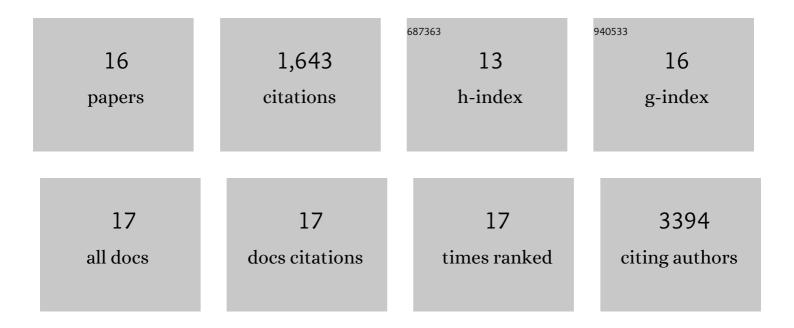
## Ivan Borozan

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Molecular and Pathology Features of Colorectal Tumors and Patient Outcomes Are Associated with <i>Fusobacterium nucleatum</i> and Its Subspecies <i>animalis</i> . Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 210-220.	2.5	19
2	Landscape of somatic single nucleotide variants and indels in colorectal cancer and impact on survival. Nature Communications, 2020, 11, 3644.	12.8	55
3	The landscape of viral associations in human cancers. Nature Genetics, 2020, 52, 320-330.	21.4	261
4	Epstein–Barr virus BORF2 inhibits cellular APOBEC3B to preserve viral genome integrity. Nature Microbiology, 2019, 4, 78-88.	13.3	95
5	Analysis of Epstein-Barr Virus Genomes and Expression Profiles in Gastric Adenocarcinoma. Journal of Virology, 2018, 92, .	3.4	47
6	Association of Distinct Mutational Signatures With Correlates of Increased Immune Activity in Pancreatic Ductal Adenocarcinoma. JAMA Oncology, 2017, 3, 774.	7.1	221
7	CSSSCL: a python package that uses combined sequence similarity scores for accurate taxonomic classification of long and short sequence reads. Bioinformatics, 2016, 32, 453-455.	4.1	4
8	Evaluation of Alignment Algorithms for Discovery and Identification of Pathogens Using RNA-Seq. PLoS ONE, 2013, 8, e76935.	2.5	37
9	CaPSID: A bioinformatics platform for computational pathogen sequence identification in human genomes and transcriptomes. BMC Bioinformatics, 2012, 13, 206.	2.6	41
10	Cell-Type Specific Gene Expression Signature in Liver Underlies Response to Interferon Therapy in Chronic Hepatitis C Infection. Gastroenterology, 2010, 138, 1123-1133.e3.	1.3	99
11	Gene expression profiling of early primary biliary cirrhosis: possible insights into the mechanism of action of ursodeoxycholic acid. Liver International, 2008, 28, 997-1010.	3.9	12
12	MAID : An effect size based model for microarray data integration across laboratories and platforms. BMC Bioinformatics, 2008, 9, 305.	2.6	12
13	Hepatic gene expression and prediction of therapy response in chronic hepatitis C patients. Journal of Hepatology, 2008, 48, 708-713.	3.7	20
14	Genome-wide Analysis of Substrate Specificities of the Escherichia coli Haloacid Dehalogenase-like Phosphatase Family. Journal of Biological Chemistry, 2006, 281, 36149-36161.	3.4	249
15	Hepatic Gene Expression Discriminates Responders and Nonresponders in Treatment of Chronic Hepatitis C Viral Infection. Gastroenterology, 2005, 128, 1437-1444.	1.3	433
16	Thermalization times of positrons in molecular gases. Journal of Physics B: Atomic, Molecular and Optical Physics, 2000, 33, 2725-2732.	1.5	31