Lindsay J Hall

List of Publications by Year in descending order

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Ιμπραγιμαι

#	Article	IF	CITATIONS
1	<i>>Bifidobacterium castoris</i> strains isolated from wild mice show evidence of frequent host switching and diverse carbohydrate metabolism potential. ISME Communications, 2022, 2, .	1.7	0
2	Macrophage metabolism in the intestine is compartment specific and regulated by the microbiota. Immunology, 2022, 166, 138-152.	2.0	10
3	PRObiotics and SYNbiotics to improve gut health and growth in infants in western Kenya (PROSYNK) Tj ETQq1 1	0.784314 0.7	rgBT /Overlo
4	A systems genomics approach to uncover patient-specific pathogenic pathways and proteins in ulcerative colitis. Nature Communications, 2022, 13, 2299.	5.8	9
5	Maternal gut microbiota Bifidobacterium promotes placental morphogenesis, nutrient transport and fetal growth in mice. Cellular and Molecular Life Sciences, 2022, 79, .	2.4	19
6	Bacterial strains augment cancer therapeutics. Nature Microbiology, 2021, 6, 275-276.	5.9	2
7	Exploring the impact of gut microbiota and diet on breast cancer risk and progression. International Journal of Cancer, 2021, 149, 494-504.	2.3	22
8	Improved molecular characterization of the Klebsiella oxytoca complex reveals the prevalence of the kleboxymycin biosynthetic gene cluster. Microbial Genomics, 2021, 7, .	1.0	10
9	The Pregnancy and EARly Life study (PEARL) - a longitudinal study to understand how gut microbes contribute to maintaining health during pregnancy and early life. BMC Pediatrics, 2021, 21, 357.	0.7	2
10	Antibiotic-induced disturbances of the gut microbiota result in accelerated breast tumor growth. IScience, 2021, 24, 103012.	1.9	41
11	Exploring the Genomic Diversity and Antimicrobial Susceptibility of Bifidobacterium pseudocatenulatum in a Vietnamese Population. Microbiology Spectrum, 2021, 9, e0052621.	1.2	6
12	Microbes, human milk, and prebiotics. , 2021, , 197-237.		2
13	Enterococcus innesii sp. nov., isolated from the wax moth Galleria mellonella. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	0.8	9
14	Incidence of necrotising enterocolitis before and after introducing routine prophylactic <i>Lactobacillus</i> and <i>Bifidobacterium</i> probiotics. Archives of Disease in Childhood: Fetal and Neonatal Edition, 2020, 105, 380-386.	1.4	70
15	Breast milk-derived human milk oligosaccharides promote <i>Bifidobacterium</i> interactions within a single ecosystem. ISME Journal, 2020, 14, 635-648.	4.4	220
16	Rapid MinION profiling of preterm microbiota and antimicrobial-resistant pathogens. Nature Microbiology, 2020, 5, 430-442.	5.9	113
17	Microbiota Supplementation with Bifidobacterium and Lactobacillus Modifies the Preterm Infant Gut Microbiota and Metabolome: An Observational Study. Cell Reports Medicine, 2020, 1, 100077.	3.3	119
18	Preterm Infants Harbour a Rapidly Changing Mycobiota That Includes Candida Pathobionts. Journal of Fungi (Basel, Switzerland), 2020, 6, 273.	1.5	21

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19	Succession of Bifidobacterium longum Strains in Response to a Changing Early Life Nutritional Environment Reveals Dietary Substrate Adaptations. IScience, 2020, 23, 101368.	1.9	26
20	Bifidobacterium breve UCC2003 Induces a Distinct Global Transcriptomic Program in Neonatal Murine Intestinal Epithelial Cells. IScience, 2020, 23, 101336.	1.9	23
21	The early life microbiota protects neonatal mice from pathological small intestinal epithelial cell shedding. FASEB Journal, 2020, 34, 7075-7088.	0.2	27
22	Bifidobacterium breve UCC2003 Exopolysaccharide Modulates the Early Life Microbiota by Acting as a Potential Dietary Substrate. Nutrients, 2020, 12, 948.	1.7	22
23	Setting the agenda for social science research on the human microbiome. Palgrave Communications, 2020, 6, .	4.7	39
24	Preterm infants harbour diverse Klebsiella populations, including atypical species that encode and produce an array of antimicrobial resistance- and virulence-associated factors. Microbial Genomics, 2020, 6, .	1.0	35
25	Recent advances in understanding the neonatal microbiome. F1000Research, 2020, 9, 422.	0.8	22
26	Antibiotic use and the risk of rheumatoid arthritis: a population-based case-control study. BMC Medicine, 2019, 17, 154.	2.3	23
27	Genomic Analysis of Clostridium perfringens BEC/CPILE-Positive, Toxinotype D and E Strains Isolated from Healthy Children. Toxins, 2019, 11, 543.	1.5	11
28	Integrative analysis of Paneth cell proteomic and transcriptomic data from intestinal organoids reveals functional processes dependent on autophagy. DMM Disease Models and Mechanisms, 2019, 12, .	1.2	20
29	Streaming histogram sketching for rapid microbiome analytics. Microbiome, 2019, 7, 40.	4.9	18
30	Genomic analysis on broiler-associated Clostridium perfringens strains and exploratory caecal microbiome investigation reveals key factors linked to poultry necrotic enteritis. Animal Microbiome, 2019, 1, 12.	1.5	29
31	Draft Genome Sequences of Citrobacter freundii and Citrobacter murliniae Strains Isolated from the Feces of Preterm Infants. Microbiology Resource Announcements, 2019, 8, .	0.3	6
32	Phylogenomic analysis of gastroenteritis-associated Clostridium perfringens in England and Wales over a 7-year period indicates distribution of clonal toxigenic strains in multiple outbreaks and extensive involvement of enterotoxin-encoding (CPE) plasmids. Microbial Genomics, 2019, 5, .	1.0	16
33	Draft Genome Sequence of Raoultella ornithinolytica P079F W, Isolated from the Feces of a Preterm Infant. Microbiology Resource Announcements, 2019, 8, .	0.3	2
34	Improving causality in microbiome research: can human genetic epidemiology help?. Wellcome Open Research, 2019, 4, 199.	0.9	21
35	Improving causality in microbiome research: can human genetic epidemiology help?. Wellcome Open Research, 2019, 4, 199.	0.9	28
36	The microbiota, antibiotics and breast cancer. Breast Cancer Management, 2019, 8, BMT29.	0.2	8

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37	The impact of storage conditions on human stool 16S rRNA microbiome composition and diversity. PeerJ, 2019, 7, e8133.	0.9	20
38	Response: Commentary: Probing Genomic Aspects of the Multi-Host Pathogen Clostridium perfringens Reveals Significant Pangenome Diversity, and a Diverse Array of Virulence Factors. Frontiers in Microbiology, 2018, 9, 1857.	1.5	1
39	Antibiotics induce sustained dysregulation of intestinal T cell immunity by perturbing macrophage homeostasis. Science Translational Medicine, 2018, 10, .	5.8	200
40	An update on the human and animal enteric pathogen <i>Clostridium perfringens</i> . Emerging Microbes and Infections, 2018, 7, 1-15.	3.0	262
41	<i>Bifidobacterium breve</i> reduces apoptotic epithelial cell shedding in an exopolysaccharide and MyD88-dependent manner. Open Biology, 2017, 7, 160155.	1.5	65
42	The microbiome beyond the horizon of ecological and evolutionary theory. Nature Ecology and Evolution, 2017, 1, 1606-1615.	3.4	216
43	Preterm Infant-Associated Clostridium tertium, Clostridium cadaveris, and Clostridium paraputrificum Strains: Genomic and Evolutionary Insights. Genome Biology and Evolution, 2017, 9, 2707-2714.	1.1	39
44	Optimisation of 16S rRNA gut microbiota profiling of extremely low birth weight infants. BMC Genomics, 2017, 18, 841.	1.2	47
45	Probing Genomic Aspects of the Multi-Host Pathogen Clostridium perfringens Reveals Significant Pangenome Diversity, and a Diverse Array of Virulence Factors. Frontiers in Microbiology, 2017, 8, 2485.	1.5	70
46	Exploring the role of the microbiota member <i>Bifidobacterium</i> in modulating immune-linked diseases. Emerging Topics in Life Sciences, 2017, 1, 333-349.	1.1	78
47	Gut Microbiome in New-Onset Crohn's Disease. Gastroenterology, 2014, 147, 932-934.	0.6	18
48	Development and characterization of an enhanced nonviral expression vector for electroporation cancer treatment. Molecular Therapy - Methods and Clinical Development, 2014, 1, 14012.	1.8	6
49	Regulation of Host Gene Expression by Gut Microbiota. Gastroenterology, 2013, 144, 841-844.	0.6	2
50	Natural killer cells protect mice from DSS-induced colitis by regulating neutrophil function via the NKG2A receptor. Mucosal Immunology, 2013, 6, 1016-1026.	2.7	55
51	A mouse model of pathological small intestinal epithelial cell apoptosis and shedding induced by systemic administration of lipopolysaccharide. DMM Disease Models and Mechanisms, 2013, 6, 1388-99.	1.2	137
52	Natural Killer Cells Protect against Mucosal and Systemic Infection with the Enteric Pathogen Citrobacter rodentium. Infection and Immunity, 2013, 81, 460-469.	1.0	53
53	Bifidobacterium breve UCC2003 surface exopolysaccharide production is a beneficial trait mediating commensal-host interaction through immune modulation and pathogen protection. Gut Microbes, 2012, 3, 420-425.	4.3	67
54	The Sphingosine-1-Phosphate Analogue FTY720 Impairs Mucosal Immunity and Clearance of the Enteric Pathogen Citrobacter rodentium. Infection and Immunity, 2012, 80, 2712-2723.	1.0	23

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55	Role of Autophagy in NOD2-Induced Inflammation in Crohn's Disease. Gastroenterology, 2012, 142, 1032-1034.	0.6	4
56	Cell Shedding: Old Questions Answered. Gastroenterology, 2012, 143, 1389-1391.	0.6	7
57	Bifidobacterial surface-exopolysaccharide facilitates commensal-host interaction through immune modulation and pathogen protection. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 2108-2113.	3.3	450
58	Induction and Activation of Adaptive Immune Populations During Acute and Chronic Phases of a Murine Model of Experimental Colitis. Digestive Diseases and Sciences, 2011, 56, 79-89.	1.1	88
59	Live Vaccines and Their Role in Modern Vaccinology. , 2011, , 3-14.		0
60	A Salmonella Typhimurium-Typhi Genomic Chimera: A Model to Study Vi Polysaccharide Capsule Function In Vivo. PLoS Pathogens, 2011, 7, e1002131.	2.1	41
61	Probing local innate immune responses after mucosal immunisation. Journal of Immune Based Therapies and Vaccines, 2010, 8, 5.	2.4	4
62	Use of bioluminescence imaging to track neutrophil migration and its inhibition in experimental colitis. Clinical and Experimental Immunology, 2010, 162, 188-196.	1.1	30
63	NK Cells Influence Both Innate and Adaptive Immune Responses after Mucosal Immunization with Antigen and Mucosal Adjuvant. Journal of Immunology, 2010, 184, 4327-4337.	0.4	35
64	Characterisation of a live Salmonella vaccine stably expressing the Mycobacterium tuberculosis Ag85B–ESAT6 fusion protein. Vaccine, 2009, 27, 6894-6904.	1.7	25
65	Candidate Live, Attenuated Salmonella enterica Serotype Typhimurium Vaccines with Reduced Fecal Shedding Are Immunogenic and Effective Oral Vaccines, Infection and Immunity, 2007, 75, 1835-1842.	1.0	47