Hirokazu Tsuji

List of Publications by Year in descending order

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ΗΙΡΟΚΑΖΗ ΤΟΙΗ

#	Article	IF	CITATIONS
1	Possible association of Bifidobacterium and Lactobacillus in the gut microbiota of patients with major depressive disorder. Journal of Affective Disorders, 2016, 202, 254-257.	2.0	419
2	Intestinal Dysbiosis and Lowered Serum Lipopolysaccharide-Binding Protein in Parkinson's Disease. PLoS ONE, 2015, 10, e0142164.	1.1	381
3	Probiotic Bifidobacterium breve Induces IL-10-Producing Tr1 Cells in the Colon. PLoS Pathogens, 2012, 8, e1002714.	2.1	277
4	Sensitive Quantitative Detection of Commensal Bacteria by rRNA-Targeted Reverse Transcription-PCR. Applied and Environmental Microbiology, 2007, 73, 32-39.	1.4	257
5	Establishment of an Analytical System for the Human Fecal Microbiota, Based on Reverse Transcription-Quantitative PCR Targeting of Multicopy rRNA Molecules. Applied and Environmental Microbiology, 2009, 75, 1961-1969.	1.4	237
6	Diversity in gut bacterial community of school-age children in Asia. Scientific Reports, 2015, 5, 8397.	1.6	221
7	Gut dysbiosis is associated with metabolism and systemic inflammation in patients with ischemic stroke. PLoS ONE, 2017, 12, e0171521.	1.1	205
8	Progression of Parkinson's disease is associated with gut dysbiosis: Two-year follow-up study. PLoS ONE, 2017, 12, e0187307.	1.1	195
9	Identification of phenol- and p-cresol-producing intestinal bacteria by using media supplemented with tyrosine and its metabolites. FEMS Microbiology Ecology, 2018, 94, .	1.3	182
10	Gut Dysbiosis in Patients with Anorexia Nervosa. PLoS ONE, 2015, 10, e0145274.	1.1	179
11	Key bacterial taxa and metabolic pathways affecting gut short-chain fatty acid profiles in early life. ISME Journal, 2021, 15, 2574-2590.	4.4	131
12	A Single Species of Clostridium Subcluster XIVa Decreased in Ulcerative Colitis Patients. Inflammatory Bowel Diseases, 2016, 22, 2802-2810.	0.9	126
13	Sensitive Quantitative Analysis of the Meconium Bacterial Microbiota in Healthy Term Infants Born Vaginally or by Cesarean Section. Frontiers in Microbiology, 2016, 7, 1997.	1.5	125
14	Ontogenesis of the Gut Microbiota Composition in Healthy, Full-Term, Vaginally Born and Breast-Fed Infants over the First 3 Years of Life: A Quantitative Bird's-Eye View. Frontiers in Microbiology, 2017, 8, 1388.	1.5	103
15	Molecular monitoring of the development of intestinal microbiota in Japanese infants. Beneficial Microbes, 2012, 3, 113-125.	1.0	81
16	Role of probiotic in preventing acute diarrhoea in children: a community-based, randomized, double-blind placebo-controlled field trial in an urban slum. Epidemiology and Infection, 2011, 139, 919-926.	1.0	79
17	Fusicatenibacter saccharivorans gen. nov., sp. nov., isolated from human faeces. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 3691-3696.	0.8	78
18	Bifidobacterium and Lactobacillus Counts in the Gut Microbiota of Patients With Bipolar Disorder and Healthy Controls. Frontiers in Psychiatry, 2018, 9, 730.	1.3	73

Hirokazu Tsuji

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19	Isolation and characterization of the equol-producing bacterium Slackia sp. strain NATTS. Archives of Microbiology, 2010, 192, 279-287.	1.0	69
20	Evolution of gut Bifidobacterium population in healthy Japanese infants over the first three years of life: a quantitative assessment. Scientific Reports, 2017, 7, 10097.	1.6	67
21	Detection of Human Intestinal Catalase-Negative, Gram-Positive Cocci by rRNA-Targeted Reverse Transcription-PCR. Applied and Environmental Microbiology, 2010, 76, 5440-5451.	1.4	63
22	Prostate Cancer Chemoprevention Study: An investigative randomized control study using purified isoflavones in men with rising prostateâ€specific antigen. Cancer Science, 2012, 103, 125-130.	1.7	59
23	Metabolic endotoxemia promotes neuroinflammation after focal cerebral ischemia. Journal of Cerebral Blood Flow and Metabolism, 2020, 40, 2505-2520.	2.4	58
24	Identification of an Enzyme System for Daidzein-to-Equol Conversion in Slackia sp. Strain NATTS. Applied and Environmental Microbiology, 2012, 78, 1228-1236.	1.4	57
25	Bacterial rRNA-Targeted Reverse Transcription-PCR Used To Identify Pathogens Responsible for Fever with Neutropenia. Journal of Clinical Microbiology, 2010, 48, 1624-1628.	1.8	56
26	Diversity of Intestinal Clostridium coccoides Group in the Japanese Population, as Demonstrated by Reverse Transcription-Quantitative PCR. PLoS ONE, 2015, 10, e0126226.	1.1	54
27	Association between Yogurt Consumption and Intestinal Microbiota in Healthy Young Adults Differs by Host Gender. Frontiers in Microbiology, 2017, 8, 847.	1.5	54
28	A Hydrogen Peroxide-Forming NADH Oxidase That Functions as an Alkyl Hydroperoxide Reductase in Amphibacillus xylanus. Journal of Bacteriology, 2000, 182, 5046-5051.	1.0	48
29	Sensitive Quantification of Clostridium difficile Cells by Reverse Transcription-Quantitative PCR Targeting rRNA Molecules. Applied and Environmental Microbiology, 2012, 78, 5111-5118.	1.4	46
30	Influence of Isoflavone Intake and Equol-producing Intestinal Flora on Prostate Cancer Risk. Asian Pacific Journal of Cancer Prevention, 2013, 14, 1-4.	0.5	42
31	Gut dysbiosis following C-section instigates higher colonisation of toxigenic Clostridium perfringens in infants. Beneficial Microbes, 2017, 8, 353-365.	1.0	39
32	Sensitive quantification of Clostridium perfringens in human feces by quantitative real-time PCR targeting alpha-toxin and enterotoxin genes. BMC Microbiology, 2015, 15, 219.	1.3	35
33	Counting the Countless: Bacterial Quantification by Targeting rRNA Molecules to Explore the Human Gut Microbiota in Health and Disease. Frontiers in Microbiology, 2018, 9, 1417.	1.5	35
34	Intestinal Microbiota Profiles of Healthy Pre-School and School-Age Children and Effects of Probiotic Supplementation. Annals of Nutrition and Metabolism, 2015, 67, 257-266.	1.0	34
35	Habitual intake of fermented milk products containing Lactobacillus casei strain Shirota and a reduced risk of hypertension in older people. Beneficial Microbes, 2017, 8, 23-29.	1.0	34
36	Independent and Interactive Effects of Habitually Ingesting Fermented Milk Products Containing Lactobacillus casei Strain Shirota and of Engaging in Moderate Habitual Daily Physical Activity on the Intestinal Health of Older People. Frontiers in Microbiology, 2019, 10, 1477.	1.5	28

Hirokazu Tsuji

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37	Repeated-batch production of galactooligosaccharides from lactose at high concentration by using alginate-immobilized cells of Sporobolomyces singularis YIT 10047. Journal of General and Applied Microbiology, 2008, 54, 285-293.	0.4	25
38	Establishment of a sensitive system for analysis of human vaginal microbiota on the basis of rRNA-targeted reverse transcription-quantitative PCR. Journal of Microbiological Methods, 2015, 111, 93-104.	0.7	23
39	Characterization of the Gut Microbiota of Papua New Guineans Using Reverse Transcription Quantitative PCR. PLoS ONE, 2015, 10, e0117427.	1.1	22
40	<i>Bifidobacterium</i> Supplementation of Colostrum and Breast Milk Enhances Weight Gain and Metabolic Responses Associated with Microbiota Establishment in Very-Preterm Infants. Biomedicine Hub, 2020, 4, 1-10.	0.4	22
41	Development of a sensitive rRNAâ€ŧargeted reverse transcriptionâ€quantitative polymerase chain reaction for detection of <i>Vibrio cholerae/mimicus</i> , <i>V. parahaemolyticus/alginolyticus</i> and <i>Campylobacter jejuni/coli</i> . Microbiology and Immunology, 2012, 56, 10-20.	0.7	20
42	Multiple Transporters and Glycoside Hydrolases Are Involved in Arabinoxylan-Derived Oligosaccharide Utilization in Bifidobacterium pseudocatenulatum. Applied and Environmental Microbiology, 2020, 86, .	1.4	18
43	Intestinal Enterobacteriaceae and Escherichia coli populations in Japanese adults demonstrated by the reverse transcription-quantitative PCR and the clone library analyses. Journal of Microbiological Methods, 2013, 92, 213-219.	0.7	16
44	Relationship of serum levels and dietary intake of isoflavone, and the novel bacterium Slackia sp. strain NATTS with the risk of prostate cancer: a case–control study among Japanese men. International Urology and Nephrology, 2016, 48, 1453-1460.	0.6	16
45	Association of life habits and fermented milk intake with stool frequency, defecatory symptoms and intestinal microbiota in healthy Japanese adults. Beneficial Microbes, 2019, 10, 841-854.	1.0	16
46	A practical random mutagenesis system for probiotic <i>Lactobacillus casei</i> using Tn <i>5</i> transposition complexes. Journal of Applied Microbiology, 2010, 109, 657-666.	1.4	15
47	Up to Species-level Community Analysis of Human Gut Microbiota by 16S rRNA Amplicon Pyrosequencing. Bioscience of Microbiota, Food and Health, 2013, 32, 69-76.	0.8	15
48	Counts of Slackia sp. strain NATTS in Intestinal Flora are Correlated to Serum Concentrations of Equol both in Prostate Cancer Cases and Controls in Japanese Men. Asian Pacific Journal of Cancer Prevention, 2014, 15, 2693-2697.	0.5	14
49	Transposon Mutagenesis of Probiotic Lactobacillus casei Identifies asnH, an Asparagine Synthetase Gene Involved in Its Immune-Activating Capacity. PLoS ONE, 2014, 9, e83876.	1.1	13
50	Stool preparation under anaerobic conditions contributes to retention of obligate anaerobes: potential improvement for fecal microbiota transplantation. BMC Microbiology, 2021, 21, 275.	1.3	13
51	Sensitive and rapid RT-qPCR quantification of pathogenic Candida species in human blood. Journal of Microbiological Methods, 2015, 117, 128-135.	0.7	12
52	Yearly changes in the composition of gut microbiota in the elderly, and the effect of lactobacilli intake on these changes. Scientific Reports, 2021, 11, 12765.	1.6	12
53	Sensitive Quantitative Detection of Commensal Bacteria by rRNA-Targeted Reverse Transcription-PCR. Applied and Environmental Microbiology, 2007, 73, 6695-6695.	1.4	8
54	Impacts of Habitual Diets Intake on Gut Microbial Counts in Healthy Japanese Adults. Nutrients, 2020, 12, 2414.	1.7	7

HIROKAZU TSUJI

#	Article	IF	CITATIONS
55	Development of a rapid and sensitive analytical system for Pseudomonas aeruginosa based on reverse transcription quantitative PCR targeting of rRNA molecules. Emerging Microbes and Infections, 2021, 10, 677-686.	3.0	7
56	Yakult Intestinal Flora-SCAN: A Novel Culture-Independent Analytical Method for Detection of Bacteria in the Bloodstream. Annals of Nutrition and Metabolism, 2017, 71, 4-10.	1.0	5
57	Metabolomics profile of Japanese female patients with restricting-type anorexia nervosa. Physiology and Behavior, 2021, 228, 113204.	1.0	5
58	Synthesis and Immunestimulating Activity of Lactobacilli-Originated Polysaccharide–Polymeric Microparticle Conjugates. Langmuir, 2015, 31, 1489-1495.	1.6	4
59	Gut dysbiosis induces lipopolysaccharide-mediated inflammation after cerebral ischemia in type 2 diabetic mice. Journal of the Neurological Sciences, 2017, 381, 875.	0.3	3
60	Higher enterococcus counts indicate a lower risk of colorectal adenomas: a prospective cohort study. Oncotarget, 2018, 9, 21459-21467.	0.8	3
61	Critical roles of a housekeeping sortase of probiotic Bifidobacterium bifidum in bacterium–host cell crosstalk. IScience, 2021, 24, 103363.	1.9	2
62	Enhanced Immunostimulating Activity of Lactobacilli-Mimicking Materials by Controlling Size. Bioconjugate Chemistry, 2015, 26, 1775-1781.	1.8	1
63	Intestinal dysbiosis and lowered serum lipopolysaccharide-binding protein in PD. Parkinsonism and Related Disorders, 2016, 22, e32.	1.1	1
64	Evolution of gut Bifidobacterium population in healthy Japanese infants over the first three years of life: a quantitative assessment. Scientific Reports, 2017, 7, .	1.6	1
65	Isoflavones in Japanese and Caucasian men with prostate cancer in Hawaii Journal of Clinical Oncology, 2014, 32, 241-241.	0.8	1

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