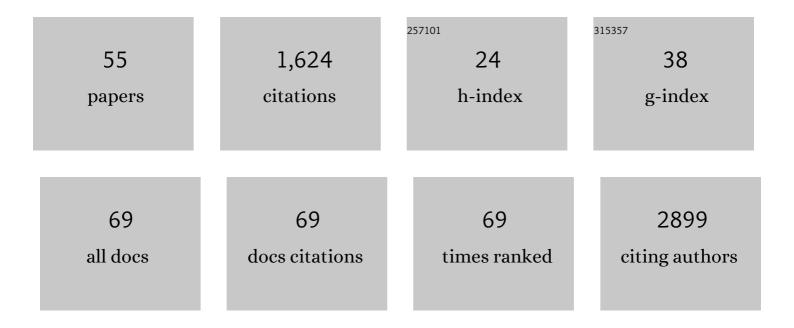
Chiara Piubelli

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5143034/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	SARSâ€CoVâ€2 rapid antigen test in comparison to RTâ€PCR targeting different genes: A realâ€life evaluation among unselected patients in a regional hospital of Italy. Journal of Medical Virology, 2022, 94, 1190-1195.	2.5	17
2	SARS-CoV-2 vaccination elicits unconventional IgM specific responses in naÃ ⁻ ve and previously COVID-19-infected individuals. EBioMedicine, 2022, 77, 103888.	2.7	39
3	Tropheryma whipplei, Helicobacter pylori, and Intestinal Protozoal Co-Infections in Italian and Immigrant Populations: A Cross-Sectional Study. Microorganisms, 2022, 10, 769.	1.6	4
4	Overall decrease in SARS-CoV-2 viral load and reduction in clinical burden: the experience of a hospital in northern Italy. Clinical Microbiology and Infection, 2021, 27, 131.e1-131.e3.	2.8	17
5	SARS-CoV-2 Epitope Mapping on Microarrays Highlights Strong Immune-Response to N Protein Region. Vaccines, 2021, 9, 35.	2.1	38
6	Antibody Response to the BNT162b2 mRNA COVID-19 Vaccine in Subjects with Prior SARS-CoV-2 Infection. Viruses, 2021, 13, 422.	1.5	140
7	Preliminary comparison between an in-house real-time PCR vs microscopy for the diagnosis of Loa loa and Mansonella perstans. Acta Tropica, 2021, 216, 105838.	0.9	4
8	Molecular techniques for the genomic viral RNA detection of West Nile, Dengue, Zika and Chikungunya arboviruses: a narrative review. Expert Review of Molecular Diagnostics, 2021, 21, 591-612.	1.5	5
9	ACoRE: Accurate SARS-CoV-2 genome reconstruction for the characterization of intra-host and inter-host viral diversity in clinical samples and for the evaluation of re-infections. Genomics, 2021, 113, 1628-1638.	1.3	8
10	Antibody response induced by the BNT162b2 mRNA COVID-19 vaccine in a cohort of health-care workers, with or without prior SARS-CoV-2 infection: a prospective study. Clinical Microbiology and Infection, 2021, 27, 1845-1850.	2.8	44
11	Preliminary evaluation of a new Schistosoma Immunochromatographic Test. Acta Tropica, 2021, 219, 105893.	0.9	0
12	Antibody response in individuals infected with SARS-CoV-2 early after the first dose of the BNT162b2 mRNA vaccine. Journal of Infection, 2021, , .	1.7	3
13	Impact of Full Vaccination with mRNA BNT162b2 on SARS-CoV-2 Infection: Genomic and Subgenomic Viral RNAs Detection in Nasopharyngeal Swab and Saliva of Health Care Workers. Microorganisms, 2021, 9, 1738.	1.6	4
14	Monoclonal Antibodies for Protozoan Infections: A Future Reality or a Utopic Idea?. Frontiers in Medicine, 2021, 8, 745665.	1.2	4
15	Serology study after BTN162b2 vaccination in participants previously infected with SARS-CoV-2 in two different waves versus naÃ ⁻ ve. Communications Medicine, 2021, 1, .	1.9	18
16	Prosthetic joint infection due to Mycobacterium xenopi: a review of the literature with a new case report. Infection, 2020, 48, 165-171.	2.3	7
17	Leprosy in Refugees and Migrants in Italy and a Literature Review of Cases Reported in Europe between 2009 and 2018. Microorganisms, 2020, 8, 1113.	1.6	7
18	Assessment of the direct quantitation of SARS-CoV-2 by droplet digital PCR. Scientific Reports, 2020, 10, 18764.	1.6	38

CHIARA PIUBELLI

#	Article	IF	CITATIONS
19	Sensitivity, Specificity and Predictive Values of Molecular and Serological Tests for COVID-19: A Longitudinal Study in Emergency Room. Diagnostics, 2020, 10, 669.	1.3	34
20	Systemic profile of immune factors in an elderly Italian population affected by chronic strongyloidiasis. Parasites and Vectors, 2020, 13, 515.	1.0	4
21	Genetic and Clinical Heterogeneity in Thirteen New Cases with Aceruloplasminemia. Atypical Anemia as a Clue for an Early Diagnosis. International Journal of Molecular Sciences, 2020, 21, 2374.	1.8	25
22	Dynamics of anti-malarial antibodies in non-immune patients during and after a first and unique Plasmodium falciparum malaria episode. Malaria Journal, 2020, 19, 228.	0.8	5
23	Droplet Digital PCR for the Detection of Plasmodium falciparum DNA in Whole Blood and Serum: A Comparative Analysis with Other Molecular Methods. Pathogens, 2020, 9, 478.	1.2	8
24	Digital PCR: a new technology for diagnosis of parasitic infections. Clinical Microbiology and Infection, 2019, 25, 1510-1516.	2.8	49
25	Blastocystis prevalence and subtypes in autochthonous and immigrant patients in a referral centre for parasitic infections in Italy. PLoS ONE, 2019, 14, e0210171.	1.1	24
26	An orally available, brain-penetrant CAMKK2 inhibitor reduces food intake in rodent model. Bioorganic and Medicinal Chemistry Letters, 2018, 28, 1958-1963.	1.0	20
27	A comprehensive analysis of the faecal microbiome and metabolome of Strongyloides stercoralis infected volunteers from a non-endemic area. Scientific Reports, 2018, 8, 15651.	1.6	51
28	Preliminary Comparison of an in-House Real-Time PCR with the Automated BD Max Enteric Parasite Panel for the Detection of <i>Giardia intestinalis</i> . Journal of Parasitology, 2018, 104, 702-704.	0.3	0
29	The arrhythmogenic cardiomyopathy-specific coding and non-coding transcriptome in human cardiac stromal cells. BMC Genomics, 2018, 19, 491.	1.2	21
30	HDAC Inhibition Improves the Sarcoendoplasmic Reticulum Ca2+-ATPase Activity in Cardiac Myocytes. International Journal of Molecular Sciences, 2018, 19, 419.	1.8	21
31	Accuracy of molecular biology techniques for the diagnosis of Strongyloides stercoralis infection—A systematic review and meta-analysis. PLoS Neglected Tropical Diseases, 2018, 12, e0006229.	1.3	104
32	A decade of progress on the genetic basis of coronary artery disease. Practical insights for the internist. European Journal of Internal Medicine, 2017, 41, 10-17.	1.0	14
33	Identification of new BMP6 proâ€peptide mutations in patients with iron overload. American Journal of Hematology, 2017, 92, 562-568.	2.0	35
34	Identification of New BMP6 Pro-Peptide Mutations in Patients with Unexplained Iron-Overload. Blood, 2016, 128, 264-264.	0.6	0
35	microRNAs and Cardiac Cell Fate. Cells, 2014, 3, 802-823.	1.8	38
36	Proteomics of rat hypothalamus, hippocampus and pre-frontal/frontal cortex after central administration of the neuropeptide PACAP. Molecular Biology Reports, 2012, 39, 2921-2935.	1.0	13

CHIARA PIUBELLI

#	Article	IF	CITATIONS
37	Escitalopram modulates neuron-remodelling proteins in a rat gene–environment interaction model of depression as revealed by proteomics. Part I: genetic background. International Journal of Neuropsychopharmacology, 2011, 14, 796-833.	1.0	20
38	Regulation of cytoskeleton machinery, neurogenesis and energy metabolism pathways in a rat gene-environment model of depression revealed by proteomic analysis. Neuroscience, 2011, 176, 349-380.	1.1	42
39	Nortriptyline influences protein pathways involved in carbohydrate metabolism and actin-related processes in a rat gene–environment model of depression. European Neuropsychopharmacology, 2011, 21, 545-562.	0.3	21
40	Escitalopram affects cytoskeleton and synaptic plasticity pathways in a rat gene–environment interaction model of depression as revealed by proteomics. Part II: environmental challenge. International Journal of Neuropsychopharmacology, 2011, 14, 834-855.	1.0	26
41	Proteome Effects of Antidepressant Medications. Advances in Neurobiology, 2011, , 399-441.	1.3	Ο
42	Early-life stress and antidepressants modulate peripheral biomarkers in a gene–environment rat model of depression. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2010, 34, 1037-1048.	2.5	78
43	Expression Profiling of a Genetic Animal Model of Depression Reveals Novel Molecular Pathways Underlying Depressive-Like Behaviours. PLoS ONE, 2010, 5, e12596.	1.1	33
44	P.2.d.007 Proteomic analysis of hippocampus and frontal cortex in a rat model of depression with gene-environment interaction and antidepressant treatment. European Neuropsychopharmacology, 2006, 16, S336-S337.	0.3	1
45	Proteomic analysis of rat hippocampus and frontal cortex after chronic treatment with fluoxetine or putative novel antidepressants: CRF1 and NK1 receptor antagonists. European Neuropsychopharmacology, 2006, 16, 521-537.	0.3	56
46	Proteomic analysis of rat hippocampus after repeated psychosocial stress. Neuroscience, 2006, 137, 1237-1246.	1.1	70
47	Searching for markers of Creutzfeldt-Jakob disease in cerebrospinal fluid by two-dimensional mapping. Proteomics, 2006, 6, S256-S261.	1.3	36
48	Proteomic changes in rat serum, polymorphonuclear and mononuclear leukocytes after chronic nicotine administration. Proteomics, 2005, 5, 1382-1394.	1.3	16
49	Proteome analysis in the clinical chemistry laboratory: Myth or reality?. Clinica Chimica Acta, 2005, 357, 123-139.	0.5	99
50	Critical survey of quantitative proteomics in two-dimensional electrophoretic approaches. Journal of Chromatography A, 2004, 1051, 3-17.	1.8	100
51	The Proteome: Anno Domini 2002. Clinical Chemistry and Laboratory Medicine, 2003, 41, 425-38.	1.4	31
52	Development of a High-Throughput Scintillation Proximity Assay for the Identification of C-Domain Translational Initiation Factor 2 Inhibitors. Journal of Biomolecular Screening, 2002, 7, 541-546.	2.6	6
53	Proteome analysis of rat polymorphonuclear leukocytes: A two-dimensional electrophoresis/ mass spectrometry approach. Electrophoresis, 2002, 23, 298-310.	1.3	35
54	Proteomic analysis of rat brain tissue: Comparison of protocols for two-dimensional gel electrophoresis analysis based on different solubilizing agents. Electrophoresis, 2002, 23, 4132-4141.	1.3	43

#	Article	IF	CITATIONS
55	Effect of experimental conditions on the analysis of sodium dodecyl sulphate polyacrylamide gel electrophoresis separated proteins by matrix-assisted laser desorption/ ionisation mass spectrometry. , 2000, 14, 18-25.		35