James Butcher

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5769877/publications.pdf Version: 2024-02-01



| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Altered intestinal microbiota–host mitochondria crosstalk in new onset Crohn's disease. Nature Communications, 2016, 7, 13419. | 12.8 | 326 |
| 2 | Metaproteomics reveals associations between microbiome and intestinal extracellular vesicle proteins in pediatric inflammatory bowel disease. Nature Communications, 2018, 9, 2873. | 12.8 | 209 |
| 3 | Advancing functional and translational microbiome research using meta-omics approaches. Microbiome, 2019, 7, 154. | 11.1 | 177 |
| 4 | MetaPro-IQ: a universal metaproteomic approach to studying human and mouse gut microbiota. Microbiome, 2016, 4, 31. | 11.1 | 154 |
| 5 | Characterization of the oxidative stress stimulon and PerR regulon of Campylobacter jejuni. BMC Genomics, 2009, 10, 481. | 2.8 | 144 |
| 6 | Structure and regulon of <i>Campylobacter jejuni</i> ferric uptake regulator Fur define apo-Fur regulation. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 10047-10052. | 7.1 | 114 |
| 7 | Functional Impacts of the Intestinal Microbiome in the Pathogenesis of Inflammatory Bowel Disease. Inflammatory Bowel Diseases, 2015, 21, 139-153. | 1.9 | 112 |
| 8 | Disruption of maternal gut microbiota during gestation alters offspring microbiota and immunity. Microbiome, 2018, 6, 124. | 11.1 | 109 |
| 9 | Nutrient Acquisition and Metabolism by Campylobacter jejuni. Frontiers in Cellular and Infection Microbiology, 2012, 2, 5. | 3.9 | 108 |
| 10 | Refined analysis of the Campylobacter jejuni iron-dependent/independent Fur- and PerR-transcriptomes. BMC Genomics, 2015, 16, 498. | 2.8 | 49 |
| 11 | Transcriptomic Analysis of the Campylobacter jejuni Response to T4-Like Phage NCTC 12673 Infection. Viruses, 2018, 10, 332. | 3.3 | 46 |
| 12 | Mucosa-Associated Ileal Microbiota in New-Onset Pediatric Crohn's Disease. Inflammatory Bowel Diseases, 2016, 22, 1533-1539. | 1.9 | 43 |
| 13 | Evaluating in Vitro Culture Medium of Gut Microbiome with Orthogonal Experimental Design and a Metaproteomics Approach. Journal of Proteome Research, 2018, 17, 154-163. | 3.7 | 41 |
| 14 | <i>In Vitro</i> Metabolic Labeling of Intestinal Microbiota for Quantitative Metaproteomics. Analytical Chemistry, 2016, 88, 6120-6125. | 6.5 | 40 |
| 15 | The Transcriptional Landscape of Campylobacter jejuni under Iron Replete and Iron Limited Growth Conditions. PLoS ONE, 2013, 8, e79475. | 2.5 | 39 |
| 16 | Phenotypic Screening of a Targeted Mutant Library Reveals Campylobacter jejuni Defenses against Oxidative Stress. Infection and Immunity, 2014, 82, 2266-2275. | 2.2 | 38 |
| 17 | Campylobacter jejuni ferricâ \in enterobactin receptor CfrA is TonB3 dependent and mediates iron acquisition from structurally different catechol siderophores. Metallomics, 2013, 5, 988. | 2.4 | 32 |
| 18 | The mucosal–luminal interface: an ideal sample to study the mucosa-associated microbiota and the intestinal microbial biogeography. Pediatric Research, 2019, 85, 895-903. | 2.3 | 32 |

JAMES BUTCHER

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 19 | Mothers of Preterm Infants Have Individualized Breast Milk Microbiota that Changes Temporally Based on Maternal Characteristics. Cell Host and Microbe, 2020, 28, 669-682.e4. | 11.0 | 31 |
| 20 | Variation on a theme: investigating the structural repertoires used by ferric uptake regulators to control gene expression. BioMetals, 2018, 31, 681-704. | 4.1 | 27 |
| 21 | The gastrointestinal pathogen Campylobacter jejuni metabolizes sugars with potential help from commensal Bacteroides vulgatus. Communications Biology, 2020, 3, 2. | 4.4 | 26 |
| 22 | Stress Responses, Adaptation, and Virulence of Bacterial Pathogens During Host Gastrointestinal Colonization. Microbiology Spectrum, 2016, 4, . | 3.0 | 25 |
| 23 | 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. Journal of Nutrition, 2021, 151, 320-329. | 2.9 | 24 |
| 24 | Independent of Birth Mode or Gestational Age, Very-Low-Birth-Weight Infants Fed Their Mothers' Milk Rapidly Develop Personalized Microbiotas Low in Bifidobacterium. Journal of Nutrition, 2018, 148, 326-335. | 2.9 | 22 |
| 25 | 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. BMC Microbiology, 2020, 20, 219. | 3.3 | 20 |
| 26 | Stress Responses, Adaptation, and Virulence of Bacterial Pathogens During Host Gastrointestinal Colonization. , 0, , 385-411. | | 18 |
| 27 | The impact of probiotics and lactoferrin supplementation on piglet gastrointestinal microbial communities. BioMetals, 2019, 32, 533-543. | 4.1 | 18 |
| 28 | Methods and Strategies to Examine the Human Breastmilk Microbiome. Methods in Molecular Biology, 2018, 1849, 63-86. | 0.9 | 15 |
| 29 | Virome Sequencing of the Human Intestinal Mucosal–Luminal Interface. Frontiers in Cellular and Infection Microbiology, 2020, 10, 582187. | 3.9 | 14 |
| 30 | Binding of Phage-Encoded FlaGrab to Motile Campylobacter jejuni Flagella Inhibits Growth, Downregulates Energy Metabolism, and Requires Specific Flagellar Glycans. Frontiers in Microbiology, 2020, 11, 397. | 3.5 | 14 |
| 31 | Functional insights into the interplay between DNA interaction and metal coordination in ferric uptake regulators. Scientific Reports, 2018, 8, 7140. | 3.3 | 13 |
| 32 | Use of a Rabbit Soft Tissue Chamber Model to Investigate Campylobacter Jejuni–Host Interactions. Frontiers in Microbiology, 2010, 1, 126. | 3.5 | 12 |
| 33 | 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. Journal of Nutrition, 2021, 151, 3431-3441. | 2.9 | 10 |
| 34 | NuA4 Lysine Acetyltransferase Complex Contributes to Phospholipid Homeostasis in Saccharomyces cerevisiae. G3: Genes, Genomes, Genetics, 2017, 7, 1799-1809. | 1.8 | 7 |
| 35 | Micromanaging Oligodendrocyte Differentiation by Noncoding RNA: Toward a Better Understanding of the Lineage Commitment Process. Journal of Neuroscience, 2009, 29, 5365-5366. | 3.6 | 6 |
| 36 | Crystal structure of <i>Campylobacter jejuni</i> peroxide regulator. FEBS Letters, 2018, 592, 2351-2360. | 2.8 | 6 |

JAMES BUTCHER

| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 37 | Bovine Lactoferrin Supplementation Does Not Disrupt Microbiota Development in Preterm Infants Receiving Probiotics. Journal of Pediatric Gastroenterology and Nutrition, 2020, 71, 216-222. | 1.8 | 5 |
| 38 | Examining the Effects of an Anti-Salmonella Bacteriophage Preparation, BAFASAL®, on Ex-Vivo Human Gut Microbiome Composition and Function Using a Multi-Omics Approach. Viruses, 2021, 13, 1734. | 3.3 | 5 |
| 39 | Reduced Infection Efficiency of Phage NCTC 12673 on Non-Motile Campylobacter jejuni Strains Is Related to Oxidative Stress. Viruses, 2021, 13, 1955. | 3.3 | 4 |
| 40 | Analyzing Prokaryotic RNA-Seq Data: A Case Study Identifying Holo-Fur Regulated Genes in Campylobacter jejuni. Methods in Molecular Biology, 2017, 1512, 245-256. | 0.9 | 0 |
| 41 | Characterization of gastrointestinal pathologies in the dystonia musculorum mouse model for hereditary sensory and autonomic neuropathy type VI. Neurogastroenterology and Motility, 2020, 32, e13773. | 3.0 | 0 |