Mireia Valles-Colomer

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3347788/publications.pdf

Version: 2024-02-01

23 papers 6,629 citations

361296 20 h-index 677027 22 g-index

26 all docs

26 does citations

26 times ranked 9747 citing authors

#	Article	lF	CITATIONS
1	Exploring the relationship between the gut microbiome and mental health outcomes in a posttraumatic stress disorder cohort relative to trauma-exposed controls. European Neuropsychopharmacology, 2022, 56, 24-38.	0.3	26
2	Variation and transmission of the human gut microbiota across multiple familial generations. Nature Microbiology, 2022, 7, 87-96.	5.9	32
3	Metagenomic and metabolomic remodeling in nonagenarians and centenarians and its association with genetic and socioeconomic factors. Nature Aging, 2022, 2, 438-452.	5.3	17
4	Integrating taxonomic, functional, and strain-level profiling of diverse microbial communities with bioBakery 3. ELife, $2021,10,10$	2.8	808
5	Genomic diversity and ecology of human-associated Akkermansia species in the gut microbiome revealed by extensive metagenomic assembly. Genome Biology, 2021, 22, 209.	3.8	65
6	Microbiome connections with host metabolism and habitual diet from 1,098 deeply phenotyped individuals. Nature Medicine, 2021, 27, 321-332.	15.2	477
7	Treponema peruense sp. nov., a commensal spirochaete isolated from human faeces. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	0.8	10
8	Combinatorial, additive and dose-dependent drug–microbiome associations. Nature, 2021, 600, 500-505.	13.7	102
9	Successional Stages in Infant Gut Microbiota Maturation. MBio, 2021, 12, e0185721.	1.8	48
10	Depression and suicidality: A link to premature T helper cell aging and increased Th17 cells. Brain, Behavior, and Immunity, 2020, 87, 603-609.	2.0	57
11	Statin therapy is associated with lower prevalence of gut microbiota dysbiosis. Nature, 2020, 581, 310-315.	13.7	283
12	Analysis of 1321 Eubacterium rectale genomes from metagenomes uncovers complex phylogeographic population structure and subspecies functional adaptations. Genome Biology, 2020, 21, 138.	3.8	72
13	Gut microbiome variation is associated to Multiple Sclerosis phenotypic subtypes. Annals of Clinical and Translational Neurology, 2020, 7, 406-419.	1.7	59
14	Quantitative microbiome profiling disentangles inflammation- and bile duct obstruction-associated microbiota alterations across PSC/IBD diagnoses. Nature Microbiology, 2019, 4, 1826-1831.	5.9	149
15	The neuroactive potential of the human gut microbiota in quality of life and depression. Nature Microbiology, 2019, 4, 623-632.	5.9	1,206
16	Practical guidelines for gut microbiome analysis in microbiota-gut-brain axis research. Behavioral and Brain Sciences, 2019, 42, .	0.4	1
17	The Gut Microbiome and Mental Health: Implications for Anxiety- and Trauma-Related Disorders. OMICS A Journal of Integrative Biology, 2018, 22, 90-107.	1.0	110
18	Butyrate Producers as Potential Next-Generation Probiotics: Safety Assessment of the Administration of <i>Butyricicoccus pullicaecorum</i> to Healthy Volunteers. MSystems, 2018, 3, .	1.7	99

#	Article	IF	CITATIONS
19	A low-gluten diet induces changes in the intestinal microbiome of healthy Danish adults. Nature Communications, 2018, 9, 4630.	5.8	124
20	Quantitative microbiome profiling links gut community variation to microbial load. Nature, 2017, 551, 507-511.	13.7	791
21	Population-level analysis of gut microbiome variation. Science, 2016, 352, 560-564.	6.0	1,716
22	Species–function relationships shape ecological properties of the human gut microbiome. Nature Microbiology, 2016, 1, 16088.	5.9	279
23	Meta-omics in Inflammatory Bowel Disease Research: Applications, Challenges, and Guidelines. Journal of Crohn's and Colitis, 2016, 10, 735-746.	0.6	37