Heekuk Park

List of Publications by Year
in descending order

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44 papers	1,654 citations	394421 19 h-index	302126 39 g-index
44 all docs	44 docs citations	44 times ranked	3212 citing authors

#	Article	IF	Citations
1	Hepatic pathology in patients dying of COVID-19: a series of 40 cases including clinical, histologic, and virologic data. Modern Pathology, 2020, 33, 2147-2155.	5.5	193
2	Improved Glucose Homeostasis in Obese Mice Treated With Resveratrol Is Associated With Alterations in the Gut Microbiome. Diabetes, 2017, 66, 418-425.	0.6	189
3	Fecal Microbial Transplants Reduce Antibiotic-resistant Genes in Patients With RecurrentClostridium difficileInfection. Clinical Infectious Diseases, 2016, 62, 1479-1486.	5.8	166
4	A high-sugar diet rapidly enhances susceptibility to colitis via depletion of luminal short-chain fatty acids in mice. Scientific Reports, 2019, 9, 12294.	3.3	115
5	Fecal microbiota transplantation in the management of hepatic encephalopathy. Hepatology, 2016, 63, 339-340.	7.3	109
6	Characterization of the Fungal Microbiota (Mycobiome) in Healthy and Dandruff-Afflicted Human Scalps. PLoS ONE, 2012, 7, e32847.	2.5	105
7	Virulence factors of uropathogenic Escherichia coli of urinary tract infections and asymptomatic bacteriuria in children. Journal of Microbiology, Immunology and Infection, 2014, 47, 455-461.	3.1	100
8	Inulinâ€type fructans and whey protein both modulate appetite but only fructans alter gut microbiota in adults with overweight/obesity: A randomized controlled trial. Molecular Nutrition and Food Research, 2017, 61, 1700484.	3.3	91
9	Microbial Communities in the Upper Respiratory Tract of Patients with Asthma and Chronic Obstructive Pulmonary Disease. PLoS ONE, 2014, 9, e109710.	2.5	74
10	Amylose resistant starch (HAMâ€RS2) supplementation increases the proportion of Faecalibacterium bacteria in endâ€stage renal disease patients: Microbial analysis from a randomized placeboâ€controlled trial. Hemodialysis International, 2019, 23, 343-347.	0.9	61
11	Effect of chicory inulin-type fructanâ€“containing snack bars on the human gut microbiota in low dietary fiber consumers in a randomized crossover trial. American Journal of Clinical Nutrition, 2020, 111, 1286-1296.	4.7	47
12	The success of fecal microbial transplantation in Clostridium difficile infection correlates with bacteriophage relative abundance in the donor: a retrospective cohort study. Gut Microbes, 2019, 10, 676-687.	9.8	35
13	Real-time PCR assays for the detection and quantification of Streptococcus pneumoniae. FEMS Microbiology Letters, 2010, 310, 48-53.	1.8	34
14	Roles of the gut virome and mycobiome in faecal microbiota transplantation. The Lancet Gastroenterology and Hepatology, 2022, 7, 472-484.	8.1	34
15	Identification of the cpsA gene as a specific marker for the discrimination of Streptococcus pneumoniae from viridans group streptococci. Journal of Medical Microbiology, 2010, 59, 1146-1152.	1.8	32
16	Subgingival microbiome and clinical periodontal status in an elderly cohort: The WHICAP ancillary study of oral health. Journal of Periodontology, 2020, 91, S56-S67.	3.4	31
17	Oral Microbiome Alterations and SARS-CoV-2 Saliva Viral Load in Patients with COVID-19. Microbiology Spectrum, 2021, 9, e0005521.	3.0	31
18	A Distinctive Urinary Metabolomic Fingerprint Is Linked With Endoscopic Postoperative Disease Recurrence in Crohnâ€™s Disease Patients. Inflammatory Bowel Diseases, 2018, 24, 861-870.	1.9	24

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19	rpoA is a useful gene for identification and classification of Streptococcus pneumoniae from the closely related viridans group streptococci. FEMS Microbiology Letters, 2010, 305, 58-64.	1.8	22
20	Sex-Specific Differences in the Gut Microbiome in Response to Dietary Fiber Supplementation in IL-10-Deficient Mice. Nutrients, 2020, 12, 2088.	4.1	20
21	Development of a novel PCR assay based on the 16Sâ€“23S rRNA internal transcribed spacer region for the detection of Lactococcus garvieae. Journal of Fish Diseases, 2012, 35, 481-487.	1.9	19
22	Roux-en-Y gastric bypass and sleeve gastrectomy induce substantial and persistent changes in microbial communities and metabolic pathways. Gut Microbes, 2022, 14, 2050636.	9.8	16
23	Simultaneous Detection of Streptococcus pneumoniae, S. mitis, and S. oralis by a Novel Multiplex PCR Assay Targeting the gyrB Gene. Journal of Clinical Microbiology, 2013, 51, 835-840.	3.9	14
24	Analysis of Oropharyngeal Microbiota between the Patients with Bronchial Asthma and the Non-Asthmatic Persons. Journal of Bacteriology and Virology, 2013, 43, 270.	0.1	14
25	Antibiotic Exposure, Not Alloreactivity, Is the Major Driver of Microbiome Changes in Hematopoietic Cell Transplantation. Transplantation and Cellular Therapy, 2022, 28, 135-144.	1.2	11
26	Prebiotic Supplementation Following Ileocecal Resection in a Murine Model is Associated With a Loss of Microbial Diversity and Increased Inflammation. Inflammatory Bowel Diseases, 2018, 24, 101-110.	1.9	10
27	Endospore forming bacteria may be associated with maintenance of surgically-induced remission in Crohnâ€™s disease. Scientific Reports, 2018, 8, 9734.	3.3	10
28	Ileal microbial shifts after Roux-en-Y gastric bypass orchestrate changes in glucose metabolism through modulation of bile acids and L-cell adaptation. Scientific Reports, 2021, 11, 23813.	3.3	10
29	Population structure ofblaKPC-harbouring IncN plasmids at a New York City medical centre and evidence for multi-species horizontal transmission. Journal of Antimicrobial Chemotherapy, 2022, 77, 1873-1882.	3.0	7
30	The rgg Gene is a Specific Marker for Streptococcus oralis. Journal of Dental Research, 2010, 89, 1299-1303.	5.2	6
31	Development of a 16S-23S rRNA intergenic spacer-based quantitative PCR assay for improved detection and enumeration ofLactococcus garvieae. FEMS Microbiology Letters, 2013, 339, 10-16.	1.8	5
32	Lack of Effect of Gluten Challenge on Fecal Microbiome in Patients With Celiac Disease and Non-Celiac Gluten Sensitivity. Clinical and Translational Gastroenterology, 2021, 12, e00441.	2.5	4
33	Identification of a pheA Gene Associated with Streptococcus mitis by Using Suppression Subtractive Hybridization. Applied and Environmental Microbiology, 2012, 78, 3004-3009.	3.1	3
34	Salivary microbiome differences in prepubertal children with and without adrenal androgen excess. Pediatric Research, 2022, 91, 1797-1803.	2.3	3
35	Comparative genome analysis of Lactococcus garvieae using a suppression subtractive hybridization library: discovery of novel DNA signatures. FEMS Microbiology Letters, 2011, 325, 77-84.	1.8	2
36	Endospore-Forming Bacteria are Associated with Maintenance of Remission following Intestinal Resection in Crohnâ€™s Disease. Gastroenterology, 2017, 152, S192-S193.	1.3	2

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37	Timing of Tributyrin Supplementation Differentially Modulates Gastrointestinal Inflammation and Gut Microbial Recolonization Following Murine Ileocecal Resection. Nutrients, 2021, 13, 2069.	4.1	2
38	A Metaviromic Analysis of Viral Communities in the Feces of Unexplained Acute Gastroenteritis. Journal of Bacteriology and Virology, 2013, 43, 290.	0.1	1
39	Su1888 Rifaximin in Combination With a Western-Style Diet Induces Ileal Inflammation and Enhances Growth of Proteobacteria in IL-10-/- Mice. Gastroenterology, 2016, 150, S580.	1.3	1
40	Oral Acetate Reduces a High Sugar Diet Induced Increased Susceptibility to Colitis. Gastroenterology, 2017, 152, S997.	1.3	1
41	546 High Sugar Diets Promote an Inflammatory Microbiota and Reduce Gene Expression Related to Intestinal Barrier Function. Gastroenterology, 2016, 150, S114-S115.	1.3	0
42	Altered Phage Diversity and Increased Growth Rate of Escherichia Coli are Associated with Fecal Transplantation Failure in Patients with Clostridium Difficile Infection. Gastroenterology, 2017, 152, S191.	1.3	0
43	Fecal Microbiota Transplantation in the Management of Mild Overt Hepatic Encephalopathy: A Case Report: Presidential Poster. American Journal of Gastroenterology, 2015, 110, S312-S313.	0.4	0
44	Autologous Transplant Recipients Have a Healthier Gut Microbiota at Baseline and Faster Recovery from Microbiome Injury Compared to Allogeneic Transplant Recipients. Blood, 2019, 134, 4491-4491.	1.4	0