Tal Korem

List of Publications by Year in descending order

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Version: 2024-02-01

430442 552369 10,585 26 18 26 h-index citations g-index papers 30 30 30 15304 docs citations times ranked citing authors all docs

#	Article	lF	Citations
1	Prediction of Personal Glycemic Responses to Food for Individuals With Type 1 Diabetes Through Integration of Clinical and Microbial Data. Diabetes Care, 2022, 45, 502-511.	4.3	15
2	The Gut Microbiome of Adults With Type 1 Diabetes and Its Association With the Host Glycemic Control. Diabetes Care, 2022, 45, 555-563.	4.3	19
3	Accurate and robust inference of microbial growth dynamics from metagenomic sequencing reveals personalized growth rates. Genome Research, 2022, 32, 558-568.	2.4	23
4	Deciphering metabolism, one microbe at a time. Nature, 2021, 595, 355-357.	13.7	0
5	Embracing Metagenomic Complexity with a Genome-Free Approach. MSystems, 2021, 6, e0081621.	1.7	9
6	Minimal Associations between Short-Term Dietary Intake and Salivary Microbiome Composition. Microorganisms, 2021, 9, 1739.	1.6	2
7	The hygiene hypothesis, the COVID pandemic, and consequences for the human microbiome. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	100
8	A non-optimal cervicovaginal microbiota in pregnancy is associated with a distinct metabolomic signature among non-Hispanic Black individuals. Scientific Reports, 2021, 11, 22794.	1.6	8
9	Longitudinal Multi-omics Reveals Subset-Specific Mechanisms Underlying Irritable Bowel Syndrome. Cell, 2020, 182, 1460-1473.e17.	13.5	217
10	A reference map of potential determinants for the human serum metabolome. Nature, 2020, 588, 135-140.	13.7	230
11	The Cancer Microbiome: Distinguishing Direct and Indirect Effects Requires a Systemic View. Trends in Cancer, 2020, 6, 192-204.	3.8	162
12	Relationship of the Esophageal Microbiome and Tissue Gene Expression and Links to the Oral Microbiome: A Randomized Clinical Trial. Clinical and Translational Gastroenterology, 2020, 11, e00235.	1.3	13
13	Structural variation in the gut microbiome associates with host health. Nature, 2019, 568, 43-48.	13.7	244
14	Environment dominates over host genetics in shaping human gut microbiota. Nature, 2018, 555, 210-215.	13.7	1,958
15	Post-Antibiotic Gut Mucosal Microbiome Reconstitution Is Impaired by Probiotics and Improved by Autologous FMT. Cell, 2018, 174, 1406-1423.e16.	13.5	752
16	Involvement of a gut–retina axis in protection against dietary glycemia-induced age-related macular degeneration. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E4472-E4481.	3.3	179
17	Bread Affects Clinical Parameters and Induces Gut Microbiome-Associated Personal Glycemic Responses. Cell Metabolism, 2017, 25, 1243-1253.e5.	7.2	233
18	Microbiota Diurnal Rhythmicity Programs Host Transcriptome Oscillations. Cell, 2016, 167, 1495-1510.e12.	13.5	591

#	Article	IF	CITATIONS
19	Taking it Personally: Personalized Utilization of the Human Microbiome in Health and Disease. Cell Host and Microbe, 2016, 19, 12-20.	5.1	192
20	Microbiota-Modulated Metabolites Shape the Intestinal Microenvironment by Regulating NLRP6 Inflammasome Signaling. Cell, 2015, 163, 1428-1443.	13.5	728
21	Growth dynamics of gut microbiota in health and disease inferred from single metagenomic samples. Science, 2015, 349, 1101-1106.	6.0	382
22	Non-caloric artificial sweeteners and the microbiome: findings and challenges. Gut Microbes, 2015, 6, 149-155.	4.3	152
23	Artificial Sweeteners Induce Glucose Intolerance by Altering the Gut Microbiota. Obstetrical and Gynecological Survey, 2015, 70, 31-32.	0.2	6
24	Personalized Nutrition by Prediction of Glycemic Responses. Cell, 2015, 163, 1079-1094.	13.5	1,816
25	Transkingdom Control of Microbiota Diurnal Oscillations Promotes Metabolic Homeostasis. Cell, 2014, 159, 514-529.	13.5	984
26	Artificial sweeteners induce glucose intolerance by altering the gut microbiota. Nature, 2014, 514, 181-186.	13.7	1,529