Shinji Fukuda

List of Publications by Year in descending order

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72 papers 12,195 citations

147566 31 h-index 63 g-index

84 all docs

84 docs citations

84 times ranked 16166 citing authors

#	Article	IF	Citations
1	Gut environment changes due to androgen deprivation therapy in patients with prostate cancer. Prostate Cancer and Prostatic Diseases, 2023, 26, 323-330.	2.0	9
2	Aspergillus-Derived Cellulase Preparation Exhibits Prebiotic-like Effects on Gut Microbiota in Rats. Fermentation, 2022, 8, 71.	1.4	4
3	Surgical Treatment for Colorectal Cancer Partially Restores Gut Microbiome and Metabolome Traits. MSystems, 2022, 7, e0001822.	1.7	3
4	Rap1 prevents colitogenic Th17 cell expansion and facilitates Treg cell differentiation and distal TCR signaling. Communications Biology, 2022, 5, 206.	2.0	5
5	Resistant Maltodextrin Intake Reduces Virulent Metabolites in the Gut Environment: A Randomized Control Study in a Japanese Cohort. Frontiers in Microbiology, 2022, 13, .	1.5	4
6	Gasdermin D–mediated release of IL-33 from senescent hepatic stellate cells promotes obesity-associated hepatocellular carcinoma. Science Immunology, 2022, 7, .	5.6	43
7	Effects of Psychotropics on the Microbiome in Patients With Depression and Anxiety: Considerations in a Naturalistic Clinical Setting. International Journal of Neuropsychopharmacology, 2021, 24, 97-107.	1.0	24
8	Positive Effects of Oral Antibiotic Administration in Murine Chronic Graft-Versus-Host Disease. International Journal of Molecular Sciences, 2021, 22, 3745.	1.8	8
9	Symbiotic polyamine metabolism regulates epithelial proliferation and macrophage differentiation in the colon. Nature Communications, 2021, 12, 2105.	5.8	105
10	Complete Genome Sequence of Adlercreutzia equolifaciens subsp. celatus DSM 18785. Microbiology Resource Announcements, 2021, 10, .	0.3	5
11	The Nutritional Efficacy of Chlorella Supplementation Depends on the Individual Gut Environment: A Randomised Control Study. Frontiers in Nutrition, 2021, 8, 648073.	1.6	7
12	Energy landscape analysis elucidates the multistability of ecological communities across environmental gradients. Ecological Monographs, 2021, 91, e01469.	2.4	15
13	Complete Genome Sequence of Atopobiaceae Bacterium Strain P1, Isolated from Mouse Feces. Microbiology Resource Announcements, 2021, 10, e0062721.	0.3	1
14	Fecal Microbial and Metabolomic Change during treatment course for depression: An Observational Study. Journal of Psychiatric Research, 2021, 140, 45-52.	1.5	10
15	Seaweed Dietary Fiber Sodium Alginate Suppresses the Migration of Colonic Inflammatory Monocytes and Diet-Induced Metabolic Syndrome via the Gut Microbiota. Nutrients, 2021, 13, 2812.	1.7	13
16	Metabolic Effects of Bee Larva-Derived Protein in Mice: Assessment of an Alternative Protein Source. Foods, 2021, 10, 2642.	1.9	0
17	Lactoferrin Ameliorates Dry Eye Disease Potentially through Enhancement of Short-Chain Fatty Acid Production by Gut Microbiota in Mice. International Journal of Molecular Sciences, 2021, 22, 12384.	1.8	8
18	Decoding gut microbiota by imaging analysis of fecal samples. IScience, 2021, 24, 103481.	1.9	2

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19	Supplemental Aspergillus Lipase and Protease Preparations Display Powerful Bifidogenic Effects and Modulate the Gut Microbiota Community of Rats. Fermentation, 2021, 7, 294.	1.4	6
20	The guanylate cyclase C agonist linaclotide ameliorates the gut–cardio–renal axis in an adenine-induced mouse model of chronic kidney disease. Nephrology Dialysis Transplantation, 2020, 35, 250-264.	0.4	35
21	Dietary l-serine confers a competitive fitness advantage to Enterobacteriaceae in the inflamed gut. Nature Microbiology, 2020, 5, 116-125.	5.9	93
22	Conjunctival Injection Reduction in Patients with Atopic Keratoconjunctivitis Due to Synergic Effect of Bovine Enteric-Coated Lactoferrin in 0.1% Tacrolimus Ophthalmic Suspension. Journal of Clinical Medicine, 2020, 9, 3093.	1.0	5
23	Asperuloside Improves Obesity and Type 2 Diabetes through Modulation of Gut Microbiota and Metabolic Signaling. IScience, 2020, 23, 101522.	1.9	21
24	Gut microbiota depletion by chronic antibiotic treatment alters the sleep/wake architecture and sleep EEG power spectra in mice. Scientific Reports, 2020, 10, 19554.	1.6	59
25	An Enteric Pathogen Subverts Colonization Resistance by Evading Competition for Amino Acids in the Gut. Cell Host and Microbe, 2020, 28, 526-533.e5.	5.1	29
26	Germ-Free Conditions Modulate Host Purine Metabolism, Exacerbating Adenine-Induced Kidney Damage. Toxins, 2020, 12, 547.	1.5	23
27	Paternal restraint stress affects offspring metabolism via ATF-2 dependent mechanisms in Drosophila melanogaster germ cells. Communications Biology, 2020, 3, 208.	2.0	16
28	Scoring Species for Synthetic Community Design: Network Analyses of Functional Core Microbiomes. Frontiers in Microbiology, 2020, 11, 1361.	1.5	26
29	Complete Genome Sequence of Bifidobacterium longum Strain Jih1, Isolated from Human Feces. Microbiology Resource Announcements, 2020, 9, .	0.3	1
30	Interleukin-22-mediated host glycosylation prevents Clostridioides difficile infection by modulating the metabolic activity of the gut microbiota. Nature Medicine, 2020, 26, 608-617.	15.2	136
31	Influence of gastrectomy for gastric cancer treatment on faecal microbiome and metabolome profiles. Gut, 2020, 69, 1404-1415.	6.1	84
32	Does the Gut Microbiota Modulate Host Physiology through Polymicrobial Biofilms?. Microbes and Environments, 2020, 35, n/a .	0.7	13
33	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice. , 2020, 15, e0228358.		0
34	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice. , 2020, 15, e0228358.		0
35	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice. , 2020, 15, e0228358.		0
36	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice. , 2020, 15, e0228358.		0

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37	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice., 2020, 15, e0228358.		О
38	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice. , 2020, 15, e0228358.		0
39	Protective effects of bifidobacteria against enteropathogens. Microbial Biotechnology, 2019, 12, 1097-1100.	2.0	17
40	Metagenomic and metabolomic analyses reveal distinct stage-specific phenotypes of the gut microbiota in colorectal cancer. Nature Medicine, 2019, 25, 968-976.	15.2	748
41	Gut microbiome-derived phenyl sulfate contributes to albuminuria in diabetic kidney disease. Nature Communications, 2019, 10, 1835.	5.8	173
42	Effects of bowel preparation on the human gut microbiome and metabolome. Scientific Reports, 2019, 9, 4042.	1.6	78
43	Cutibacterium acnes (Propionibacterium acnes) 16S rRNA Genotyping of Microbial Samples from Possessions Contributes to Owner Identification. MSystems, 2019, 4, .	1.7	33
44	Effects of the 1975 Japanese diet on the gut microbiota in younger adults. Journal of Nutritional Biochemistry, 2019, 64, 121-127.	1.9	27
45	Draft Genome Sequences of Enterococcus faecalis Strains Isolated from Healthy Japanese Individuals. Microbiology Resource Announcements, 2019, 8, .	0.3	2
46	Proton Pump Inhibitors Increase the Susceptibility of Mice to Oral Infection with Enteropathogenic Bacteria. Digestive Diseases and Sciences, 2018, 63, 881-889.	1.1	15
47	Understanding the role of the gut ecosystem in diabetes mellitus. Journal of Diabetes Investigation, 2018, 9, 5-12.	1.1	110
48	Canagliflozin reduces plasma uremic toxins and alters the intestinal microbiota composition in a chronic kidney disease mouse model. American Journal of Physiology - Renal Physiology, 2018, 315, F824-F833.	1.3	84
49	A Metabologenomic Approach Reveals Changes in the Intestinal Environment of Mice Fed on American Diet. International Journal of Molecular Sciences, 2018, 19, 4079.	1.8	41
50	A Metabolomic-Based Evaluation of the Role of Commensal Microbiota throughout the Gastrointestinal Tract in Mice. Microorganisms, 2018, 6, 101.	1.6	24
51	Urban Diets Linked to Gut Microbiome and Metabolome Alterations in Children: A Comparative Cross-Sectional Study in Thailand. Frontiers in Microbiology, 2018, 9, 1345.	1.5	55
52	Neonatal acquisition of <i>Clostridia</i> species protects against colonization by bacterial pathogens. Science, 2017, 356, 315-319.	6.0	199
53	Evaluation of the impact of gut microbiota on uremic solute accumulation by a CE-TOFMS–based metabolomics approach. Kidney International, 2017, 92, 634-645.	2.6	173
54	Functional Characterization of Inflammatory Bowel Disease–Associated Gut Dysbiosis in Gnotobiotic Mice. Cellular and Molecular Gastroenterology and Hepatology, 2016, 2, 468-481.	2.3	189

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55	Probiotic Bifidobacterium longum alters gut luminal metabolism through modification of the gut microbial community. Scientific Reports, 2015, 5, 13548.	1.6	126
56	An Integrated Outlook on the Metagenome and Metabolome of Intestinal Diseases. Diseases (Basel,) Tj ETQq0 C	O rgBT /O	verlock 10 Tf
57	The Consumption of Bicarbonate-Rich Mineral Water Improves Glycemic Control. Evidence-based Complementary and Alternative Medicine, 2015, 2015, 1-10.	0.5	42
58	The Role of Integrated Omics in Elucidating the Gut Microbiota Health Potentials. Microbiology Monographs, 2015, , 73-100.	0.3	2
59	Alteration of the Intestinal Environment by Lubiprostone Is Associated with Amelioration of Adenine-Induced CKD. Journal of the American Society of Nephrology: JASN, 2015, 26, 1787-1794.	3.0	162
60	Toward the comprehensive understanding of the gut ecosystem via metabolomics-based integrated omics approach. Seminars in Immunopathology, 2015, 37, 5-16.	2.8	46
61	<i>Lactobacillus acidophilus</i> L-92 Cells Activate Expression of Immunomodulatory Genes in THP-1 Cells. Bioscience of Microbiota, Food and Health, 2014, 33, 157-164.	0.8	6
62	A novel mucosal vaccine targeting Peyer's patch M cells induces protective antigen-specific IgA responses. International Immunology, 2014, 26, 619-625.	1.8	62
63	Multiple Omics Uncovers Host–Gut Microbial Mutualism During Prebiotic Fructooligosaccharide Supplementation. DNA Research, 2014, 21, 469-480.	1.5	101
64	Gut microbiome and metabolic diseases. Seminars in Immunopathology, 2014, 36, 103-114.	2.8	121
65	Treg induction by a rationally selected mixture of Clostridia strains from the human microbiota. Nature, 2013, 500, 232-236.	13.7	2,339
66	Commensal microbe-derived butyrate induces the differentiation of colonic regulatory T cells. Nature, 2013, 504, 446-450.	13.7	3,901
67	Microbiota-derived lactate accelerates colon epithelial cell turnover in starvation-refed mice. Nature Communications, 2013, 4, 1654.	5.8	111
68	Robustness of Gut Microbiota of Healthy Adults in Response to Probiotic Intervention Revealed by High-Throughput Pyrosequencing. DNA Research, 2013, 20, 241-253.	1.5	272
69	Acetate-producing bifidobacteria protect the host from enteropathogenic infection via carbohydrate transporters. Gut Microbes, 2012, 3, 449-454.	4.3	174
70	The Epithelia-Specific Membrane Trafficking Factor AP-1B Controls Gut Immune Homeostasis in Mice. Gastroenterology, 2011, 141, 621-632.	0.6	51
71	Bifidobacteria can protect from enteropathogenic infection through production of acetate. Nature, 2011, 469, 543-547.	13.7	1,836
72	A new strain of Butyrivibrio fibrisolvens that has high ability to isomerize linoleic acid to conjugated linoleic acid. Journal of General and Applied Microbiology, 2005, 51, 105-113.	0.4	41