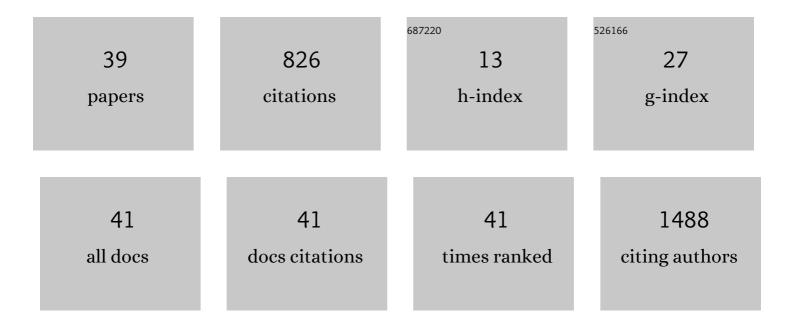
## Georgij Arapidi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Apoptotic Cell-Derived Extracellular Vesicles Promote Malignancy of Glioblastoma Via Intercellular Transfer of Splicing Factors. Cancer Cell, 2018, 34, 119-135.e10.	7.7	222
2	Proteome–Metabolome Profiling of Ovarian Cancer Ascites Reveals Novel Components Involved in Intercellular Communication. Molecular and Cellular Proteomics, 2014, 13, 3558-3571.	2.5	100
3	Quantitative proteomic analysis of Vietnamese krait venoms: Neurotoxins are the major components in Bungarus multicinctus and phospholipases A2 in Bungarus fasciatus. Toxicon, 2015, 107, 197-209.	0.8	55
4	Distinct types of short open reading frames are translated in plant cells. Genome Research, 2019, 29, 1464-1477.	2.4	43
5	Specific pools of endogenous peptides are present in gametophore, protonema, and protoplast cells of the moss Physcomitrella patens. BMC Plant Biology, 2015, 15, 87.	1.6	40
6	Therapy-induced stress response is associated with downregulation of pre-mRNA splicing in cancer cells. Genome Medicine, 2018, 10, 49.	3.6	40
7	The Pathogenesis of the Demyelinating Form of Guillain-Barre Syndrome (GBS): Proteo-peptidomic and Immunological Profiling of Physiological Fluids. Molecular and Cellular Proteomics, 2016, 15, 2366-2378.	2.5	39
8	Benchmarking germline CNV calling tools from exome sequencing data. Scientific Reports, 2021, 11, 14416.	1.6	36
9	Exposure to the Epstein–Barr Viral Antigen Latent Membrane Protein 1 Induces Myelin-Reactive Antibodies In Vivo. Frontiers in Immunology, 2017, 8, 777.	2.2	22
10	New method for peptide desorption from abundant blood proteins for plasma/serum peptidome analyses by mass spectrometry. Journal of Proteomics, 2011, 74, 595-606.	1.2	20
11	Alternative splicing shapes transcriptome but not proteome diversity in Physcomitrella patens. Scientific Reports, 2017, 7, 2698.	1.6	17
12	The Physcomitrella patens Chloroplast Proteome Changes in Response to Protoplastation. Frontiers in Plant Science, 2016, 7, 1661.	1.7	16
13	Critical Review of Existing MHC I Immunopeptidome Isolation Methods. Molecules, 2020, 25, 5409.	1.7	15
14	Peptidomics dataset: Blood plasma and serum samples of healthy donors fractionated on a set of chromatography sorbents. Data in Brief, 2018, 18, 1204-1211.	0.5	14
15	Identification of Antimicrobial Peptides from Novel Lactobacillus fermentum Strain. Protein Journal, 2020, 39, 73-84.	0.7	13
16	Expression and Intracellular Localization of Paraoxonase 2 in Different Types of Malignancies. Acta Naturae, 2018, 10, 92-99.	1.7	12
17	In Silico Analysis of Peptide Potential Biological Functions. Russian Journal of Bioorganic Chemistry, 2018, 44, 367-385.	0.3	11
18	Proteogenomic analysis of Mycobacterium tuberculosis Beijing B0/W148 cluster strains. Journal of Proteomics, 2019, 192, 18-26.	1.2	11

GEORGIJ ARAPIDI

#	Article	IF	CITATIONS
19	Metabolic Changes of Mycobacterium tuberculosis during the Anti-Tuberculosis Therapy. Pathogens, 2020, 9, 131.	1.2	11
20	Comprehensive Atlas of the Myelin Basic Protein Interaction Landscape. Biomolecules, 2021, 11, 1628.	1.8	11
21	Autoimmune Effect of Antibodies against the SARS-CoV-2 Nucleoprotein. Viruses, 2022, 14, 1141.	1.5	10
22	Progress in Methods for Copy Number Variation Profiling. International Journal of Molecular Sciences, 2022, 23, 2143.	1.8	9
23	Comprehensive analysis of draft genomes of two closely related pseudomonas syringae phylogroup 2b strains infecting mono- and dicotyledon host plants. BMC Genomics, 2016, 17, 1010.	1.2	8
24	Antimicrobial activity of endogenous peptides of the moss Physcomitrella patens. Russian Journal of Bioorganic Chemistry, 2017, 43, 248-254.	0.3	8
25	Peptidome profiling dataset of ovarian cancer and non-cancer proximal fluids: Ascites and blood sera. Data in Brief, 2019, 22, 557-562.	0.5	8
26	System OMICs analysis of Mycobacterium tuberculosis Beijing B0/W148 cluster. Scientific Reports, 2019, 9, 19255.	1.6	7
27	LogLoss-BERAF: An ensemble-based machine learning model for constructing highly accurate diagnostic sets of methylation sites accounting for heterogeneity in prostate cancer. PLoS ONE, 2018, 13, e0204371.	1.1	6
28	Methylation profile of induced pluripotent stem cells generated by integration and integration-free approaches. Data in Brief, 2018, 17, 662-666.	0.5	5
29	Serum proteome profiling for diagnostics of ovarian cancer using ClinProt magnetic technique and MALDI-TOF mass spectrometry. Biochemistry (Moscow) Supplement Series B: Biomedical Chemistry, 2008, 2, 335-342.	0.2	4
30	Chromatin Trapping of Factors Involved in DNA Replication and Repair Underlies Heat-Induced Radio- and Chemosensitization. Cells, 2020, 9, 1423.	1.8	3
31	Multiomic Profiling Identified EGF Receptor Signaling as a Potential Inhibitor of Type I Interferon Response in Models of Oncolytic Therapy by Vesicular Stomatitis Virus. International Journal of Molecular Sciences, 2022, 23, 5244.	1.8	3
32	Scope and limitations of MALDI-TOF MS blood serum peptide profiling in cancer diagnostics. Russian Journal of Bioorganic Chemistry, 2016, 42, 497-505.	0.3	2
33	Proteogenomic Approach for Mycobacterium tuberculosis Investigation. Methods in Molecular Biology, 2021, 2259, 191-201.	0.4	2
34	The Role of Intercellular Communication in Cancer Progression. Russian Journal of Bioorganic Chemistry, 2018, 44, 473-480.	0.3	1
35	The Diverse Roles of Spliceosomal Proteins in the Regulation of Cell Processes. Russian Journal of Bioorganic Chemistry, 2019, 45, 1-8.	0.3	1
36	A Role of Vesicular Transduction of Intercellular Signals in Cancer Development. Russian Journal of Bioorganic Chemistry, 2018, 44, 129-139.	0.3	0

#	Article	IF	CITATIONS
37	Spliceosomal componentsâ€mediated intercellular communication: role in chemoresistance acquisition of ovarian cancer cells. FASEB Journal, 2021, 35, .	0.2	0
38	Identification and analysis of exogenous peptides in human blood serum and plasma: Search for potential agents of interaction between the intestinal microbiota and the human body. FASEB Journal, 2021, 35, .	0.2	0
39	Substitutions in SurA and BamA Lead to Reduced Susceptibility to Broad Range Antibiotics in Gonococci. Genes, 2021, 12, 1312.	1.0	Ο