Christel Chehoud

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10769351/publications.pdf

Version: 2024-02-01

12 papers

2,912 citations

759055 12 h-index 1199470 12 g-index

12 all docs 12 docs citations

times ranked

12

5571 citing authors

#	Article	IF	CITATIONS
1	Inflammation, Antibiotics, and Diet as Environmental Stressors of the Gut Microbiome in Pediatric Crohn's Disease. Cell Host and Microbe, 2015, 18, 489-500.	5.1	646
2	Rapid evolution of the human gut virome. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 12450-12455.	3.3	489
3	Comparative metabolomics in vegans and omnivores reveal constraints on diet-dependent gut microbiota metabolite production. Gut, 2016, 65, 63-72.	6.1	428
4	Optimizing methods and dodging pitfalls in microbiome research. Microbiome, 2017, 5, 52.	4.9	420
5	Fungal Signature in the Gut Microbiota of Pediatric Patients With Inflammatory Bowel Disease. Inflammatory Bowel Diseases, 2015, 21, 1948-1956.	0.9	194
6	Engineering the gut microbiota to treat hyperammonemia. Journal of Clinical Investigation, 2015, 125, 2841-2850.	3.9	154
7	Prevalence of Apical Periodontitis in Endodontically Treated Premolars and Molars with Untreated Canal: A Cone-beam Computed Tomography Study. Journal of Endodontics, 2016, 42, 538-541.	1.4	152
8	Comparative Effectiveness of Nutritional and Biological Therapy in North American Children with Active Crohn $\hat{E}\frac{1}{4}$ s Disease. Inflammatory Bowel Diseases, 2015, 21, 1786-1793.	0.9	141
9	Complement modulates the cutaneous microbiome and inflammatory milieu. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 15061-15066.	3.3	138
10	Transfer of Viral Communities between Human Individuals during Fecal Microbiota Transplantation. MBio, 2016, 7, e00322.	1.8	90
11	Associations of the vaginal microbiota with HIV infection, bacterial vaginosis, and demographic factors. Aids, 2017, 31, 895-904.	1.0	44
12	Dietary Regulation of the Gut Microbiota Engineered by a Minimal Defined Bacterial Consortium. PLoS ONE, 2016, 11, e0155620.	1.1	16