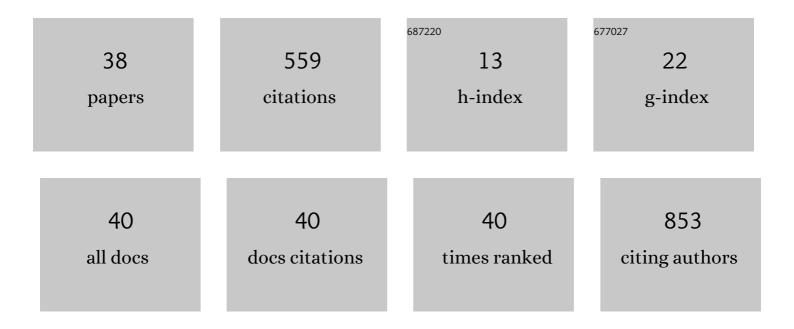
Fani Chatzopoulou

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Differential expression of the ascorbate oxidase multigene family during fruit development and in response to stress. Planta, 2007, 225, 873-885.	1.6	67
2	Towards Elucidating Carnosic Acid Biosynthesis in Lamiaceae: Functional Characterization of the Three First Steps of the Pathway in Salvia fruticosa and Rosmarinus officinalis. PLoS ONE, 2015, 10, e0124106.	1.1	67
3	Genetic justification of severe COVID-19 using a rigorous algorithm. Clinical Immunology, 2021, 226, 108726.	1.4	47
4	Genetic prediction of ICU hospitalization and mortality in COVIDâ€19 patients using artificial neural networks. Journal of Cellular and Molecular Medicine, 2022, 26, 1445-1455.	1.6	45
5	EST analysis and annotation of transcripts derived from a trichome-specific cDNA library from Salvia fruticosa. Plant Cell Reports, 2010, 29, 523-534.	2.8	33
6	Salivary mRNA markers having the potential to detect oral squamous cell carcinoma segregated from oral leukoplakia with dysplasia. Cancer Epidemiology, 2016, 43, 112-118.	0.8	25
7	Silencing of ascorbate oxidase results in reduced growth, altered ascorbic acid levels and ripening pattern in melon fruit. Plant Physiology and Biochemistry, 2020, 156, 291-303.	2.8	21
8	First Report of Isolation and Genome Sequence of L. petauri Strain from a Rainbow Trout Lactococcosis Outbreak. Current Microbiology, 2020, 77, 1089-1096.	1.0	21
9	Antibacterial activity of griseofulvin analogues as an example of drug repurposing. International Journal of Antimicrobial Agents, 2020, 55, 105884.	1.1	19
10	Increased thrombotic and impaired fibrinolytic response to acute exercise in patients with essential hypertension: The effect of treatment with an angiotensin II receptor blocker. Journal of Human Hypertension, 2014, 28, 606-609.	1.0	17
11	Towards analyzing the potential of exosomes to deliver microRNA therapeutics. Journal of Cellular Physiology, 2021, 236, 1529-1544.	2.0	17
12	Serum humanin concentrations in women with pre-eclampsia compared to women with uncomplicated pregnancies. Journal of Maternal-Fetal and Neonatal Medicine, 2018, 31, 305-311.	0.7	15
13	OPG/RANK/RANKL signaling axis in patients with type I diabetes: Associations with parathormone and vitamin D. Italian Journal of Pediatrics, 2019, 45, 161.	1.0	14
14	Association Between Cardiotrophin 1 Levels and Central Blood Pressure in Untreated Patients With Essential Hypertension. American Journal of Hypertension, 2014, 27, 651-655.	1.0	13
15	Emergence of Klebsiella pneumoniae ST11 co-producing NDM-1 and OXA-48 carbapenemases in Greece. Journal of Global Antimicrobial Resistance, 2019, 19, 81-82.	0.9	12
16	Identification of complement-related host genetic risk factors associated with influenza A(H1N1)pdm09 outcome: challenges ahead. Medical Microbiology and Immunology, 2019, 208, 631-640.	2.6	12
17	The GEnetic Syntax Score: a genetic risk assessment implementation tool grading the complexity of coronary artery disease—rationale and design of the GESS study. BMC Cardiovascular Disorders, 2021, 21, 284.	0.7	11
18	A Risk-Stratification Machine Learning Framework for the Prediction of Coronary Artery Disease Severity: Insights From the GESS Trial. Frontiers in Cardiovascular Medicine, 2021, 8, 812182.	1.1	11

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19	Dissecting miRNA–Gene Networks to Map Clinical Utility Roads of Pharmacogenomics-Guided Therapeutic Decisions in Cardiovascular Precision Medicine. Cells, 2022, 11, 607.	1.8	11
20	Whole Genome Sequencing of NDM-1-Producing ST11 <i>Klebsiella pneumoniae</i> Isolated in a Private Laboratory in Greece. Microbial Drug Resistance, 2019, 25, 80-86.	0.9	10
21	Th2/Th17 cytokine profile in phenotyped Greek asthmatics and relationship to biomarkers of inflammation. Respiratory Medicine, 2019, 151, 102-110.	1.3	9
22	Immune Response of Adult Sickle Cell Disease Patients after COVID-19 Vaccination: The Experience of a Greek Center. Journal of Clinical Medicine, 2022, 11, 937.	1.0	7
23	Uncovering the pharmacological response of novel sesquiterpene derivatives that differentially alter gene expression and modulate the cell cycle in cancer cells. International Journal of Oncology, 2018, 53, 2167-2179.	1.4	6
24	Whole-genome sequencing study of KPC-encoding Klebsiella pneumoniae isolated in Greek private laboratories from non-hospitalised patients. Journal of Global Antimicrobial Resistance, 2020, 20, 78-81.	0.9	6
25	Neutralizing antibody and T cell responses to SARS-CoV-2 vaccination in hematopoietic cell transplant recipients. Bone Marrow Transplantation, 2022, 57, 1183-1186.	1.3	6
26	mcr Genes Conferring Colistin Resistance in Enterobacterales; a Five Year Overview. Acta Medica Academica, 2022, 50, 365.	0.3	6
27	Robust SARS-COV-2-specific T-cell immune memory persists long-term in immunocompetent individuals post BNT162b2 double shot. Heliyon, 2022, 8, e09863.	1.4	5
28	Epidemiological Study and Classification of ICU Infections, Using the Carrier State Criterion. European Journal of Inflammation, 2010, 8, 181-188.	0.2	4
29	Clonal outbreak caused by VIM-4-producing Proteus mirabilis in a Greek tertiary-care hospital. International Journal of Antimicrobial Agents, 2020, 56, 106060.	1.1	4
30	An initially unidentified case of urinary tract infection due to Aerococcus urinae. New Microbiologica, 2017, 40, 221-222.	0.1	4
31	Repeated Negative Serological Testing in Otherwise Healthy Patients With Coronavirus Disease 2019. Journal of Infectious Diseases, 2021, 223, 924-926.	1.9	3
32	Pathogen Burden and Hypertension: More Questions Than Answers. American Journal of Hypertension, 2013, 26, 1459-1459.	1.0	2
33	Whole-genome sequencing of a CTX-M-11-encoding and quinolone-non-susceptible Klebsiella pneumoniae ST194 isolate from a hospitalised dog in Greece. Journal of Global Antimicrobial Resistance, 2018, 14, 126-128.	0.9	1
34	Prevalence and correlates of persistent intracellular HIV transcription in individuals on efavirenz versus atazanavir-based regimens: A prospective cohort study. PLoS ONE, 2018, 13, e0194262.	1.1	1
35	Humoral and T Cell Immune Responses to Sars-Cov-2 Vaccination in Hematopoietic Cell Transplant Recipients. Blood, 2021, 138, 2896-2896.	0.6	1
36	MP254VDR GENE EXPRESSION AND GLOBAL DNA METHYLATION ARE ASSOCIATED WITH INSULIN RESISTANCE IN ESRD PATIENTS ON CHRONIC HAEMODIALYSIS. Nephrology Dialysis Transplantation, 2016, 31, i424-i425.	0.4	0

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37	Vascular access malfunction: towards a more genecentric view. Clinical Nephrology, 2013, 80, 370-376.	0.4	0
38	Thrombotic Microangiopathy Variants Are Independently Associated with Critical Disease in COVID-19 Patients. Blood, 2020, 136, 21-22.	0.6	0