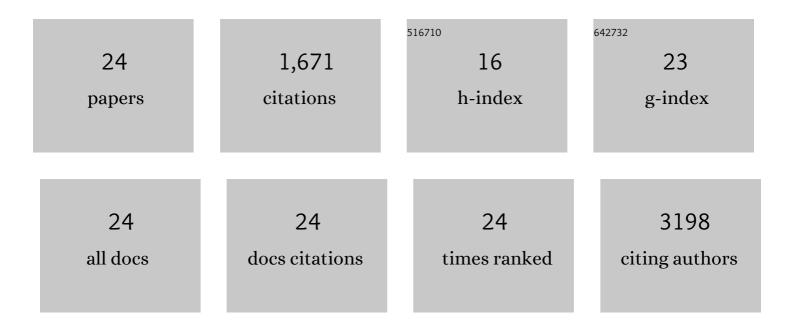
Roberto Cusano

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/12142051/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Gut Microbiota Markers and Dietary Habits Associated with Extreme Longevity in Healthy Sardinian Centenarians. Nutrients, 2022, 14, 2436.	4.1	18
2	Gut microbiota and metabolome distinctive features in Parkinson disease: Focus on levodopa and levodopa arbidopa intrajejunal gel. European Journal of Neurology, 2021, 28, 1198-1209.	3.3	20
3	Clinical Phenotypes of Parkinson's Disease Associate with Distinct Gut Microbiota and Metabolome Enterotypes. Biomolecules, 2021, 11, 144.	4.0	33
4	Gut microbiota markers associated with obesity and overweight in Italian adults. Scientific Reports, 2021, 11, 5532.	3.3	169
5	Genetic variants of TAS2R38 bitter taste receptor associate with distinct gut microbiota traits in Parkinson's disease: A pilot study. International Journal of Biological Macromolecules, 2020, 165, 665-674.	7.5	23
6	Gut Microbiota and Metabolome Alterations Associated with Parkinson's Disease. MSystems, 2020, 5, .	3.8	161
7	Impact of a Moderately Hypocaloric Mediterranean Diet on the Gut Microbiota Composition of Italian Obese Patients. Nutrients, 2020, 12, 2707.	4.1	33
8	Population- and individual-specific regulatory variation in Sardinia. Nature Genetics, 2017, 49, 700-707.	21.4	38
9	Overexpression of the Cytokine BAFF and Autoimmunity Risk. New England Journal of Medicine, 2017, 376, 1615-1626.	27.0	301
10	Deep genomic analysis of the Chlorella sorokiniana SAG 211-8k chloroplast. European Journal of Phycology, 2017, 52, 320-329.	2.0	3
11	Complete genome sequence of mitochondrial DNA (mtDNA) ofChlorella sorokiniana. Mitochondrial DNA, 2016, 27, 1539-1541.	0.6	5
12	Complete genome sequence of chloroplast DNA (cpDNA) of <i>Chlorella sorokiniana</i> . Mitochondrial DNA, 2016, 27, 838-839.	0.6	12
13	Complete sequence and characterization of mitochondrial and chloroplast genome of <i>Chlorella variabilis</i> NC64A. Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 3128-3130.	0.7	7
14	<i>ALDH18A1</i> gene mutations cause dominant spastic paraplegia SPG9: loss of function effect and plausibility of a dominant negative mechanism. Brain, 2016, 139, e3-e3.	7.6	42
15	Genetic Variants Regulating Immune Cell Levels in Health and Disease. Cell, 2013, 155, 242-256.	28.9	295
16	Genetic heterogeneity in inherited spastic paraplegia associated with epilepsy. , 2003, 117A, 116-121.		7
17	Title is missing!. Medicine (United States), 2003, 82, 203-215.	1.0	30
18	MYH9-Related Disease. Medicine (United States), 2003, 82, 203-215.	1.0	255

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#	Article	IF	CITATIONS
19	Epstein syndrome: another renal disorder with mutations in the nonmuscle myosin heavy chainÂ9 gene. Human Genetics, 2002, 110, 182-186.	3.8	45
20	Linkage Analysis in Families with Recurrent Neuroblastoma. Annals of the New York Academy of Sciences, 2002, 963, 74-84.	3.8	17
21	A refined physical and transcriptional map of the SPG9 locus on 10q23.3–q24.2. European Journal of Human Genetics, 2000, 8, 777-782.	2.8	17
22	Neuroblastoma in Two Siblings Supports the Role of 1p36 Deletion in Tumor Development. Cancer Genetics and Cytogenetics, 1999, 109, 126-130.	1.0	13
23	Genetic Mapping to 10q23.3-q24.2, in a Large Italian Pedigree, of a New Syndrome Showing Bilateral Cataracts, Gastroesophageal Reflux, and Spastic Paraparesis with Amyotrophy. American Journal of Human Genetics, 1999, 64, 586-593.	6.2	108
24	Interstitial and large chromosome 1p deletion occurs in localized and disseminated neuroblastomas and predicts an unfavourable outcome. Cancer Letters, 1998, 130, 83-92.	7.2	19