molecular-mass trypsin isoinhibitors from oil-rape (Brassica napus var.) Tj ETQq1 1 0.78	4314 rgBT 0.2	/Oyerlock 10
28	The first case of mitochondrial acetoacetyl-CoA thiolase deficiency identified by expanded newborn metabolic screening in Italy: the importance of an integrated diagnostic approach. Journal of Inherited Metabolic Disease, 2010, 33, 91-94.	3.6	30
29	The proteome of methylmalonic acidemia (MMA): the elucidation of altered pathways in patient livers. Molecular BioSystems, 2016, 12, 566-574.	2.9	30
30	Proteomic Analysis of Mucopolysaccharidosis IIIB Mouse Brain. Biomolecules, 2020, 10, 355.	4.0	30
31	Challenges in Metabolomics-Based Tests, Biomarkers Revealed by Metabolomic Analysis, and the Promise of the Application of Metabolomics in Precision Medicine. International Journal of Molecular Sciences, 2022, 23, 5213.	4.1	30
32	Long-term follow-up of patients with phenylketonuria treated with tetrahydrobiopterin: a seven years experience. Orphanet Journal of Rare Diseases, 2015, 10, 14.	2.7	29
33	Urine Proteomics Revealed a Significant Correlation Between Urine-Fibronectin Abundance and Estimated-GFR Decline in Patients with Bardet-Biedl Syndrome. Kidney and Blood Pressure Research, 2018, 43, 389-405.	2.0	28
34	Integration of Proteomics and Metabolomics in Exploring Genetic and Rare Metabolic Diseases. Kidney Diseases (Basel, Switzerland), 2017, 3, 66-77.	2.5	26
35	Citrulline Blood Levels as Indicators of Residual Intestinal Absorption in Patients with Short Bowel Syndrome. Annals of Nutrition and Metabolism, 2008, 53, 137-142.	1.9	25
36	Transcription Factor TBX1 Overexpression Induces Downregulation of Proteins Involved in Retinoic Acid Metabolism: A Comparative Proteomic Analysis. Journal of Proteome Research, 2009, 8, 1515-1526.	3.7	25

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37	Insulinâ€resistance in glycogen storage disease type Ia: linking carbohydrates and mitochondria?. Journal of Inherited Metabolic Disease, 2018, 41, 985-995.	3.6	24
38	The first knock-in rat model for glutaric aciduria type I allows further insights into pathophysiology in brain and periphery. Molecular Genetics and Metabolism, 2021, 133, 157-181.	1.1	22
39	Differential proteomic analysis in human cells subjected to ribosomal stress. Proteomics, 2013, 13, 1220-1227.	2.2	20
40	Optimization of an HPLC method for phenylalanine and tyrosine quantization in dried blood spot. Clinical Biochemistry, 2013, 46, 1892-1895.	1.9	18
41	The saturation degree of fatty acids and their derived acylcarnitines determines the direct effect of metabolically active thyroid hormones on insulin sensitivity in skeletal muscle cells. FASEB Journal, 2019, 33, 1811-1823.	0.5	18
42	Dataset of a comparative proteomics experiment in a methylmalonyl-CoA mutase knockout HEK 293 cell model. Data in Brief, 2020, 33, 106453.	1.0	18
43	Proteomic and Bioinformatic Investigation of Altered Pathways in Neuroglobin-Deficient Breast Cancer Cells. Molecules, 2021, 26, 2397.	3.8	18
44	Sex Affects Human Premature Neonates' Blood Metabolome According to Gestational Age, Parenteral Nutrition, and Caffeine Treatment. Metabolites, 2021, 11, 158.	2.9	17
45	A knock-in rat model unravels acute and chronic renal toxicity in glutaric aciduria type I. Molecular Genetics and Metabolism, 2021, 134, 287-300.	1.1	17
46	Analysis of the interactome of ribosomal protein S19 mutants. Proteomics, 2014, 14, 2286-2296.	2.2	16
47	Proteomics and metabolomics studies exploring the pathophysiology of renal dysfunction in autosomal dominant polycystic kidney disease and other ciliopathies. Nephrology Dialysis Transplantation, 2020, 35, 1853-1861.	0.7	16
48	Gaining insights into the Bcr-Abl activity-independent mechanisms of resistance to imatinib mesylate in KCL22 cells: A comparative proteomic approach. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2010, 1804, 1974-1987.	2.3	15
49	Exercise with food withdrawal at thermoneutrality impacts fuel use, the microbiome, AMPK phosphorylation, muscle fibers, and thyroid hormone levels in rats. Physiological Reports, 2020, 8, e14354.	1.7	15
50	Galactosemia: Biochemistry, Molecular Genetics, Newborn Screening, and Treatment. Biomolecules, 2022, 12, 968.	4.0	15
51	Overexpression of Neuroglobin Promotes Energy Metabolism and Autophagy Induction in Human Neuroblastoma SH-SY5Y Cells. Cells, 2021, 10, 3394.	4.1	14
52	Biochemical and molecular characterization of 3-Methylcrotonylglycinuria in an Italian asymptomatic girl. Genetics and Molecular Biology, 2018, 41, 379-385.	1.3	8
53	Proteome data of neuroblastoma cells overexpressing Neuroglobin. Data in Brief, 2022, 41, 107843.	1.0	8
54	Unravelling pathways downstream Sox6 induction in K562 erythroid cells by proteomic analysis. Scientific Reports, 2017, 7, 14088.	3.3	6

#	Article	IF	CITATIONS
55	Deregulation of microtubule organization and RNA metabolism in <i>Arx</i> models for lissencephaly and developmental epileptic encephalopathy. Human Molecular Genetics, 2022, 31, 1884-1908.	2.9	6
56	Quantification of Imatinib Plasma Levels in Patients with Chronic Myeloid Leukemia: Comparison Between HPLC–UV and LC–MS/MS. International Journal of Peptide Research and Therapeutics, 2013, 19, 109-116.	1.9	5
57	<scp>ZSCAN</scp> 4 <sup>+</sup> mouse embryonic stem cells have an oxidative and flexible metabolic profile. EMBO Reports, 2020, 21, e48942.	4.5	5
58	Targeted metabolomics. , 2022, , 219-236.		4
59	Hypermethioninemia in Campania: Results from 10†years of newborn screening. Molecular Genetics and Metabolism Reports, 2019, 21, 100520.	1.1	2
60	Beneficial Effects of Slow-Release Large Neutral Amino Acids after a Phenylalanine Oral Load in Patients with Phenylketonuria. Nutrients, 2021, 13, 4012.	4.1	2
61	Editorial for Special Issue: Neuroglobin from Brain Protection to Cancer Progression. Cells, 2022, 11, 2181.	4.1	0