

James Butcher

List of Publications by Year in descending order

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41
papers

2,139
citations

331670

21
h-index

315739

38
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43
all docs

43
docs citations

43
times ranked

3157
citing authors

#	ARTICLE	IF	CITATIONS
1	Maternal Diet and Infant Feeding Practices Are Associated with Variation in the Human Milk Microbiota at 3 Months Postpartum in a Cohort of Women with High Rates of Gestational Glucose Intolerance. <i>Journal of Nutrition</i> , 2021, 151, 320-329.	2.9	24
2	Examining the Effects of an Anti-Salmonella Bacteriophage Preparation, BAFASALÂ®, on Ex-Vivo Human Gut Microbiome Composition and Function Using a Multi-Omics Approach. <i>Viruses</i> , 2021, 13, 1734.	3.3	5
3	Oligosaccharides and Microbiota in Human Milk Are Interrelated at 3 Months Postpartum in a Cohort of Women with a High Prevalence of Gestational Impaired Glucose Tolerance. <i>Journal of Nutrition</i> , 2021, 151, 3431-3441.	2.9	10
4	Reduced Infection Efficiency of Phage NCTC 12673 on Non-Motile <i>Campylobacter jejuni</i> Strains Is Related to Oxidative Stress. <i>Viruses</i> , 2021, 13, 1955.	3.3	4
5	The gastrointestinal pathogen <i>Campylobacter jejuni</i> metabolizes sugars with potential help from commensal <i>Bacteroides vulgatus</i> . <i>Communications Biology</i> , 2020, 3, 2.	4.4	26
6	Characterization of gastrointestinal pathologies in the dystonia musculorum mouse model for hereditary sensory and autonomic neuropathy type VI. <i>Neurogastroenterology and Motility</i> , 2020, 32, e13773.	3.0	0
7	Examining the relationship between maternal body size, gestational glucose tolerance status, mode of delivery and ethnicity on human milk microbiota at three months post-partum. <i>BMC Microbiology</i> , 2020, 20, 219.	3.3	20
8	Virome Sequencing of the Human Intestinal Mucosalâ€“Luminal Interface. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020, 10, 582187.	3.9	14
9	Mothers of Preterm Infants Have Individualized Breast Milk Microbiota that Changes Temporally Based on Maternal Characteristics. <i>Cell Host and Microbe</i> , 2020, 28, 669-682.e4.	11.0	31
10	Bovine Lactoferrin Supplementation Does Not Disrupt Microbiota Development in Preterm Infants Receiving Probiotics. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2020, 71, 216-222.	1.8	5
11	Binding of Phage-Encoded FlaGrab to Motile <i>Campylobacter jejuni</i> Flagella Inhibits Growth, Downregulates Energy Metabolism, and Requires Specific Flagellar Glycans. <i>Frontiers in Microbiology</i> , 2020, 11, 397.	3.5	14
12	The impact of probiotics and lactoferrin supplementation on piglet gastrointestinal microbial communities. <i>BioMetals</i> , 2019, 32, 533-543.	4.1	18
13	The mucosalâ€“luminal interface: an ideal sample to study the mucosa-associated microbiota and the intestinal microbial biogeography. <i>Pediatric Research</i> , 2019, 85, 895-903.	2.3	32
14	Advancing functional and translational microbiome research using meta-omics approaches. <i>Microbiome</i> , 2019, 7, 154.	11.1	177
15	Independent of Birth Mode or Gestational Age, Very-Low-Birth-Weight Infants Fed Their Mothers' Milk Rapidly Develop Personalized Microbiotas Low in <i>Bifidobacterium</i> . <i>Journal of Nutrition</i> , 2018, 148, 326-335.	2.9	22
16	Evaluating in Vitro Culture Medium of Gut Microbiome with Orthogonal Experimental Design and a Metaproteomics Approach. <i>Journal of Proteome Research</i> , 2018, 17, 154-163.	3.7	41
17	Methods and Strategies to Examine the Human Breastmilk Microbiome. <i>Methods in Molecular Biology</i> , 2018, 1849, 63-86.	0.9	15
18	Crystal structure of <i>Campylobacter jejuni</i> peroxide regulator. <i>FEBS Letters</i> , 2018, 592, 2351-2360.	2.8	6

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19	Variation on a theme: investigating the structural repertoires used by ferric uptake regulators to control gene expression. <i>BioMetals</i> , 2018, 31, 681-704.	4.1	27
20	Transcriptomic Analysis of the <i>Campylobacter jejuni</i> Response to T4-Like Phage NCTC 12673 Infection. <i>Viruses</i> , 2018, 10, 332.	3.3	46
21	Disruption of maternal gut microbiota during gestation alters offspring microbiota and immunity. <i>Microbiome</i> , 2018, 6, 124.	11.1	109
22	Metaproteomics reveals associations between microbiome and intestinal extracellular vesicle proteins in pediatric inflammatory bowel disease. <i>Nature Communications</i> , 2018, 9, 2873.	12.8	209
23	Functional insights into the interplay between DNA interaction and metal coordination in ferric uptake regulators. <i>Scientific Reports</i> , 2018, 8, 7140.	3.3	13
24	NuA4 Lysine Acetyltransferase Complex Contributes to Phospholipid Homeostasis in <i>Saccharomyces cerevisiae</i> . <i>G3: Genes, Genomes, Genetics</i> , 2017, 7, 1799-1809.	1.8	7
25	Analyzing Prokaryotic RNA-Seq Data: A Case Study Identifying Holo-Fur Regulated Genes in <i>Campylobacter jejuni</i> . <i>Methods in Molecular Biology</i> , 2017, 1512, 245-256.	0.9	0
26	Mucosa-Associated Ileal Microbiota in New-Onset Pediatric Crohn's Disease. <i>Inflammatory Bowel Diseases</i> , 2016, 22, 1533-1539.	1.9	43
27	Stress Responses, Adaptation, and Virulence of Bacterial Pathogens During Host Gastrointestinal Colonization. <i>Microbiology Spectrum</i> , 2016, 4, .	3.0	25
28	Altered intestinal microbiota-host mitochondria crosstalk in new onset Crohn's disease. <i>Nature Communications</i> , 2016, 7, 13419.	12.8	326
29	<i>In Vitro</i> Metabolic Labeling of Intestinal Microbiota for Quantitative Metaproteomics. <i>Analytical Chemistry</i> , 2016, 88, 6120-6125.	6.5	40
30	MetaPro-IQ: a universal metaproteomic approach to studying human and mouse gut microbiota. <i>Microbiome</i> , 2016, 4, 31.	11.1	154
31	Functional Impacts of the Intestinal Microbiome in the Pathogenesis of Inflammatory Bowel Disease. <i>Inflammatory Bowel Diseases</i> , 2015, 21, 139-153.	1.9	112
32	Refined analysis of the <i>Campylobacter jejuni</i> iron-dependent/independent Fur- and PerR-transcriptomes. <i>BMC Genomics</i> , 2015, 16, 498.	2.8	49
33	Phenotypic Screening of a Targeted Mutant Library Reveals <i>Campylobacter jejuni</i> Defenses against Oxidative Stress. <i>Infection and Immunity</i> , 2014, 82, 2266-2275.	2.2	38
34	<i>Campylobacter jejuni</i> ferric-enterobactin receptor CfrA is TonB3 dependent and mediates iron acquisition from structurally different catechol siderophores. <i>Metallomics</i> , 2013, 5, 988.	2.4	32
35	The Transcriptional Landscape of <i>Campylobacter jejuni</i> under Iron Replete and Iron Limited Growth Conditions. <i>PLoS ONE</i> , 2013, 8, e79475.	2.5	39
36	Structure and regulon of <i>Campylobacter jejuni</i> ferric uptake regulator Fur define apo-Fur regulation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 10047-10052.	7.1	114

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37	Nutrient Acquisition and Metabolism by <i>Campylobacter jejuni</i> . <i>Frontiers in Cellular and Infection Microbiology</i> , 2012, 2, 5.	3.9	108
38	Use of a Rabbit Soft Tissue Chamber Model to Investigate <i>Campylobacter Jejuni</i> –Host Interactions. <i>Frontiers in Microbiology</i> , 2010, 1, 126.	3.5	12
39	Micromanaging Oligodendrocyte Differentiation by Noncoding RNA: Toward a Better Understanding of the Lineage Commitment Process. <i>Journal of Neuroscience</i> , 2009, 29, 5365-5366.	3.6	6
40	Characterization of the oxidative stress stimulon and PerR regulon of <i>Campylobacter jejuni</i> . <i>BMC Genomics</i> , 2009, 10, 481.	2.8	144
41	Stress Responses, Adaptation, and Virulence of Bacterial Pathogens During Host Gastrointestinal Colonization. , 0, , 385-411.		18