

# Daniele Braga

## List of Publications by Year in descending order

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Version: 2024-02-01

17  
papers

937  
citations

687220

13  
h-index

887953

17  
g-index

17  
all docs

17  
docs citations

17  
times ranked

2780  
citing authors

#	ARTICLE	IF	CITATIONS
1	Gut vascular barrier impairment leads to intestinal bacteria dissemination and colorectal cancer metastasis to liver. <i>Cancer Cell</i> , 2021, 39, 708-724.e11.	7.7	175
2	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. <i>Nature Communications</i> , 2017, 8, 14977.	5.8	169
3	Genomewide Association Study Using a High-Density Single Nucleotide Polymorphism Array and Case-Control Design Identifies a Novel Essential Hypertension Susceptibility Locus in the Promoter Region of Endothelial NO Synthase. <i>Hypertension</i> , 2012, 59, 248-255.	1.3	144
4	Identification of a choroid plexus vascular barrier closing during intestinal inflammation. <i>Science</i> , 2021, 374, 439-448.	6.0	115
5	Self-renewal and phenotypic conversion are the main physiological responses of macrophages to the endogenous estrogen surge. <i>Scientific Reports</i> , 2017, 7, 44270.	1.6	58
6	Target Sequencing, Cell Experiments, and a Population Study Establish Endothelial Nitric Oxide Synthase ( <i>eNOS</i> ) Gene as Hypertension Susceptibility Gene. <i>Hypertension</i> , 2013, 62, 844-852.	1.3	48
7	Analysis of immune, microbiota and metabolome maturation in infants in a clinical trial of <i>Lactobacillus paracasei</i> CBAÅL74-fermented formula. <i>Nature Communications</i> , 2020, 11, 2703.	5.8	45
8	Interaction between polyphenols intake and PON1 gene variants on markers of cardiovascular disease: a nutrigenetic observational study. <i>Journal of Translational Medicine</i> , 2016, 14, 186.	1.8	38
9	TET2 and CSMD1 genes affect SBP response to hydrochlorothiazide in never-treated essential hypertensives. <i>Journal of Hypertension</i> , 2015, 33, 1301-1309.	0.3	29
10	Genome-wide association study identifies CAMKID variants involved in blood pressure response to losartan: the SOPHIA study. <i>Pharmacogenomics</i> , 2014, 15, 1643-1652.	0.6	27
11	Exome sequencing identifies variants in two genes encoding the LIM-proteins NRAP and FHL1 in an Italian patient with BAG3 myofibrillar myopathy. <i>Journal of Muscle Research and Cell Motility</i> , 2016, 37, 101-115.	0.9	23
12	A longitudinal study highlights shared aspects of the transcriptomic response to cardiogenic and septic shock. <i>Critical Care</i> , 2019, 23, 414.	2.5	20
13	Transcriptome Analysis of iPSC-Derived Neurons from Rubinstein-Taybi Patients Reveals Deficits in Neuronal Differentiation. <i>Molecular Neurobiology</i> , 2020, 57, 3685-3701.	1.9	15
14	Identification of a transcriptome profile associated with improvement of organ function in septic shock patients after early supportive therapy. <i>Critical Care</i> , 2018, 22, 312.	2.5	14
15	The burden of multiple sclerosis variants in continental Italians and Sardinians. <i>Multiple Sclerosis Journal</i> , 2015, 21, 1385-1395.	1.4	10
16	Left ventricular diastolic function associated with common genetic variation in ATP12A in a general population. <i>BMC Medical Genetics</i> , 2014, 15, 121.	2.1	4
17	Application of an Exploratory Knowledge-Discovery Pipeline Based on Machine Learning to Multi-Scale OMICS Data to Characterise Myocardial Injury in a Cohort of Patients with Septic Shock: An Observational Study. <i>Journal of Clinical Medicine</i> , 2021, 10, 4354.	1.0	3