

Vincent B Young

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

219 papers	17,795 citations	75 h-index	130 g-index
265 ext. papers	22,022 ext. citations	7.4 avg, IF	7.14 L-index

#	Paper	IF	Citations
219	A Dietary Fiber-Deprived Gut Microbiota Degrades the Colonic Mucus Barrier and Enhances Pathogen Susceptibility. <i>Cell</i> , 2016 , 167, 1339-1353.e21	56.2	1149
218	Decreased diversity of the fecal Microbiome in recurrent Clostridium difficile-associated diarrhea. <i>Journal of Infectious Diseases</i> , 2008 , 197, 435-8	7	764
217	The gut microbiome in health and in disease. <i>Current Opinion in Gastroenterology</i> , 2015 , 31, 69-75	3	721
216	Analysis of the lung microbiome in the "healthy" smoker and in COPD. <i>PLoS ONE</i> , 2011 , 6, e16384	3.7	614
215	Antibiotic-induced shifts in the mouse gut microbiome and metabolome increase susceptibility to Clostridium difficile infection. <i>Nature Communications</i> , 2014 , 5, 3114	17.4	568
214	Comparison of the respiratory microbiome in healthy nonsmokers and smokers. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2013 , 187, 1067-75	10.2	501
213	Defining a healthy human gut microbiome: current concepts, future directions, and clinical applications. <i>Cell Host and Microbe</i> , 2012 , 12, 611-22	23.4	448
212	Decade-long bacterial community dynamics in cystic fibrosis airways. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 5809-14	11.5	431
211	Analysis of the upper respiratory tract microbiotas as the source of the lung and gastric microbiotas in healthy individuals. <i>MBio</i> , 2015 , 6, e00037	7.8	429
210	Reproducible community dynamics of the gastrointestinal microbiota following antibiotic perturbation. <i>Infection and Immunity</i> , 2009 , 77, 2367-75	3.7	418
209	Gut microbiome-derived metabolites modulate intestinal epithelial cell damage and mitigate graft-versus-host disease. <i>Nature Immunology</i> , 2016 , 17, 505-513	19.1	366
208	Role of the intestinal microbiota in resistance to colonization by Clostridium difficile. <i>Gastroenterology</i> , 2014 , 146, 1547-53	13.3	266
207	From structure to function: the ecology of host-associated microbial communities. <i>Microbiology and Molecular Biology Reviews</i> , 2010 , 74, 453-76	13.2	251
206	Antibiotic-associated diarrhea accompanied by large-scale alterations in the composition of the fecal microbiota. <i>Journal of Clinical Microbiology</i> , 2004 , 42, 1203-6	9.7	241
205	The role of the microbiome in human health and disease: an introduction for clinicians. <i>BMJ, The</i> , 2017 , 356, j831	5.9	238
204	Application of a neutral community model to assess structuring of the human lung microbiome. <i>MBio</i> , 2015 , 6,	7.8	237
203	The microbiome of the lung. <i>Translational Research</i> , 2012 , 160, 258-66	11	236

202	Alteration of the murine gut microbiota during infection with the parasitic helminth <i>Heligmosomoides polygyrus</i> . <i>Inflammatory Bowel Diseases</i> , 2010 , 16, 1841-9	4.5	236
201	Persistence and toxin production by <i>Clostridium difficile</i> within human intestinal organoids result in disruption of epithelial paracellular barrier function. <i>Infection and Immunity</i> , 2015 , 83, 138-45	3.7	219
200	Antibiotic-Induced Alterations of the Gut Microbiota Alter Secondary Bile Acid Production and Allow for <i>Clostridium difficile</i> Spore Germination and Outgrowth in the Large Intestine. <i>MSphere</i> , 2016 , 1,	5	216
199	Recovery of the gut microbiome following fecal microbiota transplantation. <i>MBio</i> , 2014 , 5, e00893-14	7.8	209
198	Suppression of <i>Clostridium difficile</i> in the gastrointestinal tracts of germfree mice inoculated with a murine isolate from the family Lachnospiraceae. <i>Infection and Immunity</i> , 2012 , 80, 3786-94	3.7	201
197	The interplay between microbiome dynamics and pathogen dynamics in a murine model of <i>Clostridium difficile</i> Infection. <i>Gut Microbes</i> , 2011 , 2, 145-58	8.8	192
196	Microbiome data distinguish patients with <i>Clostridium difficile</i> infection and non-C. <i>difficile</i> -associated diarrhea from healthy controls. <i>MBio</i> , 2014 , 5, e01021-14	7.8	185
195	Interaction between the intestinal microbiota and host in <i>Clostridium difficile</i> colonization resistance. <i>Trends in Microbiology</i> , 2012 , 20, 313-9	12.4	171
194	<i>Candida albicans</i> and bacterial microbiota interactions in the cecum during recolonization following broad-spectrum antibiotic therapy. <i>Infection and Immunity</i> , 2012 , 80, 3371-80	3.7	170
193	The gut microbiome composition associates with bipolar disorder and illness severity. <i>Journal of Psychiatric Research</i> , 2017 , 87, 23-29	5.2	167
192	Interactions Between the Gastrointestinal Microbiome and <i>Clostridium difficile</i> . <i>Annual Review of Microbiology</i> , 2015 , 69, 445-61	17.5	167
191	Changes in cystic fibrosis airway microbiota at pulmonary exacerbation. <i>Annals of the American Thoracic Society</i> , 2013 , 10, 179-87	4.7	161
190	Significance of the microbiome in obstructive lung disease. <i>Thorax</i> , 2012 , 67, 456-63	7.3	161
189	Evolutionary genetics of a new pathogenic <i>Escherichia</i> species: <i>Escherichia albertii</i> and related <i>Shigella boydii</i> strains. <i>Journal of Bacteriology</i> , 2005 , 187, 619-28	3.5	153
188	NLRP12 attenuates colon inflammation by maintaining colonic microbial diversity and promoting protective commensal bacterial growth. <i>Nature Immunology</i> , 2017 , 18, 541-551	19.1	151
187	<i>Clostridium difficile</i> ribotype does not predict severe infection. <i>Clinical Infectious Diseases</i> , 2012 , 55, 1661-8	11.6	144
186	Microbial ecology of the murine gut associated with the development of dextran sodium sulfate-induced colitis. <i>Inflammatory Bowel Diseases</i> , 2011 , 17, 917-26	4.5	143
185	<i>Clostridium difficile</i> and the microbiota. <i>Journal of Clinical Investigation</i> , 2014 , 124, 4182-9	15.9	142

184	The role of the microbiota in infectious diseases. <i>Nature Microbiology</i> , 2019 , 4, 35-45	26.6	142
183	Widespread colonization of the lung by <i>Tropheryma whippelii</i> in HIV infection. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2013 , 187, 1110-7	10.2	140
182	Gut Microbiota and Colonization Resistance against Bacterial Enteric Infection. <i>Microbiology and Molecular Biology Reviews</i> , 2019 , 83,	13.2	126
181	Functional Characterization of Inflammatory Bowel Disease-Associated Gut Dysbiosis in Gnotobiotic Mice. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2016 , 2, 468-481	7.9	123
180	Cytotoxic distending toxin sequence and activity in the enterohepatic pathogen <i>Helicobacter hepaticus</i> . <i>Infection and Immunity</i> , 2000 , 68, 184-91	3.7	122
179	Antibiotic administration alters the community structure of the gastrointestinal microbiota. <i>Gut Microbes</i> , 2010 , 1, 279-284	8.8	117
178	The intestinal microbiota in health and disease. <i>Current Opinion in Gastroenterology</i> , 2012 , 28, 63-9	3	117
177	Cefoperazone-treated mice as an experimental platform to assess differential virulence of <i>Clostridium difficile</i> strains. <i>Gut Microbes</i> , 2011 , 2, 326-34	8.8	113
176	In vitro and in vivo characterization of <i>Helicobacter hepaticus</i> cytotoxic distending toxin mutants. <i>Infection and Immunity</i> , 2004 , 72, 2521-7	3.7	113
175	Metabolic Model-Based Integration of Microbiome Taxonomic and Metabolomic Profiles Elucidates Mechanistic Links between Ecological and Metabolic Variation. <i>MSystems</i> , 2016 , 1,	7.6	108
174	Standard colonic lavage alters the natural state of mucosal-associated microbiota in the human colon. <i>PLoS ONE</i> , 2012 , 7, e32545	3.7	106
173	Perturbation of the small intestine microbial ecology by streptomycin alters pathology in a <i>Salmonella enterica</i> serovar typhimurium murine model of infection. <i>Infection and Immunity</i> , 2009 , 77, 2691-702	3.7	104
172	The nasal cavity microbiota of healthy adults. <i>Microbiome</i> , 2014 , 2, 27	16.6	103
171	Stress-induced corticotropin-releasing hormone-mediated NLRP6 inflammasome inhibition and transmissible enteritis in mice. <i>Gastroenterology</i> , 2013 , 144, 1478-87, 1487.e1-8	13.3	101
170	Dynamics and establishment of <i>Clostridium difficile</i> infection in the murine gastrointestinal tract. <i>Infection and Immunity</i> , 2015 , 83, 934-41	3.7	100
169	Multicenter Comparison of Lung and Oral Microbiomes of HIV-infected and HIV-uninfected Individuals. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2015 , 192, 1335-44	10.2	97
168	Bacterial colonization stimulates a complex physiological response in the immature human intestinal epithelium. <i>ELife</i> , 2017 , 6,	8.9	97
167	Interplay between the gastric bacterial microbiota and <i>Candida albicans</i> during postantibiotic recolonization and gastritis. <i>Infection and Immunity</i> , 2012 , 80, 150-8	3.7	95

166	A gene-targeted approach to investigate the intestinal butyrate-producing bacterial community. <i>Microbiome</i> , 2013 , 1, 8	16.6	94
165	Comparison of stool versus rectal swab samples and storage conditions on bacterial community profiles. <i>BMC Microbiology</i> , 2017 , 17, 78	4.5	94
164	<i>Clostridium difficile</i> ribotype 027: relationship to age, detectability of toxins A or B in stool with rapid testing, severe infection, and mortality. <i>Clinical Infectious Diseases</i> , 2015 , 61, 233-41	11.6	93
163	Identification of <i>cdtB</i> homologues and cytolethal distending toxin activity in enterohepatic <i>Helicobacter</i> spp. <i>Journal of Medical Microbiology</i> , 2000 , 49, 525-534	3.2	93
162	C57BL/6 and congenic interleukin-10-deficient mice can serve as models of <i>Campylobacter jejuni</i> colonization and enteritis. <i>Infection and Immunity</i> , 2007 , 75, 1099-115	3.7	88
161	Stabilization of the murine gut microbiome following weaning. <i>Gut Microbes</i> , 2012 , 3, 383-93	8.8	86
160	The Inhibitory Innate Immune Sensor NLRP12 Maintains a Threshold against Obesity by Regulating Gut Microbiota Homeostasis. <i>Cell Host and Microbe</i> , 2018 , 24, 364-378.e6	23.4	86
159	Disruption of the human gut microbiota following Norovirus infection. <i>PLoS ONE</i> , 2012 , 7, e48224	3.7	83
158	Colonizes Alternative Nutrient Niches during Infection across Distinct Murine Gut Microbiomes. <i>MSystems</i> , 2017 , 2,	7.6	82
157	Restoration of short chain fatty acid and bile acid metabolism following fecal microbiota transplantation in patients with recurrent <i>Clostridium difficile</i> infection. <i>Anaerobe</i> , 2018 , 53, 64-73	2.8	81
156	Cytolethal distending toxin in avian and human isolates of <i>Helicobacter pullorum</i> . <i>Journal of Infectious Diseases</i> , 2000 , 182, 620-3	7	80
155	The Cancer Microbiome: Distinguishing Direct and Indirect Effects Requires a Systemic View. <i>Trends in Cancer</i> , 2020 , 6, 192-204	12.5	79
154	3343 Identification of host-microbial interaction networks that mediate intestinal epithelial barrier function in necrotizing enterocolitis. <i>Journal of Clinical and Translational Science</i> , 2019 , 3, 13-13	0.4	78
153	2236. Stool-Derived Inflammatory Mediators Serve as Biomarkers of Severity in <i>Clostridium difficile</i> Infection. <i>Open Forum Infectious Diseases</i> , 2019 , 6, S764-S765	1	78
152	2403. <i>Clostridium difficile</i> ribotypes and human microbiota differ in Taiwan and the United States with respect to diarrheal patients. <i>Open Forum Infectious Diseases</i> , 2019 , 6, S829-S830	1	78
151	2849. Gut Microbiota Differences at the Time of Medical Intensive Care Unit (MICU) Admission Are Associated with Acquisition of Multi-drug-Resistant Organisms (MDROs) Among Patients Not Already Colonized with an MDRO. <i>Open Forum Infectious Diseases</i> , 2019 , 6, S71-S72	1	78
150	2355. The Association Between Diagnostic Testing Method and <i>Clostridium difficile</i> Infection Severity. <i>Open Forum Infectious Diseases</i> , 2019 , 6, S811-S811	1	78
149	2424. Shedding of Viable <i>Clostridiodes difficile</i> in Patients Admitted to a Medical Intensive Care Unit. <i>Open Forum Infectious Diseases</i> , 2019 , 6, S837-S838	1	78

148	2409. External Validation and Comparison of Clostridioides difficile Severity Scoring Systems. <i>Open Forum Infectious Diseases</i> , 2019 , 6, S831-S832	1	78
147	3185 A Randomized Controlled Trial Comparing the Nonabsorbable Antibiotic Rifaximin vs. Dietary Intervention Low in Fermentable Sugars (FODMAP) in Irritable Bowel Syndrome. <i>Journal of Clinical and Translational Science</i> , 2019 , 3, 31-31	0.4	78
146	Fecal Microbiota Transplantation Eliminates Clostridium difficile in a Murine Model of Relapsing Disease. <i>Infection and Immunity</i> , 2015 , 83, 3838-46	3.7	76
145	A Generalizable, Data-Driven Approach to Predict Daily Risk of Clostridium difficile Infection at Two Large Academic Health Centers. <i>Infection Control and Hospital Epidemiology</i> , 2018 , 39, 425-433	2	75
144	Dynamics of the fecal microbiome in patients with recurrent and nonrecurrent Clostridium difficile infection. <i>Genome Medicine</i> , 2016 , 8, 47	14.4	75
143	Comparison of brush and biopsy sampling methods of the ileal pouch for assessment of mucosa-associated microbiota of human subjects. <i>Microbiome</i> , 2014 , 2, 5	16.6	72
142	Colonization of the cecal mucosa by Helicobacter hepaticus impacts the diversity of the indigenous microbiota. <i>Infection and Immunity</i> , 2005 , 73, 6952-61	3.7	71
141	Modulation of host immune responses by the cytolethal distending toxin of Helicobacter hepaticus. <i>Infection and Immunity</i> , 2006 , 74, 4496-504	3.7	68
140	Probiotics for prevention of Clostridium difficile infection. <i>Current Opinion in Gastroenterology</i> , 2018 , 34, 3-10	3	66
139	Microbial Metabolite Signaling Is Required for Systemic Iron Homeostasis. <i>Cell Metabolism</i> , 2020 , 31, 115-130.e6	24.6	64
138	Pathogenesis of renal disease due to enterohemorrhagic Escherichia coli in germ-free mice. <i>Infection and Immunity</i> , 2008 , 76, 3054-63	3.7	59
137	Interleukin-22-mediated host glycosylation prevents Clostridioides difficile infection by modulating the metabolic activity of the gut microbiota. <i>Nature Medicine</i> , 2020 , 26, 608-617	50.5	58
136	The relationship between phenotype, ribotype, and clinical disease in human Clostridium difficile isolates. <i>Anaerobe</i> , 2013 , 24, 109-16	2.8	55
135	The anti-inflammatory drug mesalamine targets bacterial polyphosphate accumulation. <i>Nature Microbiology</i> , 2017 , 2, 16267	26.6	54
134	Alteration of the murine gastrointestinal microbiota by tigecycline leads to increased susceptibility to Clostridium difficile infection. <i>Antimicrobial Agents and Chemotherapy</i> , 2014 , 58, 2767-74	5.9	53
133	Streptococcus intermedius causing infective endocarditis and abscesses: a report of three cases and review of the literature. <i>BMC Infectious Diseases</i> , 2008 , 8, 154	4	53
132	Inflammatory bowel disease causes reversible suppression of osteoblast and chondrocyte function in mice. <i>American Journal of Physiology - Renal Physiology</i> , 2009 , 296, G1020-9	5.1	52
131	Depression, antidepressant medications, and risk of Clostridium difficile infection. <i>BMC Medicine</i> , 2013 , 11, 121	11.4	51

130	Fecal microbiota transplantation for the management of Clostridium difficile infection. <i>Infectious Disease Clinics of North America</i> , 2015 , 29, 109-22	6.5	50
129	Microbial and metabolic interactions between the gastrointestinal tract and Clostridium difficile infection. <i>Gut Microbes</i> , 2014 , 5, 86-95	8.8	50
128	Variation of the natural transformation frequency of Campylobacter jejuni in liquid shake culture. <i>Microbiology (United Kingdom)</i> , 2003 , 149, 3603-3615	2.9	49
127	A clinical and epidemiological review of non-toxigenic Clostridium difficile. <i>Anaerobe</i> , 2013 , 22, 1-5	2.8	48
126	Leptin acts independently of food intake to modulate gut microbial composition in male mice. <i>Endocrinology</i> , 2014 , 155, 748-57	4.8	45
125	Ecological succession of bacterial communities during conventionalization of germ-free mice. <i>Applied and Environmental Microbiology</i> , 2012 , 78, 2359-66	4.8	45
124	Impact of enhanced Staphylococcus DNA extraction on microbial community measures in cystic fibrosis sputum. <i>PLoS ONE</i> , 2012 , 7, e33127	3.7	44
123	Alters the Structure and Metabolism of Distinct Cecal Microbiomes during Initial Infection To Promote Sustained Colonization. <i>MSphere</i> , 2018 , 3,	5	43
122	Colitis-induced bone loss is gender dependent and associated with increased inflammation. <i>Inflammatory Bowel Diseases</i> , 2013 , 19, 1586-97	4.5	43
121	Tryptophan catabolism restricts IFN- γ -expressing neutrophils and Clostridium difficile immunopathology. <i>Journal of Immunology</i> , 2014 , 193, 807-16	5.3	42
120	Poor functional status as a risk factor for severe Clostridium difficile infection in hospitalized older adults. <i>Journal of the American Geriatrics Society</i> , 2013 , 61, 1738-42	5.6	41
119	Interleukin-23 (IL-23), independent of IL-17 and IL-22, drives neutrophil recruitment and innate inflammation during Clostridium difficile colitis in mice. <i>Immunology</i> , 2016 , 147, 114-24	7.8	39
118	Laser capture microdissection and metagenomic analysis of intact mucosa-associated microbial communities of human colon. <i>Applied Microbiology and Biotechnology</i> , 2010 , 88, 1333-42	5.7	38
117	Evaluation of portability and cost of a fluorescent PCR ribotyping protocol for Clostridium difficile epidemiology. <i>Journal of Clinical Microbiology</i> , 2015 , 53, 1192-7	9.7	36
116	Clostridium difficile ribotype diversity at six health care institutions in the United States. <i>Journal of Clinical Microbiology</i> , 2013 , 51, 1938-41	9.7	36
115	The systemic inflammatory response to Clostridium difficile infection. <i>PLoS ONE</i> , 2014 , 9, e92578	3.7	36
114	Increased Relative Abundance of Klebsiella pneumoniae Carbapenemase-producing Klebsiella pneumoniae Within the Gut Microbiota Is Associated With Risk of Bloodstream Infection in Long-term Acute Care Hospital Patients. <i>Clinical Infectious Diseases</i> , 2019 , 68, 2053-2059	11.6	36
113	Procalcitonin levels associate with severity of Clostridium difficile infection. <i>PLoS ONE</i> , 2013 , 8, e58265	3.7	32

112	Variation in germination of <i>Clostridium difficile</i> clinical isolates correlates to disease severity. <i>Anaerobe</i> , 2015 , 33, 64-70	2.8	31
111	Role of GM-CSF in the inflammatory cytokine network that regulates neutrophil influx into the colonic mucosa during <i>Clostridium difficile</i> infection in mice. <i>Gut Microbes</i> , 2014 , 5, 476-84	8.8	31
110	Therapeutic manipulation of the microbiota: past, present, and considerations for the future. <i>Clinical Microbiology and Infection</i> , 2016 , 22, 905-909	9.5	29
109	Multiphasic analysis of the temporal development of the distal gut microbiota in patients following ileal pouch anal anastomosis. <i>Microbiome</i> , 2013 , 1, 9	16.6	29
108	Spatial and Temporal Analysis of the Stomach and Small-Intestinal Microbiota in Fasted Healthy Humans. <i>MSphere</i> , 2019 , 4,	5	28
107	Murine norovirus infection does not cause major disruptions in the murine intestinal microbiota. <i>Microbiome</i> , 2013 , 1, 7	16.6	27
106	High-resolution profiling of the gut microbiome reveals the extent of burden. <i>Npj Biofilms and Microbiomes</i> , 2017 , 3, 35	8.2	27
105	Overview of the gastrointestinal microbiota. <i>Advances in Experimental Medicine and Biology</i> , 2008 , 635, 29-40	3.6	27
104	Real-time Measurement of Epithelial Barrier Permeability in Human Intestinal Organoids. <i>Journal of Visualized Experiments</i> , 2017 ,	1.6	26
103	Acute infection of mice with <i>Clostridium difficile</i> leads to eIF2 γ phosphorylation and pro-survival signalling as part of the mucosal inflammatory response. <i>Immunology</i> , 2013 , 140, 111-22	7.8	26
102	Challenges in IBD research: update on progress and prioritization of the CCFAR research agenda. <i>Inflammatory Bowel Diseases</i> , 2013 , 19, 677-82	4.5	26
101	Ulcerative typhlocolitis associated with <i>Helicobacter mastomyrinus</i> in telomerase-deficient mice. <i>Veterinary Pathology</i> , 2011 , 48, 713-25	2.8	26
100	Lethal toxin is a critical determinant of rapid mortality in rodent models of <i>Clostridium sordellii</i> endometritis. <i>Anaerobe</i> , 2010 , 16, 155-60	2.8	26
99	Lessons learned from the prenatal microbiome controversy. <i>Microbiome</i> , 2021 , 9, 8	16.6	25
98	Using Machine Learning and the Electronic Health Record to Predict Complicated Infection. <i>Open Forum Infectious Diseases</i> , 2019 , 6, ofz186	1	24
97	The effects of intestinal microbial community structure on disease manifestation in IL-10 $^{-/-}$ mice infected with <i>Helicobacter hepaticus</i> . <i>Microbiome</i> , 2013 , 1, 15	16.6	24
96	Novel therapies and preventative strategies for primary and recurrent <i>Clostridium difficile</i> infections. <i>Annals of the New York Academy of Sciences</i> , 2019 , 1435, 110-138	6.5	24
95	The Role of Fecal Microbiota Transplantation in Reducing Intestinal Colonization With Antibiotic-Resistant Organisms: The Current Landscape and Future Directions. <i>Open Forum Infectious Diseases</i> , 2019 , 6,	1	23

94	Detection of mixed populations of <i>Clostridium difficile</i> from symptomatic patients using capillary-based polymerase chain reaction ribotyping. <i>Infection Control and Hospital Epidemiology</i> , 2013 , 34, 961-966	2	23
93	Effect of sample storage conditions on culture-independent bacterial community measures in cystic fibrosis sputum specimens. <i>Journal of Clinical Microbiology</i> , 2011 , 49, 3717-8	9.7	22
92	Genetic diversity of <i>Campylobacter</i> sp. isolates from retail chicken products and humans with gastroenteritis in Central Michigan. <i>Journal of Clinical Microbiology</i> , 2005 , 43, 4221-4	9.7	22
91	Faecal microbiota transplantation for the treatment of recurrent <i>Clostridium difficile</i> infection: current promise and future needs. <i>Current Opinion in Gastroenterology</i> , 2013 , 29, 628-32	3	21
90	A randomised trial of sheathed versus standard forceps for obtaining uncontaminated biopsy specimens of microbiota from the terminal ileum. <i>Gut</i> , 2011 , 60, 1043-9	19.2	21
89	Explaining unexplained diarrhea and associating risks and infections. <i>Animal Health Research Reviews</i> , 2007 , 8, 69-80	2.1	21
88	Reducing Recurrence of <i>C. difficile</i> Infection. <i>Cell</i> , 2017 , 169, 375	56.2	20
87	The Gut Microbiota Is Associated with Clearance of <i>Clostridium difficile</i> Infection Independent of Adaptive Immunity. <i>MSphere</i> , 2019 , 4,	5	20
86	Emerging Insights into Antibiotic-Associated Diarrhea and <i>Clostridium difficile</i> Infection through the Lens of Microbial Ecology. <i>Interdisciplinary Perspectives on Infectious Diseases</i> , 2008 , 2008, 125081	1.7	20
85	Elevated fecal calprotectin associates with adverse outcomes from <i>Clostridium difficile</i> infection in older adults. <i>Infectious Diseases</i> , 2016 , 48, 663-9	3.1	20
84	Interleukin-22 and CD160 play additive roles in the host mucosal response to <i>Clostridium difficile</i> infection in mice. <i>Immunology</i> , 2015 , 144, 587-97	7.8	19
83	Murine models to study <i>Clostridium difficile</i> infection and transmission. <i>Anaerobe</i> , 2013 , 24, 94-7	2.8	19
82	Studying the Enteric Microbiome in Inflammatory Bowel Diseases: Getting through the Growing Pains and Moving Forward. <i>Frontiers in Microbiology</i> , 2011 , 2, 144	5.7	19
81	Impact of a hormone-releasing intrauterine system on the vaginal microbiome: a prospective baboon model. <i>Journal of Medical Primatology</i> , 2014 , 43, 89-99	0.7	18
80	Human microbiome science: vision for the future, Bethesda, MD, July 24 to 26, 2013. <i>Microbiome</i> , 2014 , 2,	16.6	18
79	Presence of multiple <i>Clostridium difficile</i> strains at primary infection is associated with development of recurrent disease. <i>Anaerobe</i> , 2018 , 53, 74-81	2.8	18
78	The rest of the story: the microbiome and gastrointestinal infections. <i>Current Opinion in Microbiology</i> , 2015 , 23, 121-5	7.9	17
77	Chronic atrophic gastritis in SCID mice experimentally infected with <i>Campylobacter fetus</i> . <i>Infection and Immunity</i> , 2000 , 68, 2110-8	3.7	17

76	Role of interferon- γ and inflammatory monocytes in driving colonic inflammation during acute Clostridium difficile infection in mice. <i>Immunology</i> , 2017 , 150, 468-477	7.8	16
75	A whole new ball game: Stem cell-derived epithelia in the study of host-microbe interactions. <i>Anaerobe</i> , 2016 , 37, 25-8	2.8	16
74	Increases in Colonic Bacterial Diversity after β Fatty Acid Supplementation Predict Decreased Colonic Prostaglandin E2 Concentrations in Healthy Adults. <i>Journal of Nutrition</i> , 2019 , 149, 1170-1179	4.1	15
73	Effects of intrauterine contraception on the vaginal microbiota. <i>Contraception</i> , 2017 , 96, 189-195	2.5	14
72	The role of the humoral immune response to Clostridium difficile toxins A and B in susceptibility to C. difficile infection: a case-control study. <i>Anaerobe</i> , 2014 , 27, 82-6	2.8	14
71	Non-toxigenic Clostridium sordellii: clinical and microbiological features of a case of cholangitis-associated bacteremia. <i>Anaerobe</i> , 2011 , 17, 252-6	2.8	14
70	An Observational Cohort Study of Ribotype 027 and Recurrent Infection. <i>MSphere</i> , 2018 , 3,	5	14
69	Emergence of Carbapenemase-producing Klebsiella Pneumoniae of Sequence type 258 in Michigan, USA. <i>Gastroenterology Insights</i> , 2013 , 5, e5	2.1	13
68	Relationship between COMLEX and USMLE scores among osteopathic medical students who take both examinations. <i>Teaching and Learning in Medicine</i> , 2010 , 22, 3-7	3.4	13
67	Dietary Xanthan Gum Alters Antibiotic Efficacy against the Murine Gut Microbiota and Attenuates Colonization. <i>MSphere</i> , 2020 , 5,	5	13
66	Systemic Inflammatory Mediators Are Effective Biomarkers for Predicting Adverse Outcomes in Clostridioides difficile Infection. <i>MBio</i> , 2020 , 11,	7.8	12
65	Genetic Determinants of Trehalose Utilization Are Not Associated With Severe Infection Outcome. <i>Open Forum Infectious Diseases</i> , 2020 , 7, ofz548	1	12
64	The role of Gr-1(+) cells and tumour necrosis factor- β signalling during Clostridium difficile colitis in mice. <i>Immunology</i> , 2015 , 144, 704-16	7.8	12
63	Effects of tigecycline and vancomycin administration on established Clostridium difficile infection. <i>Antimicrobial Agents and Chemotherapy</i> , 2015 , 59, 1596-604	5.9	10
62	Prevalence of human norovirus and Clostridium difficile coinfections in adult hospitalized patients. <i>Clinical Epidemiology</i> , 2016 , 8, 253-60	5.9	10
61	Impact of the Levonorgestrel-Releasing Intrauterine System on the Progression of Chlamydia trachomatis Infection to Pelvic Inflammatory Disease in a Baboon Model. <i>Journal of Infectious Diseases</i> , 2018 , 217, 656-666	7	9
60	Low prevalence of Clostridium septicum fecal carriage in an adult population. <i>Anaerobe</i> , 2015 , 32, 34-36	2.8	8
59	Temporal Gut Microbial Changes Predict Recurrent Clostridioides Difficile Infection in Patients With and Without Ulcerative Colitis. <i>Inflammatory Bowel Diseases</i> , 2020 , 26, 1748-1758	4.5	8

58	Storage duration of red blood cell transfusion and Clostridium difficile infection: a within person comparison. <i>PLoS ONE</i> , 2014 , 9, e89332	3.7	8
57	The levonorgestrel-releasing intrauterine system is associated with delayed endocervical clearance of Chlamydia trachomatis without alterations in vaginal microbiota. <i>Pathogens and Disease</i> , 2015 , 73, ftv070	4.2	7
56	Clostridium difficile-induced colitis in mice is independent of leukotrienes. <i>Anaerobe</i> , 2014 , 30, 90-8	2.8	7
55	Fecal microbiota therapy: ready for prime time?. <i>Infection Control and Hospital Epidemiology</i> , 2014 , 35, 28-30	2	7
54	Spherules, hyphae, and air-crescent sign. <i>American Journal of the Medical Sciences</i> , 2008 , 335, 504-6	2.2	7
53	The human microbiome and infectious diseases: beyond koch. <i>Interdisciplinary Perspectives on Infectious Diseases</i> , 2008 , 2008, 296873	1.7	7
52	Protection from Lethal Clostridioides difficile Infection via Intraspecies Competition for Cogerminant. <i>MBio</i> , 2021 , 12,	7.8	7
51	Microbiome therapeutics for hepatic encephalopathy. <i>Journal of Hepatology</i> , 2021 , 75, 1452-1464	13.4	7
50	Gut Microbiota and Clinical Features Distinguish Colonization With Carbapenemase-Producing at the Time of Admission to a Long-term Acute Care Hospital. <i>Open Forum Infectious Diseases</i> , 2018 , 5, ofy190	1.9	6
49	Cytolethal distending toxin: a bacterial toxin which disrupts the eukaryotic cell cycle. <i>Chemical Research in Toxicology</i> , 2000 , 13, 936-9	4	6
48	Co-cultivation of microbial sub-communities in microfluidic droplets facilitates high-resolution genomic dissection of microbial dark matter <i>Integrative Biology (United Kingdom)</i> , 2020 , 12, 263-274	3.7	6
47	Changes in the Association Between Diagnostic Testing Method, Polymerase Chain Reaction Ribotype, and Clinical Outcomes From Clostridioides difficile Infection: One Institution's Experience. <i>Clinical Infectious Diseases</i> , 2021 , 73, e2883-e2889	11.6	6
46	Aging Dampens the Intestinal Innate Immune Response during Severe Clostridioides difficile Infection and Is Associated with Altered Cytokine Levels and Granulocyte Mobilization. <i>Infection and Immunity</i> , 2020 , 88,	3.7	6
45	Gastrointestinal Microbial Ecology with Perspectives on Health and Disease 2012 , 1119-1134		5
44	Author response: Bacterial colonization stimulates a complex physiological response in the immature human intestinal epithelium 2017 ,		5
43	Salmonella enterica Serovar Typhimurium SPI-1 and SPI-2 Shape the Global Transcriptional Landscape in a Human Intestinal Organoid Model System. <i>MBio</i> , 2021 , 12,	7.8	5
42	Bacteria Detected in both Urine and Open Wounds in Nursing Home Residents: a Pilot Study. <i>MSphere</i> , 2019 , 4,	5	5
41	Methods for Characterizing Microbial Communities Associated with the Human Body		5

40	Understanding increased mortality in Clostridium difficile-infected older adults. <i>Clinical Infectious Diseases</i> , 2013 , 57, 625-6	11.6	4
39	Is Clostridium difficile infection a risk factor for subsequent bloodstream infection?. <i>Anaerobe</i> , 2017 , 48, 27-33	2.8	3
38	Serum 25-Hydroxyvitamin D Levels are not Associated with Adverse Outcomes in Clostridium Difficile Infection. <i>Gastroenterology Insights</i> , 2015 , 7, 5979	2.1	3
37	Gender Differences in Non-Toxigenic Colonization and Risk of Subsequent 2015 , 2,		3
36	A plasmid locus associated with Klebsiella clinical infections encodes a microbiome-dependent gut fitness factor. <i>PLoS Pathogens</i> , 2021 , 17, e1009537	7.6	3
35	Intestinal Inflammation and Altered Gut Microbiota Associated with Inflammatory Bowel Disease Render Mice Susceptible to Clostridioides difficile Colonization and Infection. <i>MBio</i> , 2021 , 12, e0273320	7.8	3
34	Unexpected Results From a Phase 2 Trial of a Microbiome Therapeutic for Clostridioides difficile Infection: Lessons for the Future. <i>Clinical Infectious Diseases</i> , 2021 , 72, 2141-2143	11.6	3
33	Mechanistic insights into consumption of the food additive xanthan gum by the human gut microbiota.. <i>Nature Microbiology</i> , 2022 , 7, 556-569	26.6	3
32	Old and new models for studying host-microbe interactions in health and disease: as an example. <i>American Journal of Physiology - Renal Physiology</i> , 2017 , 312, G623-G627	5.1	2
31	Outbreak of Murine Infection with Associated with the Administration of a Pre- and Perinatal Methyl Donor Diet. <i>MSphere</i> , 2019 , 4,	5	2
30	Reply to Planche et al. <i>Clinical Infectious Diseases</i> , 2015 , 61, 1211-2	11.6	2
29	Fecal Microbiota Transplantations: Where Are We, Where Are We Going, and What Is the Role of the Clinical Laboratory?. <i>Clinical Chemistry</i> , 2020 , 66, 512-517	5.5	2
28	Comparative transcriptional profiling of the early host response to infection by typhoidal and non-typhoidal Salmonella serovars in human intestinal organoids. <i>PLoS Pathogens</i> , 2021 , 17, e1009987	7.6	2
27	Clostridium difficile Infection 2015 , 2744-2756.e3		2
26	Clostridium difficile colonizes alternative nutrient niches during infection across distinct murine gut microbiomes		2
25	Clostridium difficile alters the structure and metabolism of distinct cecal microbiomes during initial infection to promote sustained colonization		2
24	The Lumen of Human Intestinal Organoids Poses Greater Stress to Bacteria Compared to the Germ-Free Mouse Intestine: Escherichia coli Deficient in RpoS as a Colonization Probe. <i>MSphere</i> , 2020 , 5,	5	2
23	Toward Accurate and Robust Environmental Surveillance Using Metagenomics. <i>Frontiers in Genetics</i> , 2021 , 12, 600111	4.5	2

22	Anti-toxin antibody is not associated with recurrent <i>Clostridium difficile</i> infection. <i>Anaerobe</i> , 2021 , 67, 102299	2.8	2
21	Gastrointestinal Microbial Ecology With Perspectives on Health and Disease 2018 , 737-753		2
20	Infection: Modulation of <i>Clostridium difficile</i> infection by dietary zinc. <i>Nature Reviews Gastroenterology and Hepatology</i> , 2016 , 13, 686-688	24.2	1
19	Reply to Walker et al. <i>Clinical Infectious Diseases</i> , 2013 , 56, 1846-7	11.6	1
18	Reply to McDonald. <i>Clinical Infectious Diseases</i> , 2013 , 56, 907-8	11.6	1
17	Inter-individual Recovery of the Microbiota and Metabolome Over Time Following Fecal Microbiota Transplantation in Patients with Recurrent <i>Clostridium difficile</i> Infection		1
16	A plasmid locus associated with <i>Klebsiella</i> clinical infections encodes a microbiome-dependent gut fitness factor		1
15	Vaginal microbiota of adolescents and their mothers: A preliminary study of vertical transmission and persistence		1
14	The vaginal microbiota, high-risk human papillomavirus infection, and cervical cytology: results from a population-based study.. <i>Gynecology and Pelvic Medicine</i> , 2020 , 3,	0.3	1
13	The State of Microbiome Science at the Intersection of Infectious Diseases and Antimicrobial Resistance. <i>Journal of Infectious Diseases</i> , 2021 , 223, S187-S193	7	1
12	A multisite genomic epidemiology study of infections in the USA supports differential roles of healthcare versus community spread for two common strains. <i>Microbial Genomics</i> , 2021 , 7,	4.4	1
11	76. Validation of Systemic Inflammatory Mediators as Biomarkers for Severity and Adverse Outcomes in <i>Clostridium difficile</i> Infection. <i>Open Forum Infectious Diseases</i> , 2019 , 6, S1-S2	1	1
10	Viewing Bacterial Colonization through the Lens of Systems Biology.. <i>MSystems</i> , 2022 , e0138321	7.6	1
9	Stem-cell-derived models: tools for studying role of microbiota in intestinal homeostasis and disease. <i>Current Opinion in Gastroenterology</i> , 2021 , 37, 15-22	3	0
8	Intestinal Damage Fades, but Insults Linger: Setting the Immunological Tone for Infection. <i>Cell Host and Microbe</i> , 2019 , 25, 636-637	23.4	
7	A data-driven approach to predict daily risk of <i>Clostridium difficile</i> infection at two large academic health centers. <i>Open Forum Infectious Diseases</i> , 2017 , 4, S403-S404	1	
6	Treatment of <i>Clostridium difficile</i> Infection With Tigecycline. <i>Infectious Diseases in Clinical Practice</i> , 2014 , 22, 216-218	0.2	
5	Blind Spots in Methods Based on Cultivation and Metagenomic Sequencing for Surface Microbiomes in a Medical Intensive Care Unit. <i>Infection Control and Hospital Epidemiology</i> , 2020 , 41, s141 ² s142		

- 4 Recurrent *Clostridioides difficile* infection can be predicted using inflammatory mediator and toxin activity levels. *Infection Control and Hospital Epidemiology*, **2020**, 41, s77-s78 2
- 3 Genomic Epidemiology of *Clostridioides difficile* Sequence Types 1 and 2 Across Three US Medical Centers. *Infection Control and Hospital Epidemiology*, **2020**, 41, s238-s238 2
- 2 496 Baseline and Longitudinal Microbial Changes Predict Response to Rifaximin and/or Diet Low in Fermentable Oligosaccharides, Disaccharides, Monosaccharides, and Polyols in Irritable Bowel Syndrome. *American Journal of Gastroenterology*, **2019**, 114, S289-S289 0.7
- 1 Structure and function of the human microbiome: implications for health and disease **2022**, 2929-2946