

Joel Dore

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6585560/publications.pdf>

Version: 2024-02-01

21
papers

28,845
citations

393982

19
h-index

676716

22
g-index

24
all docs

24
docs citations

24
times ranked

30416
citing authors

#	ARTICLE	IF	CITATIONS
1	A human gut microbial gene catalogue established by metagenomic sequencing. <i>Nature</i> , 2010, 464, 59-65.	13.7	9,342
2	Enterotypes of the human gut microbiome. <i>Nature</i> , 2011, 473, 174-180.	13.7	5,800
3	Richness of human gut microbiome correlates with metabolic markers. <i>Nature</i> , 2013, 500, 541-546.	13.7	3,641
4	<i>Faecalibacterium prausnitzii</i> is an anti-inflammatory commensal bacterium identified by gut microbiota analysis of Crohn disease patients. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 16731-16736.	3.3	3,581
5	An integrated catalog of reference genes in the human gut microbiome. <i>Nature Biotechnology</i> , 2014, 32, 834-841.	9.4	1,664
6	Dietary intervention impact on gut microbial gene richness. <i>Nature</i> , 2013, 500, 585-588.	13.7	1,485
7	Identification and assembly of genomes and genetic elements in complex metagenomic samples without using reference genomes. <i>Nature Biotechnology</i> , 2014, 32, 822-828.	9.4	909
8	Twin Study Indicates Loss of Interaction Between Microbiota and Mucosa of Patients With Ulcerative Colitis. <i>Gastroenterology</i> , 2011, 141, 227-236.	0.6	518
9	Identification of an Intestinal Microbiota Signature Associated With Severity of Irritable Bowel Syndrome. <i>Gastroenterology</i> , 2017, 152, 111-123.e8.	0.6	470
10	Gut microbiota after gastric bypass in human obesity: increased richness and associations of bacterial genera with adipose tissue genes. <i>American Journal of Clinical Nutrition</i> , 2013, 98, 16-24.	2.2	351
11	A metagenomic insight into our gut's microbiome. <i>Gut</i> , 2013, 62, 146-158.	6.1	302
12	Microbial ecology perturbation in human IgA deficiency. <i>Science Translational Medicine</i> , 2018, 10, .	5.8	206
13	The influence of diet on the gut microbiota and its consequences for health. <i>Current Opinion in Biotechnology</i> , 2015, 32, 195-199.	3.3	148
14	Dietary Patterns Differently Associate with Inflammation and Gut Microbiota in Overweight and Obese Subjects. <i>PLoS ONE</i> , 2014, 9, e109434.	1.1	111
15	Bacterial protein signals are associated with Crohn's disease. <i>Gut</i> , 2014, 63, 1566-1577.	6.1	80
16	Human intestinal metagenomics: state of the art and future. <i>Current Opinion in Microbiology</i> , 2013, 16, 232-239.	2.3	62
17	Altered host-gut microbes symbiosis in severely malnourished anorexia nervosa (AN) patients undergoing enteral nutrition: An explicative factor of functional intestinal disorders?. <i>Clinical Nutrition</i> , 2019, 38, 2304-2310.	2.3	62
18	Anorexia nervosa and gut microbiota: A systematic review and quantitative synthesis of pooled microbiological data. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2021, 106, 110114.	2.5	49

#	ARTICLE	IF	CITATIONS
19	The human gut microbiome as source of innovation for health: Which physiological and therapeutic outcomes could we expect?. <i>Therapie</i> , 2017, 72, 21-38.	0.6	28
20	Quality control of microbiota metagenomics by k-mer analysis. <i>BMC Genomics</i> , 2015, 16, 183.	1.2	22
21	Roseburia, a decreased bacterial taxon in the gut microbiota of patients suffering from anorexia nervosa. <i>European Journal of Clinical Nutrition</i> , 2022, , .	1.3	6