

Evangelos Bellos

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

9 papers	263 citations	7 h-index	10 g-index
10 ext. papers	315 ext. citations	11.1 avg, IF	1.98 L-index

#	Paper	IF	Citations
9	Identification of novel locus associated with coronary artery aneurysms and validation of loci for susceptibility to Kawasaki disease. <i>European Journal of Human Genetics</i> , 2021 , 29, 1734-1744	5.3	2
8	A Rare Mutation in SPLUNC1 Affects Bacterial Adherence and Invasion in Meningococcal Disease. <i>Clinical Infectious Diseases</i> , 2020 , 70, 2045-2053	11.6	4
7	Mycobacterium tuberculosis Exploits a Molecular Off Switch of the Immune System for Intracellular Survival. <i>Scientific Reports</i> , 2018 , 8, 661	4.9	25
6	Natural resistance to Meningococcal Disease related to CFH loci: Meta-analysis of genome-wide association studies. <i>Scientific Reports</i> , 2016 , 6, 35842	4.9	26
5	Point Mutations in Exon 1B of APC Reveal Gastric Adenocarcinoma and Proximal Polyposis of the Stomach as a Familial Adenomatous Polyposis Variant. <i>American Journal of Human Genetics</i> , 2016 , 98, 830-842	11	153
4	cnvOffSeq: detecting intergenic copy number variation using off-target exome sequencing data. <i>Bioinformatics</i> , 2014 , 30, i639-45	7.2	9
3	cnvCapSeq: detecting copy number variation in long-range targeted resequencing data. <i>Nucleic Acids Research</i> , 2014 , 42, e158	20.1	8
2	A population model for genotyping indels from next-generation sequence data. <i>Nucleic Acids Research</i> , 2013 , 41, e46	20.1	12
1	cnvHiTSeq: integrative models for high-resolution copy number variation detection and genotyping using population sequencing data. <i>Genome Biology</i> , 2012 , 13, R120	18.3	24