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	IF	CITATIONS
1	Population-level analysis of gut microbiome variation. Science, 2016, 352, 560-564.	6.0	1,716
2	The neuroactive potential of the human gut microbiota in quality of life and depression. Nature Microbiology, 2019, 4, 623-632.	5.9	1,206
3	Integrating taxonomic, functional, and strain-level profiling of diverse microbial communities with bioBakery 3. ELife, 2021, 10, .	2.8	808
4	Quantitative microbiome profiling links gut community variation to microbial load. Nature, 2017, 551, 507-511.	13.7	791
5	Microbiome connections with host metabolism and habitual diet from 1,098 deeply phenotyped individuals. Nature Medicine, 2021, 27, 321-332.	15.2	477
6	Statin therapy is associated with lower prevalence of gut microbiota dysbiosis. Nature, 2020, 581, 310-315.	13.7	283
7	Species–function relationships shape ecological properties of the human gut microbiome. Nature Microbiology, 2016, 1, 16088.	5.9	279
8	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
9	A low-gluten diet induces changes in the intestinal microbiome of healthy Danish adults. Nature Communications, 2018, 9, 4630.	5.8	124
10	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
11	Combinatorial, additive and dose-dependent drug–microbiome associations. Nature, 2021, 600, 500-505.	13.7	102
12	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
13	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
14	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
15	Gut microbiome variation is associated to Multiple Sclerosis phenotypic subtypes. Annals of Clinical and Translational Neurology, 2020, 7, 406-419.	1.7	59
16	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
17	Successional Stages in Infant Gut Microbiota Maturation. MBio, 2021, 12, e0185721.	1.8	48
18	Meta-omics in Inflammatory Bowel Disease Research: Applications, Challenges, and Guidelines. Journal of Crohn's and Colitis, 2016, 10, 735-746.	0.6	37

#	Article	IF	CITATIONS
19	Variation and transmission of the human gut microbiota across multiple familial generations. Nature Microbiology, 2022, 7, 87-96.	5.9	32
20	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
21	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
22	Treponema peruense sp. nov., a commensal spirochaete isolated from human faeces. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	0.8	10
23	Practical guidelines for gut microbiome analysis in microbiota-gut-brain axis research. Behavioral and Brain Sciences, 2019, 42, .	0.4	1