

Strain-resolved analysis of hospital rooms and infants and room microbiome

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Hospitalized Premature Infants Are Colonized by Related Bacterial Strains with Distinct Proteomic Profiles. <i>MBio</i> , 2018, 9, .	1.8	34
2	Machine Learning Leveraging Genomes from Metagenomes Identifies Influential Antibiotic Resistance Genes in the Infant Gut Microbiome. <i>MSystems</i> , 2018, 3, .	1.7	68
3	Birth mode is associated with earliest strain-conferred gut microbiome functions and immunostimulatory potential. <i>Nature Communications</i> , 2018, 9, 5091.	5.8	190
4	Genetic relatedness of Gram-negative bacteria colonizing gut and skin of neonates and mother's own milk. <i>Journal of Perinatology</i> , 2018, 38, 1503-1511.	0.9	4
5	Higher intake of coagulase-negative staphylococci from maternal milk promotes gut colonization with <i>mecA</i> -negative <i>Staphylococcus epidermidis</i> in preterm neonates. <i>Journal of Perinatology</i> , 2018, 38, 1344-1352.	0.9	3
6	The developing premature infant gut microbiome is a major factor shaping the microbiome of neonatal intensive care unit rooms. <i>Microbiome</i> , 2018, 6, 112.	4.9	65
7	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
8	Application of machine learning techniques for creating urban microbial fingerprints. <i>Biology Direct</i> , 2019, 14, 13.	1.9	29
9	What Pediatricians Should Know Before Studying Gut Microbiota. <i>Journal of Clinical Medicine</i> , 2019, 8, 1206.	1.0	8
10	Multifaceted mechanisms of colistin resistance revealed by genomic analysis of multidrug-resistant <i>Klebsiella pneumoniae</i> isolates from individual patients before and after colistin treatment. <i>Journal of Infection</i> , 2019, 79, 312-321.	1.7	24
11	Hybrid metagenomic assembly enables high-resolution analysis of resistance determinants and mobile elements in human microbiomes. <i>Nature Biotechnology</i> , 2019, 37, 937-944.	9.4	216
12	The role of the preterm intestinal microbiome in sepsis and necrotising enterocolitis. <i>Early Human Development</i> , 2019, 138, 104854.	0.8	48
13	Feeding intolerance alters the gut microbiota of preterm infants. <i>PLoS ONE</i> , 2019, 14, e0210609.	1.1	19
14	Prospective surveillance of bacterial colonization and primary sepsis: findings of a tertiary neonatal intensive and intermediate care unit. <i>Journal of Hospital Infection</i> , 2019, 102, 325-331.	1.4	24
15	Longitudinal Gut Bacterial Colonization and Its Influencing Factors of Low Birth Weight Infants During the First 3 Months of Life. <i>Frontiers in Microbiology</i> , 2019, 10, 1105.	1.5	18
16	Longitudinal homogenization of the microbiome between both occupants and the built environment in a cohort of United States Air Force Cadets. <i>Microbiome</i> , 2019, 7, 70.	4.9	33
17	The Water Microbiome Through a Pilot Scale Advanced Treatment Facility for Direct Potable Reuse. <i>Frontiers in Microbiology</i> , 2019, 10, 993.	1.5	36
18	Breastmilk and NICU surfaces are potential sources of fungi for infant mycobiomes. <i>Fungal Genetics and Biology</i> , 2019, 128, 29-35.	0.9	27

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19	Using formalin fixed paraffin embedded tissue to characterize the preterm gut microbiota in necrotising enterocolitis and spontaneous isolated perforation using marginal and diseased tissue. BMC Microbiology, 2019, 19, 52.	1.3	24
20	Ten questions concerning the built environment and mental health. Building and Environment, 2019, 155, 58-69.	3.0	68
21	In-host evolution of Staphylococcus epidermidis in a pacemaker-associated endocarditis resulting in increased antibiotic tolerance. Nature Communications, 2019, 10, 1149.	5.8	64
22	The Microbiome, Metabolome, and Proteome in Preterm Neonatal Sepsis. , 2019, , 279-285.		0
23	Longitudinal Microbiome Composition and Stability Correlate with Increased Weight and Length of Very-Low-Birth-Weight Infants. MSystems, 2019, 4, .	1.7	51
24	Genome-resolved metagenomics of eukaryotic populations during early colonization of premature infants and in hospital rooms. Microbiome, 2019, 7, 26.	4.9	60
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27	Microbial transmission from mother to child: improving infant intestinal microbiota development by identifying the obstacles. Critical Reviews in Microbiology, 2019, 45, 613-648.	2.7	30
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29	Extensive Unexplored Human Microbiome Diversity Revealed by Over 150,000 Genomes from Metagenomes Spanning Age, Geography, and Lifestyle. Cell, 2019, 176, 649-662.e20.	13.5	1,087
30	A One Health Study of the Genetic Relatedness of <i>Klebsiella pneumoniae</i> and Their Mobile Elements in the East of England. Clinical Infectious Diseases, 2020, 70, 219-226.	2.9	46
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32	Genetic relationship between bacteria isolated from intraoperative air samples and surgical site infections at a major teaching hospital in Ghana. Journal of Hospital Infection, 2020, 104, 309-320.	1.4	8
33	Fungal cutaneous microbiome and host determinants in preterm and term neonates. Pediatric Research, 2020, 88, 225-233.	1.1	13
34	Acquisition and Development of the Extremely Preterm Infant Microbiota Across Multiple Anatomical Sites. Journal of Pediatric Gastroenterology and Nutrition, 2020, 70, 12-19.	0.9	16
35	Assessment of Neonatal Intensive Care Unit Practices and Preterm Newborn Gut Microbiota and 2-Year Neurodevelopmental Outcomes. JAMA Network Open, 2020, 3, e2018119.	2.8	44
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38	Continental-Scale Microbiome Study Reveals Different Environmental Characteristics Determining Microbial Richness, Composition, and Quantity in Hotel Rooms. <i>MSystems</i> , 2020, 5, .	1.7	20
39	Diversity within species: interpreting strains in microbiomes. <i>Nature Reviews Microbiology</i> , 2020, 18, 491-506.	13.6	222
40	Cartography of opportunistic pathogens and antibiotic resistance genes in a tertiary hospital environment. <i>Nature Medicine</i> , 2020, 26, 941-951.	15.2	130
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49	The effect of early probiotic exposure on the preterm infant gut microbiome development. <i>Gut Microbes</i> , 2021, 13, 1951113.	4.3	26
50	Improved metagenome binning and assembly using deep variational autoencoders. <i>Nature Biotechnology</i> , 2021, 39, 555-560.	9.4	251
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53	Comparative genomics of two <i>Shewanella xiamenensis</i> strains isolated from a pilgrim before and during travels to the Hajj. <i>Gut Pathogens</i> , 2021, 13, 9.	1.6	5
54	Impact of Probiotic <i>B. infantis</i> EVC001 Feeding in Premature Infants on the Gut Microbiome, Nosocomially Acquired Antibiotic Resistance, and Enteric Inflammation. <i>Frontiers in Pediatrics</i> , 2021, 9, 618009.	0.9	38
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66	Escaping the fate of Sisyphus: assessing resistome hybridization baits for antimicrobial resistance gene capture. <i>Environmental Microbiology</i> , 2021, 23, 7523-7537.	1.8	3
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81	Rapid methicillin resistance diversification in <i>Staphylococcus epidermidis</i> colonizing human neonates. <i>Nature Communications</i> , 2021, 12, 6062.	5.8	6
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