David Berry

List of Publications by Year in descending order

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DAVID REDOV

#	Article	IF	CITATIONS
1	Deciphering microbial interactions and detecting keystone species with co-occurrence networks. Frontiers in Microbiology, 2014, 5, 219.	1.5	1,109
2	High-fat diet alters gut microbiota physiology in mice. ISME Journal, 2014, 8, 295-308.	4.4	583
3	Barcoded Primers Used in Multiplex Amplicon Pyrosequencing Bias Amplification. Applied and Environmental Microbiology, 2011, 77, 7846-7849.	1.4	514
4	Microbial ecology of drinking water distribution systems. Current Opinion in Biotechnology, 2006, 17, 297-302.	3.3	372
5	Tracking heavy water (D ₂ O) incorporation for identifying and sorting active microbial cells. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E194-203.	3.3	359
6	Pediatric obesity is associated with an altered gut microbiota and discordant shifts in <scp><i>F</i></scp> <i>irmicutes</i> populations. Environmental Microbiology, 2017, 19, 95-105.	1.8	326
7	Genome-guided design of a defined mouse microbiota that confers colonization resistance against Salmonella enterica serovar Typhimurium. Nature Microbiology, 2017, 2, 16215.	5.9	313
8	Temporal Bacterial Community Dynamics Vary Among Ulcerative Colitis Patients After Fecal Microbiota Transplantation. American Journal of Gastroenterology, 2013, 108, 1620-1630.	0.2	298
9	Phylotype-level 16S rRNA analysis reveals new bacterial indicators of health state in acute murine colitis. ISME Journal, 2012, 6, 2091-2106.	4.4	291
10	<scp><i>NxrB</i></scp> encoding the beta subunit of nitrite oxidoreductase as functional and phylogenetic marker for nitriteâ€oxidizing <scp><i>N</i></scp> <i>itrospira</i> . Environmental Microbiology, 2014, 16, 3055-3071.	1.8	280
11	Hydrocarbon-degrading bacteria enriched by the <i>Deepwater Horizon</i> oil spill identified by cultivation and DNA-SIP. ISME Journal, 2013, 7, 2091-2104.	4.4	278
12	Microbial nutrient niches in the gut. Environmental Microbiology, 2017, 19, 1366-1378.	1.8	258
13	Cyanate as an energy source for nitrifiers. Nature, 2015, 524, 105-108.	13.7	231
14	Host-compound foraging by intestinal microbiota revealed by single-cell stable isotope probing. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 4720-4725.	3.3	210
15	Mucispirillum schaedleri Antagonizes Salmonella Virulence to Protect Mice against Colitis. Cell Host and Microbe, 2019, 25, 681-694.e8.	5.1	205
16	Rational design of a microbial consortium of mucosal sugar utilizers reduces Clostridiodes difficile colonization. Nature Communications, 2020, 11, 5104.	5.8	177
17	Longitudinal study of murine microbiota activity and interactions with the host during acute inflammation and recovery. ISME Journal, 2014, 8, 1101-1114.	4.4	174
18	An automated Raman-based platform for the sorting of live cells by functional properties. Nature Microbiology, 2019, 4, 1035-1048.	5.9	170

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19	A flexible and economical barcoding approach for highly multiplexed amplicon sequencing of diverse target genes. Frontiers in Microbiology, 2015, 6, 731.	1.5	164
20	Lifestyle and Horizontal Gene Transfer-Mediated Evolution of Mucispirillum schaedleri, a Core Member of the Murine Gut Microbiota. MSystems, 2017, 2, .	1.7	148
21	Barcoded Primers Used in Multiplex Amplicon Pyrosequencing Bias Amplification. Applied and Environmental Microbiology, 2012, 78, 612-612.	1.4	146
22	Endospores of thermophilic bacteria as tracers of microbial dispersal by ocean currents. ISME Journal, 2014, 8, 1153-1165.	4.4	139
23	Role of Bacterial Exopolysaccharides (EPS) in the Fate of the Oil Released during the Deepwater Horizon Oil Spill. PLoS ONE, 2013, 8, e67717.	1.1	135
24	Intestinal microbiota: A source of novel biomarkers in inflammatory bowel diseases?. Bailliere's Best Practice and Research in Clinical Gastroenterology, 2013, 27, 47-58.	1.0	127
25	Microbial nitrogen limitation in the mammalian large intestine. Nature Microbiology, 2018, 3, 1441-1450.	5.9	107
26	Intestinal Microbiota Signatures Associated with Inflammation History in Mice Experiencing Recurring Colitis. Frontiers in Microbiology, 2015, 6, 1408.	1.5	106
27	Long-distance electron transport in individual, living cable bacteria. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 5786-5791.	3.3	104
28	Transkingdom interactions between Lactobacilli and hepatic mitochondria attenuate western diet-induced diabetes. Nature Communications, 2021, 12, 101.	5.8	86
29	Colonization resistance and microbial ecophysiology: using gnotobiotic mouse models and single-cell technology to explore the intestinal jungle. FEMS Microbiology Reviews, 2013, 37, 793-829.	3.9	85
30	Activity and community structures of sulfate-reducing microorganisms in polar, temperate and tropical marine sediments. ISME Journal, 2016, 10, 796-809.	4.4	85
31	A fiber-deprived diet disturbs the fine-scale spatial architecture of the murine colon microbiome. Nature Communications, 2019, 10, 4366.	5.8	82
32	Polycyclic Aromatic Hydrocarbon Degradation of Phytoplankton-Associated Arenibacter spp. and Description of Arenibacter algicola sp. nov., an Aromatic Hydrocarbon-Degrading Bacterium. Applied and Environmental Microbiology, 2014, 80, 618-628.	1.4	81
33	Aberrant gut-microbiota-immune-brain axis development in premature neonates with brain damage. Cell Host and Microbe, 2021, 29, 1558-1572.e6.	5.1	80
34	Impact of microfiltration treatment of secondary wastewater effluent on biofouling of reverse osmosis membranes. Water Research, 2010, 44, 167-176.	5.3	76
35	HuR Small-Molecule Inhibitor Elicits Differential Effects in Adenomatosis Polyposis and Colorectal Carcinogenesis. Cancer Research, 2017, 77, 2424-2438.	0.4	75
36	Antioxidative activity and health benefits of anthocyanin-rich fruit juice in healthy volunteers. Free Radical Research, 2019, 53, 1045-1055.	1.5	74

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37	Intestinal Bacteria Modify Lymphoma Incidence and Latency by Affecting Systemic Inflammatory State, Oxidative Stress, and Leukocyte Genotoxicity. Cancer Research, 2013, 73, 4222-4232.	0.4	68
38	Response of the bacterial community associated with a cosmopolitan marine diatom to crude oil shows a preference for the biodegradation of aromatic hydrocarbons. Environmental Microbiology, 2016, 18, 1817-1833.	1.8	68
39	Development of reactive thin film polymer brush membranes to prevent biofouling. Journal of Membrane Science, 2010, 350, 361-370.	4.1	67
40	Handling of spurious sequences affects the outcome of high-throughput 16S rRNA gene amplicon profiling. ISME Communications, 2021, 1, .	1.7	60
41	Stable-Isotope Probing of Human and Animal Microbiome Function. Trends in Microbiology, 2018, 26, 999-1007.	3.5	57
42	Raman microspectroscopy for microbiology. Nature Reviews Methods Primers, 2021, 1, .	11.8	57
43	Mucosal Biofilms Are an Endoscopic Feature of Irritable Bowel Syndrome and Ulcerative Colitis. Gastroenterology, 2021, 161, 1245-1256.e20.	0.6	55
44	Vibrational Spectroscopy for Imaging Single Microbial Cells in Complex Biological Samples. Frontiers in Microbiology, 2017, 8, 675.	1.5	51
45	Design and performance of a single-pass bubbling bioaerosol generator. Atmospheric Environment, 2005, 39, 3521-3533.	1.9	49
46	An Economical and Flexible Dual Barcoding, Two-Step PCR Approach for Highly Multiplexed Amplicon Sequencing. Frontiers in Microbiology, 2021, 12, 669776.	1.5	48
47	Conversion of Rutin, a Prevalent Dietary Flavonol, by the Human Gut Microbiota. Frontiers in Microbiology, 2020, 11, 585428.	1.5	47
48	Optofluidic Raman-activated cell sorting for targeted genome retrieval or cultivation of microbial cells with specific functions. Nature Protocols, 2021, 16, 634-676.	5.5	41
49	Intestinal Epithelial Cell Tyrosine Kinase 2 Transduces IL-22 Signals To Protect from Acute Colitis. Journal of Immunology, 2015, 195, 5011-5024.	0.4	40
50	Type I interferons have opposing effects during the emergence and recovery phases of colitis. European Journal of Immunology, 2014, 44, 2749-2760.	1.6	39
51	Removal of Pharmaceuticals and Personal Care Products during Water Recycling: Microbial Community Structure and Effects of Substrate Concentration. Applied and Environmental Microbiology, 2014, 80, 2440-2450.	1.4	37
52	Fluorinated Gold Nanoparticles for Nanostructure Imaging Mass Spectrometry. ACS Nano, 2018, 12, 6938-6948.	7.3	37
53	Transparent soil microcosms for live-cell imaging and non-destructive stable isotope probing of soil microorganisms. ELife, 2020, 9, .	2.8	36
54	The emerging view of <scp><i>F</i></scp> <i>irmicutes</i> as key fibre degraders in the human gut. Environmental Microbiology, 2016, 18, 2081-2083.	1.8	35

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55	A 12â€week intervention with nonivamide, a TRPV1 agonist, prevents a dietaryâ€induced body fat gain and increases peripheral serotonin in moderately overweight subjects. Molecular Nutrition and Food Research, 2017, 61, 1600731.	1.5	31
56	Elucidating the role of the gut microbiota in the physiological effects of dietary fiber. Microbiome, 2022, 10, 77.	4.9	31
57	<i>Mycobacterium avium</i> Infections of <i>Acanthamoeba</i> Strains: Host Strain Variability, Grazing-Acquired Infections, and Altered Dynamics of Inactivation with Monochloramine. Applied and Environmental Microbiology, 2010, 76, 6685-6688.	1.4	29
58	Vitamin and Amino Acid Auxotrophy in Anaerobic Consortia Operating under Methanogenic Conditions. MSystems, 2017, 2, .	1.7	28
59	A dynamic and complex monochloramine stress response in Escherichia coli revealed by transcriptome analysis. Water Research, 2013, 47, 4978-4985.	5.3	26
60	Intestinal Microbiota Reduces Genotoxic Endpoints Induced By High-Energy Protons. Radiation Research, 2014, 181, 45-53.	0.7	26
61	Bottled aqua incognita: microbiota assembly and dissolved organic matter diversity in natural mineral waters. Microbiome, 2017, 5, 126.	4.9	26
62	Bacterial nutrient foraging in a mouse model of enteral nutrient deprivation: insight into the gut origin of sepsis. American Journal of Physiology - Renal Physiology, 2016, 311, G734-G743.	1.6	25
63	Evaluating the Detection of Hydrocarbon-Degrading Bacteria in 16S rRNA Gene Sequencing Surveys. Frontiers in Microbiology, 2017, 8, 896.	1.5	25
64	Effect of an Ionic Air Cleaner on Indoor/Outdoor Particle Ratios in a Residential Environment. Aerosol Science and Technology, 2007, 41, 315-328.	1.5	24
65	Allspice and Clove As Source of Triterpene Acids Activating the G Protein-Coupled Bile Acid Receptor TGR5. Frontiers in Pharmacology, 2017, 8, 468.	1.6	24
66	Effect of Growth Conditions on Inactivation of <i>Escherichia coli</i> with Monochloramine. Environmental Science & Technology, 2009, 43, 884-889.	4.6	23
67	Enrichment of Fusobacteria in Sea Surface Oil Slicks from the Deepwater Horizon Oil Spill. Microorganisms, 2016, 4, 24.	1.6	23
68	Next-generation biomonitoring of the early-life chemical exposome in neonatal and infant development. Nature Communications, 2022, 13, 2653.	5.8	23
69	Berry-Enriched Diet in Salt-Sensitive Hypertensive Rats: Metabolic Fate of (Poly)Phenols and the Role of Gut Microbiota. Nutrients, 2019, 11, 2634.	1.7	22
70	Crypt residing bacteria and proximal colonic carcinogenesis in a mouse model of Lynch syndrome. International Journal of Cancer, 2020, 147, 2316-2326.	2.3	20
71	Combined hormonal contraceptives are associated with minor changes in composition and diversity in gut microbiota of healthy women. Environmental Microbiology, 2021, 23, 3037-3047.	1.8	20
72	Comparative transcriptomics of the response of Escherichia coli to the disinfectant monochloramine and to growth conditions inducing monochloramine resistance. Water Research, 2010, 44, 4924-4931.	5.3	19

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73	Systematic Spatial Bias in DNA Microarray Hybridization Is Caused by Probe Spot Position-Dependent Variability in Lateral Diffusion. PLoS ONE, 2011, 6, e23727.	1.1	18
74	Editorial: Bifidobacteria and Their Role in the Human Gut Microbiota. Frontiers in Microbiology, 2016, 7, 2148.	1.5	17
75	Members of the Oral Microbiota Are Associated with IL-8 Release by Gingival Epithelial Cells in Healthy Individuals. Frontiers in Microbiology, 2017, 08, 416.	1.5	17
76	Polyphenol Exposure, Metabolism, and Analysis: A Global Exposomics Perspective. Annual Review of Food Science and Technology, 2021, 12, 461-484.	5.1	17
77	Differential Modulation of the European Sea Bass Gut Microbiota by Distinct Insect Meals. Frontiers in Microbiology, 2022, 13, 831034.	1.5	17
78	SRS-FISH: A high-throughput platform linking microbiome metabolism to identity at the single-cell level. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	3.3	17
79	The role of gut microbiota, butyrate and proton pump inhibitors in amyotrophic lateral sclerosis: a systematic review. International Journal of Neuroscience, 2020, 130, 727-735.	0.8	14
80	Gut microbiota and undigested food constituents modify toxin composition and suppress the genotoxicity of a naturally occurring mixture of Alternaria toxins in vitro. Archives of Toxicology, 2020, 94, 3541-3552.	1.9	13
81	Early-life chemical exposome and gut microbiome development: African research perspectives within a global environmental health context. Trends in Microbiology, 2022, 30, 1084-1100.	3.5	13
82	In vitro interactions of Alternaria mycotoxins, an emerging class of food contaminants, with the gut microbiota: a bidirectional relationship. Archives of Toxicology, 2021, 95, 2533-2549.	1.9	12
83	Long-Term Consumption of Anthocyanin-Rich Fruit Juice: Impact on Gut Microbiota and Antioxidant Markers in Lymphocytes of Healthy Males. Antioxidants, 2021, 10, 27.	2.2	11
84	Anaerobic Sulfur Oxidation Underlies Adaptation of a Chemosynthetic Symbiont to Oxic-Anoxic Interfaces. MSystems, 2021, 6, e0118620.	1.7	10
85	Ecological Processes Shaping Microbiomes of Extremely Low Birthweight Infants. Frontiers in Microbiology, 2022, 13, 812136.	1.5	5
86	Making It Stick: A Compelling Case for Precision Microbiome Reconstitution. Cell Host and Microbe, 2016, 20, 415-417.	5.1	4
87	Hidden potential: dietâ€driven changes in redox level shape the rumen microbiome. Environmental Microbiology, 2017, 19, 19-20.	1.8	4
88	Gilbert's Syndrome and the Gut Microbiota – Insights From the Case-Control BILIHEALTH Study. Frontiers in Cellular and Infection Microbiology, 2021, 11, 701109.	1.8	4
89	Persistence of the antagonistic effects of a natural mixture of Alternaria mycotoxins on the estrogen-like activity of human feces after anaerobic incubation. Toxicology Letters, 2022, 358, 88-99.	0.4	4
90	Targeting Gut Bacteria Using Inulin onjugated Mesoporous Silica Nanoparticles. Advanced Materials Interfaces, 0, , 2102558.	1.9	4

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91	Individuality of the Extremely Premature Infant Gut Microbiota Is Driven by Ecological Drift. MSystems, 2022, 7, e0016322.	1.7	4
92	Reduced alpha diversity of the oral microbiome correlates with short progressionâ€free survival in patients with relapsed/refractory multiple myeloma treated with ixazomibâ€based therapy (AGMT MM 1,) Tj ETQc	00040 rgB]	Øverlock 1
93	The unexpected versatility of the cellulosome. Environmental Microbiology, 2017, 19, 13-14.	1.8	2
94	Spotlight on how microbes influence their host's behavior. Environmental Microbiology, 2019, 21, 3185-3187.	1.8	2

95	Lipid synthesis at the trophic base as the source for energy management to build complex structures. Current Opinion in Biotechnology, 2022, 73, 364-373.	3.3	1	

Targeting Gut Bacteria Using Inulinâ \in Conjugated Mesoporous Silica Nanoparticles (Adv. Mater.) Tj ETQq0 0 0 rgBT $_{1.9}^{1/9}$ verlock 10 Tf 50 5 $_{1.9}^{1/9}$ verlock 10 96

97	Individual Sweet Taste Perception Influences Salivary Characteristics After Orosensory Stimulation With Sucrose and Noncaloric Sweeteners. Frontiers in Nutrition, 2022, 9, .	1.6	1
98	760 Bacterial Translocation Into the Mucus of Crypts Is Associated With Proximal Colonic Tumorigenesis in IL-10â^'/â^'xMSH2loxP/loxP Vill-cre (DKO) Mice. Gastroenterology, 2016, 150, S154.	0.6	0
99	Upâ€closeâ€andâ€personal with the human microbiome. Environmental Microbiology Reports, 2019, 11, 17-19.	1.0	0
100	STILLLEBEN with Symbionts. Performance Research, 2020, 25, 83-87.	0.2	0
101	A Mixed-Lipid Emulsion Containing Fish Oil for the Parenteral Nutrition of Preterm Infants: No Impact on Visual Neuronal Conduction. Nutrients, 2021, 13, 4241.	1.7	0