Marcus C De Goffau

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The Dynamics of the Human Infant Gut Microbiome in Development and in Progression toward Type 1 Diabetes. Cell Host and Microbe, 2015, 17, 260-273.	11.0	1,008
2	Human placenta has no microbiome but can contain potential pathogens. Nature, 2019, 572, 329-334.	27.8	513
3	Fecal Microbiota Composition Differs Between Children With β-Cell Autoimmunity and Those Without. Diabetes, 2013, 62, 1238-1244.	0.6	498
4	Aberrant gut microbiota composition at the onset of type 1 diabetes in young children. Diabetologia, 2014, 57, 1569-1577.	6.3	274
5	Recognizing the reagent microbiome. Nature Microbiology, 2018, 3, 851-853.	13.3	255
6	Real-time in vivo imaging of invasive- and biomaterial-associated bacterial infections using fluorescently labelled vancomycin. Nature Communications, 2013, 4, 2584.	12.8	231
7	SIRT1 stimulation by polyphenols is affected by their stability and metabolism. Mechanisms of Ageing and Development, 2006, 127, 618-627.	4.6	148
8	One Health Genomic Surveillance of Escherichia coli Demonstrates Distinct Lineages and Mobile Genetic Elements in Isolates from Humans versus Livestock. MBio, 2019, 10, .	4.1	130
9	Diversity of human small intestinal <i>Streptococcus</i> and <i>Veillonella</i> populations. FEMS Microbiology Ecology, 2013, 85, 376-388.	2.7	121
10	A Necrotizing Enterocolitis-Associated Gut Microbiota Is Present in the Meconium: Results of a Prospective Study. Clinical Infectious Diseases, 2016, 62, 863-870.	5.8	119
11	Mucus Microbiome of Anastomotic Tissue During Surgery Has Predictive Value for Colorectal Anastomotic Leakage. Annals of Surgery, 2019, 269, 911-916.	4.2	92
12	The <i>ATG16L1–T300A</i> allele impairs clearance of pathosymbionts in the inflamed ileal mucosa of Crohn's disease patients. Gut, 2015, 64, 1546-1552.	12.1	77
13	The Human Gut Microbiota. Advances in Experimental Medicine and Biology, 2016, 902, 95-108.	1.6	72
14	High level of fecal calprotectin at age 2Âmonths as a marker of intestinal inflammation predicts atopic dermatitis and asthma by age 6. Clinical and Experimental Allergy, 2015, 45, 928-939.	2.9	69
15	Genomic Surveillance of Enterococcus faecium Reveals Limited Sharing of Strains and Resistance Genes between Livestock and Humans in the United Kingdom. MBio, 2018, 9, .	4.1	63
16	Intestinal microbiota and anastomotic leakage of stapled colorectal anastomoses: a pilot study. Surgical Endoscopy and Other Interventional Techniques, 2016, 30, 2259-2265.	2.4	62
17	A longitudinal study of the infant nasopharyngeal microbiota: The effects of age, illness and antibiotic use in a cohort of South East Asian children. PLoS Neglected Tropical Diseases, 2017, 11, e0005975.	3.0	62
18	Bacterial pleomorphism and competition in a relative humidity gradient. Environmental Microbiology, 2009, 11, 809-822.	3.8	53

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19	Maturation of Gut Microbiota and Circulating Regulatory T Cells and Development of IgE Sensitization in Early Life. Frontiers in Immunology, 2019, 10, 2494.	4.8	46
20	STROBE-metagenomics: a STROBE extension statement to guide the reporting of metagenomics studies. Lancet Infectious Diseases, The, 2020, 20, e251-e260.	9.1	40
21	Standard of hygiene and immune adaptation in newborn infants. Clinical Immunology, 2014, 155, 136-147.	3.2	35
22	Batch effects account for the main findings of an in utero human intestinal bacterial colonization study. Microbiome, 2021, 9, 6.	11.1	34
23	Gut microbiomes from Gambian infants reveal the development of a non-industrialized Prevotella-based trophic network. Nature Microbiology, 2022, 7, 132-144.	13.3	30
24	Fetal inheritance of chromosomally integrated human herpesvirus 6 predisposes the mother to pre-eclampsia. Nature Microbiology, 2020, 5, 901-908.	13.3	29
25	Exploring the risk factors for differences in the cumulative incidence of coeliac disease in two neighboring countries: the prospective DIABIMMUNE study. Digestive and Liver Disease, 2016, 48, 1296-1301.	0.9	26
26	Increasing incidence of group B streptococcus neonatal infections in the Netherlands is associated with clonal expansion of CC17 and CC23. Scientific Reports, 2020, 10, 9539.	3.3	25
27	Effects of fecal microbiota transplant on DNA methylation in subjects with metabolic syndrome. Gut Microbes, 2021, 13, 1993513.	9.8	25
28	Genetic loci of Staphylococcus aureus associated with anti-neutrophil cytoplasmic autoantibody (ANCA)-associated vasculitides. Scientific Reports, 2017, 7, 12211.	3.3	24
29	Contrasting patterns of longitudinal population dynamics and antimicrobial resistance mechanisms in two priority bacterial pathogens over 7Âyears in a single center. Genome Biology, 2019, 20, 184.	8.8	22
30	Detecting eukaryotic microbiota with single-cell sensitivity in human tissue. Microbiome, 2018, 6, 151.	11.1	21
31	Low anti-staphylococcal IgG responses in granulomatosis with polyangiitis patients despite long-term Staphylococcus aureus exposure. Scientific Reports, 2015, 5, 8188.	3.3	20
32	Cold Spots in Neonatal Incubators Are Hot Spots for Microbial Contamination. Applied and Environmental Microbiology, 2011, 77, 8568-8572.	3.1	19
33	Multi-Compartment Profiling of Bacterial and Host Metabolites Identifies Intestinal Dysbiosis and Its Functional Consequences in the Critically III Child. Critical Care Medicine, 2019, 47, e727-e734.	0.9	19
34	Microbial growth on the edge of desiccation. Environmental Microbiology, 2011, 13, 2328-2335.	3.8	17
35	'Candidatus Ornithobacterium hominis': insights gained from draft genomes obtained from nasopharyngeal swabs. Microbial Genomics, 2019, 5, .	2.0	16
36	Patients With Inflammatory Bowel Disease Show IgG Immune Responses Towards Specific Intestinal Bacterial Genera. Frontiers in Immunology, 0, 13, .	4.8	12

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37	Over-celling fetal microbial exposure. Cell, 2021, 184, 5839-5841.	28.9	10
38	Biofilm formation on the Provox ActiValve: Composition and ingrowth analyzed by Illumina pairedâ€end RNA sequencing, fluorescence in situ hybridization, and confocal laser scanning microscopy. Head and Neck, 2016, 38, E432-40.	2.0	9
39	Weight shapes the intestinal microbiome in preterm infants: results of a prospective observational study. BMC Microbiology, 2021, 21, 219.	3.3	9
40	The Dynamics of the Human Infant Gut Microbiome in Development and in Progression toward Type 1 Diabetes. Cell Host and Microbe, 2016, 20, 121.	11.0	7
41	Early childhood infections and the use of antibiotics and antipyreticâ€analgesics in Finland, Estonia and Russian Karelia. Acta Paediatrica, International Journal of Paediatrics, 2019, 108, 2075-2082.	1.5	7
42	Differential expression of a prophage-encoded glycocin and its immunity protein suggests a mutualistic strategy of a phage and its host. Scientific Reports, 2019, 9, 2845.	3.3	7
43	Profiling gut microbiota and bile acid metabolism in critically ill children. Scientific Reports, 2022, 12,	3.3	5
44	Response to Comment on "Mucus Microbiome of Anastomotic Tissue During Surgery Has Predictive Value for Colorectal Anastomotic Leakage― Annals of Surgery, 2019, 269, e69-e70.	4.2	2
45	When to suspect contamination rather than colonization $\hat{a} \in$ lessons from a putative fetal sheep microbiome. Gut Microbes, 2022, 14, 2005751.	9.8	2
46	Reply to Cassir et al. Clinical Infectious Diseases, 2016, 62, 1618-1620.	5.8	1
47	1369 How Clean is Clean? Effectiveness of Disinfection of Thermometers on a Neonatal Intensive Care Unit. Pediatric Research, 2010, 68, 678-678.	2.3	Ο
48	Tu1707 Crohn's Disease Patients With the ATG16L1 T300a Allele Are Unable to Modify Their Mucosal Microbiota Profile Upon Inflammation. Gastroenterology, 2013, 144, S-827.	1.3	0
49	Sa2007 Intestinal Microbiota and Anastomotic Leakage of Stapled Colorectal Anastomoses. Gastroenterology, 2016, 150, S430.	1.3	0
50	706: LOSS OF FECAL MICROBIAL DENSITY AND INTESTINAL FERMENTATION EFFICIENCY IN CRITICALLY ILL CHILDREN. Critical Care Medicine, 2018, 46, 339-339.	0.9	0