

Strains, functions and dynamics in the expanded Huma

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Diving deeper into the communities. <i>Nature Reviews Microbiology</i> , 2017, 15, 643-643.	13.6	0
3	Subspecies in the global human gut microbiome. <i>Molecular Systems Biology</i> , 2017, 13, 960.	3.2	115
4	Microbiome measurement: Possibilities and pitfalls. <i>Bailliere's Best Practice and Research in Clinical Gastroenterology</i> , 2017, 31, 619-623.	1.0	7
5	Systematically investigating the impact of medication on the gut microbiome. <i>Current Opinion in Microbiology</i> , 2017, 39, 128-135.	2.3	65
6	The Human Gut Microbiome: From Association to Modulation. <i>Cell</i> , 2018, 172, 1198-1215.	13.5	558
7	How human microbiome talks to health and disease. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2018, 37, 1595-1601.	1.3	30
8	Deciphering Human Gut Microbiota's Nutrient Interactions: A Role for Biochemistry. <i>Biochemistry</i> , 2018, 57, 2567-2577.	1.2	19
9	No effects without causes: the Iron Dysregulation and Dormant Microbes hypothesis for chronic, inflammatory diseases. <i>Biological Reviews</i> , 2018, 93, 1518-1557.	4.7	92
10	The gut microbiota influences anticancer immunosurveillance and general health. <i>Nature Reviews Clinical Oncology</i> , 2018, 15, 382-396.	12.5	389
11	Exploring Linkages between Taxonomic and Functional Profiles of the Human Microbiome. <i>MSystems</i> , 2018, 3, .	1.7	69
12	A large-scale survey of the postmortem human microbiome, and its potential to provide insight into the living health condition. <i>Scientific Reports</i> , 2018, 8, 5724.	1.6	102
13	Impact of food additives on the gut-brain axis. <i>Physiology and Behavior</i> , 2018, 192, 173-176.	1.0	22
14	Human Microbes - The Power Within. , 2018, , .		6
15	Designer microbiomes for environmental, energy and health biotechnology. <i>Current Opinion in Microbiology</i> , 2018, 43, 117-123.	2.3	16
17	Gut microbiota modify risk for dietary glycemia-induced age-related macular degeneration. <i>Gut Microbes</i> , 2018, 9, 1-6.	4.3	18
18	Microbial diversity knows no borders. <i>Nature Reviews Microbiology</i> , 2018, 16, 66-66.	13.6	13
19	Testing association between soil bacterial diversity and soil carbon storage on the Loess Plateau. <i>Science of the Total Environment</i> , 2018, 626, 48-58.	3.9	53
20	Metatranscriptome of human faecal microbial communities in a cohort of adult men. <i>Nature Microbiology</i> , 2018, 3, 356-366.	5.9	168

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21	Core microbiomes for sustainable agroecosystems. <i>Nature Plants</i> , 2018, 4, 247-257.	4.7	639
22	Determinants of IBD Heritability: Genes, Bugs, and More. <i>Inflammatory Bowel Diseases</i> , 2018, 24, 1133-1148.	0.9	122
23	Dysbiosis in Functional Bowel Disorders. <i>Annals of Nutrition and Metabolism</i> , 2018, 72, 296-306.	1.0	46
24	Bone and the gut microbiome: a new dimension. <i>Journal of Laboratory and Precision Medicine</i> , 0, 3, 96-96.	1.1	11
25	Gut microbiome metagenomics to understand how xenobiotics impact human health. <i>Current Opinion in Toxicology</i> , 2018, 11-12, 51-58.	2.6	17
28	<i>C. elegans</i> and its bacterial diet: An interspecies model to explore the effects of microbiota on drug response. <i>Drug Discovery Today: Disease Models</i> , 2018, 28, 21-26.	1.2	5
29	Gut Microbial and Metabolic Responses to <i>Salmonella enterica</i> Serovar Typhimurium and <i>Candida albicans</i> . <i>MBio</i> , 2018, 9, .	1.8	31
30	Myalgic Encephalomyelitis/Chronic Fatigue Syndrome in the Era of the Human Microbiome: Persistent Pathogens Drive Chronic Symptoms by Interfering With Host Metabolism, Gene Expression, and Immunity. <i>Frontiers in Pediatrics</i> , 2018, 6, 373.	0.9	53
31	Qualitative and Quantitative DNA- and RNA-Based Analysis of the Bacterial Stomach Microbiota in Humans, Mice, and Gerbils. <i>MSystems</i> , 2018, 3, .	1.7	21
32	Dynamic linear models guide design and analysis of microbiota studies within artificial human guts. <i>Microbiome</i> , 2018, 6, 202.	4.9	54
33	Role of gut microbiota in intestinal wound healing and barrier function. <i>Tissue Barriers</i> , 2018, 6, 1539595.	1.6	94
34	Gut Microbiome Dysbiosis and Immunometabolism: New Frontiers for Treatment of Metabolic Diseases. <i>Mediators of Inflammation</i> , 2018, 2018, 1-12.	1.4	199
35	Antimicrobial Chemicals Associate with Microbial Function and Antibiotic Resistance Indoors. <i>MSystems</i> , 2018, 3, .	1.7	63
36	Supragingival Plaque Microbiome Ecology and Functional Potential in the Context of Health and Disease. <i>MBio</i> , 2018, 9, .	1.8	58
37	Practical considerations for sampling and data analysis in contemporary metagenomics-based environmental studies. <i>Journal of Microbiological Methods</i> , 2018, 154, 14-18.	0.7	12
38	Microbial forensics: new breakthroughs and future prospects. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 10377-10391.	1.7	76
39	Clinician Guide to Microbiome Testing. <i>Digestive Diseases and Sciences</i> , 2018, 63, 3167-3177.	1.1	22
40	A study of the correlation between obesity and intestinal flora in school-age children. <i>Scientific Reports</i> , 2018, 8, 14511.	1.6	31

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41	Microbe-derived extracellular vesicles as a smart drug delivery system. <i>Translational and Clinical Pharmacology</i> , 2018, 26, 103.	0.3	37
42	Drug pharmacomicrobiomics and toxicomicrobiomics: from scattered reports to systematic studies of drug-microbiome interactions. <i>Expert Opinion on Drug Metabolism and Toxicology</i> , 2018, 14, 1043-1055.	1.5	36
43	Compositional and Temporal Changes in the Gut Microbiome of Pediatric Ulcerative Colitis Patients Are Linked to Disease Course. <i>Cell Host and Microbe</i> , 2018, 24, 600-610.e4.	5.1	193
44	High-quality genome sequences of uncultured microbes by assembly of read clouds. <i>Nature Biotechnology</i> , 2018, 36, 1067-1075.	9.4	103
45	Modernized Tools for Streamlined Genetic Manipulation and Comparative Study of Wild and Diverse Proteobacterial Lineages. <i>MBio</i> , 2018, 9, .	1.8	65
46	Ecologically informed microbial biomarkers and accurate classification of mixed and unmixed samples in an extensive cross-study of human body sites. <i>Microbiome</i> , 2018, 6, 192.	4.9	25
47	Species-level functional profiling of metagenomes and metatranscriptomes. <i>Nature Methods</i> , 2018, 15, 962-968.	9.0	1,125
48	The metagenome of the female upper reproductive tract. <i>GigaScience</i> , 2018, 7, .	3.3	68
49	High-resolution ISR amplicon sequencing reveals personalized oral microbiome. <i>Microbiome</i> , 2018, 6, 153.	4.9	32
50	Pharmacology in the age of the holobiont. <i>Current Opinion in Systems Biology</i> , 2018, 10, 34-42.	1.3	6
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52	The Oral Microbiome Bank of China. <i>International Journal of Oral Science</i> , 2018, 10, 16.	3.6	29
53	Drivers of Microbiome Biodiversity: A Review of General Rules, Feces, and Ignorance. <i>MBio</i> , 2018, 9, .	1.8	230
54	Characterizing Microbiome Dynamics – Flow Cytometry Based Workflows from Pure Cultures to Natural Communities. <i>Journal of Visualized Experiments</i> , 2018, , .	0.2	12
55	<i>Mycobacterium tuberculosis</i> : An Adaptable Pathogen Associated With Multiple Human Diseases. <i>Frontiers in Cellular and Infection Microbiology</i> , 2018, 8, 158.	1.8	94
56	BURRITO: An Interactive Multi-Omic Tool for Visualizing Taxa-Function Relationships in Microbiome Data. <i>Frontiers in Microbiology</i> , 2018, 9, 365.	1.5	88
57	Low Maternal Microbiota Sharing across Gut, Breast Milk and Vagina, as Revealed by 16S rRNA Gene and Reduced Metagenomic Sequencing. <i>Genes</i> , 2018, 9, 231.	1.0	35
58	Temporal dynamics of the lung and plasma viromes in lung transplant recipients. <i>PLoS ONE</i> , 2018, 13, e0200428.	1.1	23

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59	Distinct Effects of Type I and III Interferons on Enteric Viruses. <i>Viruses</i> , 2018, 10, 46.	1.5	47
60	A selective gut bacterial bile salt hydrolase alters host metabolism. <i>ELife</i> , 2018, 7, .	2.8	177
61	Evaluating the potential of residual Pap test fluid as a resource for the metaproteomic analysis of the cervical-vaginal microbiome. <i>Scientific Reports</i> , 2018, 8, 10868.	1.6	12
62	Mother-to-Infant Microbial Transmission from Different Body Sites Shapes the Developing Infant Gut Microbiome. <i>Cell Host and Microbe</i> , 2018, 24, 133-145.e5.	5.1	822
63	Host genetic variation and its microbiome interactions within the Human Microbiome Project. <i>Genome Medicine</i> , 2018, 10, 6.	3.6	134
64	Profiling microbial strains in urban environments using metagenomic sequencing data. <i>Biology Direct</i> , 2018, 13, 9.	1.9	29
65	Microbial Siderophore Enterobactin Promotes Mitochondrial Iron Uptake and Development of the Host via Interaction with ATP Synthase. <i>Cell</i> , 2018, 175, 571-582.e11.	13.5	124
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67	Implication of gut microbiota metabolites in cardiovascular and metabolic diseases. <i>Cellular and Molecular Life Sciences</i> , 2018, 75, 3977-3990.	2.4	127
68	Gut microbiome and magnetic resonance spectroscopy study of subjects at ultra-high risk for psychosis may support the membrane hypothesis. <i>European Psychiatry</i> , 2018, 53, 37-45.	0.1	88
69	Comprehensive skin microbiome analysis reveals the uniqueness of human skin and evidence for phyllosymbiosis within the class Mammalia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E5786-E5795.	3.3	184
70	Unprecedented Symbiont Eukaryote Diversity Is Governed by Internal Trophic Webs in a Wild Non-Human Primate. <i>Protist</i> , 2018, 169, 307-320.	0.6	11
71	Large-Scale Analyses of Human Microbiomes Reveal Thousands of Small, Novel Genes. <i>Cell</i> , 2019, 178, 1245-1259.e14.	13.5	163
72	Full annotation of serum virome in Chinese blood donors with elevated alanine aminotransferase levels. <i>Transfusion</i> , 2019, 59, 3177-3185.	0.8	14
73	Massive metagenomic data analysis using abundance-based machine learning. <i>Biology Direct</i> , 2019, 14, 12.	1.9	29
74	Gut Microbiome Biomarkers and Functional Diversity Within an Amazonian Semi-Nomadic Hunter-Gatherer Group. <i>Frontiers in Microbiology</i> , 2019, 10, 1743.	1.5	32
75	The core microbiome of sessile ciliate <i>Stentor coeruleus</i> is not shaped by the environment. <i>Scientific Reports</i> , 2019, 9, 11356.	1.6	16
76	The Landscape of Genetic Content in the Gut and Oral Human Microbiome. <i>Cell Host and Microbe</i> , 2019, 26, 283-295.e8.	5.1	207

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78	Microbes: An Important Resource for Sustainable Agriculture. , 2019, , 53-77.		2
79	Early-life gut microbiome modulation reduces the abundance of antibiotic-resistant bacteria. <i>Antimicrobial Resistance and Infection Control</i> , 2019, 8, 131.	1.5	63
80	Tumor Microbiome Diversity and Composition Influence Pancreatic Cancer Outcomes. <i>Cell</i> , 2019, 178, 795-806.e12.	13.5	830
81	Diet-derived microbial metabolites in health and disease. <i>Nutrition Bulletin</i> , 2019, 44, 216-227.	0.8	36
82	Metabolomics for Investigating Physiological and Pathophysiological Processes. <i>Physiological Reviews</i> , 2019, 99, 1819-1875.	13.1	516
83	An integrated respiratory microbial gene catalogue to better understand the microbial aetiology of <i>Mycoplasma pneumoniae</i> pneumonia. <i>GigaScience</i> , 2019, 8, .	3.3	16
84	Consumer Safety Considerations of Skin and Oral Microbiome Perturbation. <i>Clinical Microbiology Reviews</i> , 2019, 32, .	5.7	15
85	Population Genetics of Host-Associated Microbiomes. <i>Current Molecular Biology Reports</i> , 2019, 5, 128-139.	0.8	10
86	Healthspan and lifespan extension by fecal microbiota transplantation into progeroid mice. <i>Nature Medicine</i> , 2019, 25, 1234-1242.	15.2	352
87	Quantifying spatiotemporal variability and noise in absolute microbiota abundances using replicate sampling. <i>Nature Methods</i> , 2019, 16, 731-736.	9.0	54
89	Argonaute proteins from human gastrointestinal bacteria catalyze DNA-guided cleavage of single- and double-stranded DNA at 37°C. <i>Cell Discovery</i> , 2019, 5, 38.	3.1	47
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91	Effects of Fish n-3 PUFAs on Intestinal Microbiota and Immune System. <i>Marine Drugs</i> , 2019, 17, 374.	2.2	105
92	Microbiome Is a Functional Modifier of P450 Drug Metabolism. <i>Current Pharmacology Reports</i> , 2019, 5, 481-490.	1.5	34
93	MetaMiner: A Scalable Peptidogenomics Approach for Discovery of Ribosomal Peptide Natural Products with Blind Modifications from Microbial Communities. <i>Cell Systems</i> , 2019, 9, 600-608.e4.	2.9	46
94	The Human Gut Virome Is Highly Diverse, Stable, and Individual Specific. <i>Cell Host and Microbe</i> , 2019, 26, 527-541.e5.	5.1	449
95	An assessment of Oxford Nanopore sequencing for human gut metagenome profiling: A pilot study of head and neck cancer patients. <i>Journal of Microbiological Methods</i> , 2019, 166, 105739.	0.7	13

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96	Emerging Frontiers in Microbiome Engineering. Trends in Immunology, 2019, 40, 952-973.	2.9	47
97	Coarse Grained Heat-Affected Zone Microstructure and Brittleness of Ti-Nb-B Microalloyed High Toughness and Wear Resistant Steel. Metals, 2019, 9, 289.	1.0	3
98	Human gut bacteria contain acquired interbacterial defence systems. Nature, 2019, 575, 224-228.	13.7	99
99	Graph Embedding Deep Learning Guides Microbial Biomarkers' Identification. Frontiers in Genetics, 2019, 10, 1182.	1.1	21
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106	Exact variance component tests for longitudinal microbiome studies. Genetic Epidemiology, 2019, 43, 250-262.	0.6	5
107	Challenges in Clinical Metaproteomics Highlighted by the Analysis of Acute Leukemia Patients with Gut Colonization by Multidrug-Resistant Enterobacteriaceae. Proteomes, 2019, 7, 2.	1.7	71
108	Making the microbiome public: Participatory experiments with DNA sequencing in domestic kitchens. Transactions of the Institute of British Geographers, 2019, 44, 524-541.	1.8	16
109	The Microbiota-Gut-Brain Axis. Physiological Reviews, 2019, 99, 1877-2013.	13.1	2,304
110	Host-Microbe-Drug-Nutrient Screen Identifies Bacterial Effectors of Metformin Therapy. Cell, 2019, 178, 1299-1312.e29.	13.5	186
111	Pursuing Human-Relevant Gut Microbiota-Immune Interactions. Immunity, 2019, 51, 225-239.	6.6	105
112	Inevitable future: space colonization beyond Earth with microbes first. FEMS Microbiology Ecology, 2019, 95, .	1.3	22
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115	Baseline human gut microbiota profile in healthy people and standard reporting template. <i>PLoS ONE</i> , 2019, 14, e0206484.	1.1	133
116	Priorities for the next 10 years of human microbiome research. <i>Nature</i> , 2019, 569, 623-625.	13.7	99
117	Homeostasis and dysbiosis of the gut microbiome in health and disease. <i>Journal of Biosciences</i> , 2019, 44, 1.	0.5	107
118	A comprehensive assessment of demographic, environmental, and host genetic associations with gut microbiome diversity in healthy individuals. <i>Microbiome</i> , 2019, 7, 130.	4.9	101
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120	Interplay between the human gut microbiome and host metabolism. <i>Nature Communications</i> , 2019, 10, 4505.	5.8	450
121	A metagenomic strategy for harnessing the chemical repertoire of the human microbiome. <i>Science</i> , 2019, 366, .	6.0	101
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123	Comparative genomics of Bacteria commonly identified in the built environment. <i>BMC Genomics</i> , 2019, 20, 92.	1.2	6
124	Composition and Distribution Analysis of Bioaerosols Under Different Environmental Conditions. <i>Journal of Visualized Experiments</i> , 2019, , .	0.2	5
125	A defined commensal consortium elicits CD8 T cells and anti-cancer immunity. <i>Nature</i> , 2019, 565, 600-605.	13.7	741
126	Evolutionary dynamics of bacteria in the gut microbiome within and across hosts. <i>PLoS Biology</i> , 2019, 17, e3000102.	2.6	257
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128	Fecal Microbial Transplantation and Its Potential Application in Cardiometabolic Syndrome. <i>Frontiers in Immunology</i> , 2019, 10, 1341.	2.2	63
129	A multidimensional perspective on microbial interactions. <i>FEMS Microbiology Letters</i> , 2019, 366, .	0.7	58
130	Low-Pressure Plasma Sterilization for Test Specimens to be Worn on Splints in the Oral Cavity. <i>Coatings</i> , 2019, 9, 99.	1.2	3
131	Intestinal microbiome and fitness in kidney disease. <i>Nature Reviews Nephrology</i> , 2019, 15, 531-545.	4.1	140

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132	Comparative genomics and genome biology of <i>Campylobacter showae</i> . <i>Emerging Microbes and Infections</i> , 2019, 8, 827-840.	3.0	8
133	Relationship between faecal microbiota and plasma metabolome in rats fed NK603 and MON810 GM maize from the GMO90+ study. <i>Food and Chemical Toxicology</i> , 2019, 131, 110547.	1.8	7
134	Impact of Host DNA and Sequencing Depth on the Taxonomic Resolution of Whole Metagenome Sequencing for Microbiome Analysis. <i>Frontiers in Microbiology</i> , 2019, 10, 1277.	1.5	137
135	Multi-omics of the gut microbial ecosystem in inflammatory bowel diseases. <i>Nature</i> , 2019, 569, 655-662.	13.7	1,638
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139	Biogeography of the Oral Microbiome: The Site-Specialist Hypothesis. <i>Annual Review of Microbiology</i> , 2019, 73, 335-358.	2.9	147
140	A Prospective Study of the Urinary and Gastrointestinal Microbiome in Prepubertal Males. <i>Urology</i> , 2019, 131, 204-210.	0.5	26
141	Gut microbiota in ALS: possible role in pathogenesis?. <i>Expert Review of Neurotherapeutics</i> , 2019, 19, 785-805.	1.4	30
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144	Applications of high-throughput –omics™ data in the study of frailty. <i>Translational Medicine of Aging</i> , 2019, 3, 40-51.	0.6	5
145	The gut flora modulates intestinal barrier integrity but not progression of chronic kidney disease in hyperoxaluria-related nephrocalcinosis. <i>Nephrology Dialysis Transplantation</i> , 2019, 35, 86-97.	0.4	9
146	Age- and Sex-Dependent Patterns of Gut Microbial Diversity in Human Adults. <i>MSystems</i> , 2019, 4, .	1.7	214
147	International Cancer Microbiome Consortium consensus statement on the role of the human microbiome in carcinogenesis. <i>Gut</i> , 2019, 68, 1624-1632.	6.1	173
148	<i>Helicobacter pylori</i> Infection, the Gastric Microbiome and Gastric Cancer. <i>Advances in Experimental Medicine and Biology</i> , 2019, 1149, 195-210.	0.8	49
149	Adaptive Evolution within Gut Microbiomes of Healthy People. <i>Cell Host and Microbe</i> , 2019, 25, 656-667.e8.	5.1	289

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151	Probiotic <i>Bifidobacterium lactis</i> V9 Regulates the Secretion of Sex Hormones in Polycystic Ovary Syndrome Patients through the Gut-Brain Axis. <i>MSystems</i> , 2019, 4, .	1.7	157
152	Beyond PD-L1 Markers for Lung Cancer Immunotherapy. <i>International Journal of Molecular Sciences</i> , 2019, 20, 1915.	1.8	61
153	Inverse Association Between the Skin and Oral Microbiota in Atopic Dermatitis. <i>Journal of Investigative Dermatology</i> , 2019, 139, 1779-1787.e12.	0.3	31
154	Xenobiotics Formed during Food Processing: Their Relation with the Intestinal Microbiota and Colorectal Cancer. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2051.	1.8	53
155	Characterization of microbial community profiles associated with quality of Chinese strong aromatic liquor through metagenomics. <i>Journal of Applied Microbiology</i> , 2019, 127, 750-762.	1.4	6
156	Characterization of Bacterial Communities in Breeding Waters of <i>Anopheles darlingi</i> in Manaus in the Amazon Basin Malaria-Endemic Area. <i>Microbial Ecology</i> , 2019, 78, 781-791.	1.4	21
157	Drinking Water Microbiome Project: Is it Time?. <i>Trends in Microbiology</i> , 2019, 27, 670-677.	3.5	50
158	Molecular Heterogeneity in Large-Scale Biological Data: Techniques and Applications. <i>Annual Review of Biomedical Data Science</i> , 2019, 2, 39-67.	2.8	3
159	Microbiome diurnal rhythmicity and its impact on host physiology and disease risk. <i>EMBO Reports</i> , 2019, 20, .	2.0	66
160	The Gut Microbiome and Ankylosing Spondylitis. , 2019, , 87-95.		1
161	Microbial abundance, activity and population genomic profiling with mOTUs2. <i>Nature Communications</i> , 2019, 10, 1014.	5.8	298
162	A review of 10 years of human microbiome research activities at the US National Institutes of Health, Fiscal Years 2007-2016. <i>Microbiome</i> , 2019, 7, 31.	4.9	155
163	Microbiome-MX 2018: microbiota and microbiome opportunities in Mexico, a megadiverse country. <i>Research in Microbiology</i> , 2019, 170, 235-241.	1.0	2
164	The oral microbiome of patients undergoing treatment for severe aplastic anemia: a pilot study. <i>Annals of Hematology</i> , 2019, 98, 1351-1365.	0.8	20
165	Mining the microbiota for microbial and metabolite-based immunotherapies. <i>Nature Reviews Immunology</i> , 2019, 19, 305-323.	10.6	211
166	Microbiome, Parkinson's Disease and Molecular Mimicry. <i>Cells</i> , 2019, 8, 222.	1.8	56
167	2017 NIH-wide workshop report on "The Human Microbiome: Emerging Themes at the Horizon of the 21st Century". <i>Microbiome</i> , 2019, 7, 32.	4.9	6

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168	Distinct Genetic and Functional Traits of Human Intestinal <i>Prevotella copri</i> Strains Are Associated with Different Habitual Diets. <i>Cell Host and Microbe</i> , 2019, 25, 444-453.e3.	5.1	229
169	Microbiome and Cancers, With Focus on Genitourinary Tumors. <i>Frontiers in Oncology</i> , 2019, 9, 178.	1.3	20
170	Horizontal gene transfer in human-associated microorganisms inferred by phylogenetic reconstruction and reconciliation. <i>Scientific Reports</i> , 2019, 9, 5953.	1.6	55
171	Culture-enriched human gut microbiomes reveal core and accessory resistance genes. <i>Microbiome</i> , 2019, 7, 56.	4.9	23
172	Human and Extracellular DNA Depletion for Metagenomic Analysis of Complex Clinical Infection Samples Yields Optimized Viable Microbiome Profiles. <i>Cell Reports</i> , 2019, 26, 2227-2240.e5.	2.9	104
173	The human gut Firmicute <i>Roseburia intestinalis</i> is a primary degrader of dietary β -mannans. <i>Nature Communications</i> , 2019, 10, 905.	5.8	202
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