Susan Lynch

List of Publications by Year in descending order

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18436 8835 22,879 146 62 145 citations h-index g-index papers 153 153 153 27708 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Induction of Intestinal Th17 Cells by Segmented Filamentous Bacteria. Cell, 2009, 139, 485-498.	13.5	3,818
2	The Human Intestinal Microbiome in Health and Disease. New England Journal of Medicine, 2016, 375, 2369-2379.	13.9	2,383
3	Current understanding of the human microbiome. Nature Medicine, 2018, 24, 392-400.	15.2	1,593
4	Neonatal gut microbiota associates with childhood multisensitized atopy and T cell differentiation. Nature Medicine, 2016 , 22 , 1187 - 1191 .	15.2	844
5	Comparison of the Respiratory Microbiome in Healthy Nonsmokers and Smokers. American Journal of Respiratory and Critical Care Medicine, 2013, 187, 1067-1075.	2.5	655
6	Airway microbiota and bronchial hyperresponsiveness in patients with suboptimally controlled asthma. Journal of Allergy and Clinical Immunology, 2011, 127, 372-381.e3.	1.5	598
7	Gastrointestinal Microbiome Signatures of Pediatric Patients With Irritable Bowel Syndrome. Gastroenterology, 2011, 141, 1782-1791.	0.6	579
8	Dysbiosis of the Gut Microbiota Is Associated with HIV Disease Progression and Tryptophan Catabolism. Science Translational Medicine, 2013, 5, 193ra91.	5.8	578
9	The gut microbiome: Relationships with disease and opportunities for therapy. Journal of Experimental Medicine, 2019, 216, 20-40.	4.2	547
10	The airway microbiome in patients with severe asthma: Associations with disease features and severity. Journal of Allergy and Clinical Immunology, 2015, 136, 874-884.	1.5	395
11	Airway Microbiota and Pathogen Abundance in Age-Stratified Cystic Fibrosis Patients. PLoS ONE, 2010, 5, e11044.	1.1	395
12	House dust exposure mediates gut microbiome <i>Lactobacillus</i> enrichment and airway immune defense against allergens and virus infection. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 805-810.	3.3	374
13	Sinus Microbiome Diversity Depletion and <i>Corynebacterium tuberculostearicum</i> Enrichment Mediates Rhinosinusitis. Science Translational Medicine, 2012, 4, 151ra124.	5.8	372
14	Role of the gut microbiota in defining human health. Expert Review of Anti-Infective Therapy, 2010, 8, 435-454.	2.0	339
15	Effects of early-life exposure to allergens and bacteria onÂrecurrent wheeze and atopy in urban children. Journal of Allergy and Clinical Immunology, 2014, 134, 593-601.e12.	1.5	333
16	Microbiota in Allergy and Asthma and the Emerging Relationship with the Gut Microbiome. Cell Host and Microbe, 2015, 17, 592-602.	5.1	327
17	Use of 16S rRNA Gene for Identification of a Broad Range of Clinically Relevant Bacterial Pathogens. PLoS ONE, 2015, 10, e0117617.	1.1	293
18	Airway Microbiome Dynamics in Exacerbations of Chronic Obstructive Pulmonary Disease. Journal of Clinical Microbiology, 2014, 52, 2813-2823.	1.8	272

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19	Comparative Analyses of the Bacterial Microbiota of the Human Nostril and Oropharynx. MBio, 2010, 1,	1.8	266
20	Gut microbiota in early pediatric multiple sclerosis: a caseâ^control study. European Journal of Neurology, 2016, 23, 1308-1321.	1.7	260
21	Role of the microbiota in inflammatory bowel diseases. Inflammatory Bowel Diseases, 2012, 18, 968-984.	0.9	237
22	Features of the bronchial bacterial microbiome associated with atopy, asthma, and responsiveness to inhaled corticosteroid treatment. Journal of Allergy and Clinical Immunology, 2017, 140, 63-75.	1.5	222
23	A Persistent and Diverse Airway Microbiota Present during Chronic Obstructive Pulmonary Disease Exacerbations. OMICS A Journal of Integrative Biology, 2010, 14, 9-59.	1.0	213
24	Man's best friend? The effect of pet ownership on house dust microbial communities. Journal of Allergy and Clinical Immunology, 2010, 126, 410-412.e3.	1.5	205
25	Relationship between cystic fibrosis respiratory tract bacterial communities and age, genotype, antibiotics and <i>Pseudomonas aeruginosa</i> . Environmental Microbiology, 2010, 12, 1293-1303.	1.8	203
26	The oral microbiome: Role of key organisms and complex networks in oral health and disease. Periodontology 2000, 2021, 87, 107-131.	6.3	195
27	Significance of the microbiome in obstructive lung disease. Thorax, 2012, 67, 456-463.	2.7	190
28	Alteration of the cutaneous microbiome in psoriasis and potential role in Th17 polarization. Microbiome, 2018, 6, 154.	4.9	190
29	Viable bacterial colonization is highly limited in the human intestine in utero. Nature Medicine, 2020, 26, 599-607.	15.2	180
30	Widespread Colonization of the Lung by <i>Tropheryma whipplei</i> in HIV Infection. American Journal of Respiratory and Critical Care Medicine, 2013, 187, 1110-1117.	2.5	175
31	Loss of Bacterial Diversity during Antibiotic Treatment of Intubated Patients Colonized with Pseudomonas aeruginosa. Journal of Clinical Microbiology, 2007, 45, 1954-1962.	1.8	166
32	Early-life home environment and risk of asthma among inner-city children. Journal of Allergy and Clinical Immunology, 2018, 141, 1468-1475.	1.5	160
33	Persistent Infection with <i>Pseudomonas aeruginosa</i> in Ventilator-associated Pneumonia. American Journal of Respiratory and Critical Care Medicine, 2008, 178, 513-519.	2.5	159
34	Delayed gut microbiota development in high-risk for asthma infants is temporarily modifiable by Lactobacillus supplementation. Nature Communications, 2018, 9, 707.	5.8	158
35	Compositionally and functionally distinct sinus microbiota in chronic rhinosinusitis patients have immunological and clinically divergent consequences. Microbiome, 2017, 5, 53.	4.9	151
36	Elevated faecal 12,13-diHOME concentration in neonates at high risk for asthma is produced by gut bacteria and impedes immune tolerance. Nature Microbiology, 2019, 4, 1851-1861.	5.9	148

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37	Joint effects of pregnancy, sociocultural, and environmental factors on early life gut microbiome structure and diversity. Scientific Reports, 2016, 6, 31775.	1.6	122
38	Multicenter Comparison of Lung and Oral Microbiomes of HIV-infected and HIV-uninfected Individuals. American Journal of Respiratory and Critical Care Medicine, 2015, 192, 1335-1344.	2.5	120
39	ChrR, a Soluble Quinone Reductase of Pseudomonas putida That Defends against H2O2. Journal of Biological Chemistry, 2005, 280, 22590-22595.	1.6	119
40	Distinct nasal airway bacterial microbiotas differentially relate to exacerbation in pediatric patients with asthma. Journal of Allergy and Clinical Immunology, 2019, 144, 1187-1197.	1.5	117
41	Breast-fed and bottle-fed infant rhesus macaques develop distinct gut microbiotas and immune systems. Science Translational Medicine, 2014, 6, 252ra120.	5.8	115
42	Lung Microbiota Is Related to Smoking Status and to Development of Acute Respiratory Distress Syndrome in Critically Ill Trauma Patients. American Journal of Respiratory and Critical Care Medicine, 2018, 197, 621-631.	2.5	114
43	Early Probiotic Supplementation for Eczema and Asthma Prevention: A Randomized Controlled Trial. Pediatrics, 2017, 140, .	1.0	107
44	Lactobacillus casei Abundance Is Associated with Profound Shifts in the Infant Gut Microbiome. PLoS ONE, 2010, 5, e8745.	1.1	107
45	Community ecology as a framework for human microbiome research. Nature Medicine, 2019, 25, 884-889.	15.2	96
46	Dynamics of Bacterial Colonization With Streptococcus pneumoniae, Haemophilus influenzae, and Moraxella catarrhalis During Symptomatic and Asymptomatic Viral Upper Respiratory Tract Infection. Clinical Infectious Diseases, 2018, 66, 1045-1053.	2.9	93
47	Bacterial biogeography of adult airways in atopic asthma. Microbiome, 2018, 6, 104.	4.9	93
48	Associations between the gut microbiota and host immune markers in pediatric multiple sclerosis and controls. BMC Neurology, 2016, 16, 182.	0.8	91
49	The Cystic Fibrosis Airway Microbiome. Cold Spring Harbor Perspectives in Medicine, 2013, 3, a009738-a009738.	2.9	90
50	Disease Severity and Immune Activity Relate to Distinct Interkingdom Gut Microbiome States in Ethnically Distinct Ulcerative Colitis Patients. MBio, 2016, 7, .	1.8	90
51	The emerging relationship between the airway microbiota and chronic respiratory disease: clinical implications. Expert Review of Respiratory Medicine, 2011, 5, 809-821.	1.0	89
52	Enteric Virome and Bacterial Microbiota in Children With Ulcerative Colitis and Crohn Disease. Journal of Pediatric Gastroenterology and Nutrition, 2019, 68, 30-36.	0.9	89
53	Cystic fibrosis transmembrane conductance regulator knockout mice exhibit aberrant gastrointestinal microbiota. Gut Microbes, 2013, 4, 41-47.	4.3	85
54	Characterizing the gut microbiome in trauma: significant changes in microbial diversity occur early after severe injury. Trauma Surgery and Acute Care Open, 2017, 2, e000108.	0.8	83

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55	Human gut bacterial metabolism drives Th17 activation and colitis. Cell Host and Microbe, 2022, 30, 17-30.e9.	5.1	83
56	Gut-Resident Lactobacillus Abundance Associates with IDO1 Inhibition and Th17 Dynamics in SIV-Infected Macaques. Cell Reports, 2015, 13, 1589-1597.	2.9	75
57	Lactobacillus johnsonii supplementation attenuates respiratory viral infection via metabolic reprogramming and immune cell modulation. Mucosal Immunology, 2017, 10, 1569-1580.	2.7	75
58	The microbiome and development of allergic disease. Current Opinion in Allergy and Clinical Immunology, 2016, 16, 165-171.	1.1	73
59	Nasopharyngeal microbiota composition of children is related to the frequency of upper respiratory infection and acute sinusitis. Microbiome, 2016, 4, 34.	4.9	70
60	Longitudinal Phenotypes of Respiratory Health in a High-Risk Urban Birth Cohort. American Journal of Respiratory and Critical Care Medicine, 2019, 199, 71-82.	2.5	70
61	Oral and Airway Microbiota in HIV-Infected Pneumonia Patients. Journal of Clinical Microbiology, 2012, 50, 2995-3002.	1.8	69
62	Effect of prenatal indoor pet exposure on the trajectory of total IgE levels in early childhood. Journal of Allergy and Clinical Immunology, 2011, 128, 880-885.e4.	1.5	66
63	The potential for probiotic manipulation of the gastrointestinal microbiome. Current Opinion in Biotechnology, 2012, 23, 192-201.	3.3	66
64	Increased Plasminogen Activator Inhibitor-1 Concentrations in Bronchoalveolar Lavage Fluids Are Associated with Increased Mortality in a Cohort of Patients with Pseudomonas aeruginosaÂ. Anesthesiology, 2007, 106, 252-261.	1.3	63
65	Pseudomonas aeruginosa biofilm-associated homoserine lactone C12 rapidly activates apoptosis in airway epithelia. Cellular Microbiology, 2012, 14, 698-709.	1.1	62
66	Increased mortality of ventilated patients with endotracheal Pseudomonas aeruginosa without clinical signs of infection*. Critical Care Medicine, 2008, 36, 2495-2503.	0.4	60
67	Immune Response and Mortality Risk Relate to Distinct Lung Microbiomes in Patients with HIV and Pneumonia. American Journal of Respiratory and Critical Care Medicine, 2017, 195, 104-114.	2.5	60
68	Limited engraftment of donor microbiome via one-time fecal microbial transplantation in treated HIV-infected individuals. Gut Microbes, 2017, 8, 440-450.	4.3	56
69	Gut dysbiosis in cystic fibrosis. Journal of Cystic Fibrosis, 2012, 11, 454-455.	0.3	55
70	Presence or Absence of Lipopolysaccharide O Antigens Affects Type III Secretion by Pseudomonas aeruginosa. Journal of Bacteriology, 2007, 189, 2203-2209.	1.0	53
71	The Lung Microbiome of Ugandan HIV-Infected Pneumonia Patients Is Compositionally and Functionally Distinct from That of San Franciscan Patients. PLoS ONE, 2014, 9, e95726.	1.1	53
72	PcrV antibody–antibiotic combination improves survival in Pseudomonas aeruginosa-infected mice. European Journal of Clinical Microbiology and Infectious Diseases, 2012, 31, 1837-1845.	1.3	52

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73	Fecal microbiota transplant for Crohn disease: A study evaluating safety, efficacy, and microbiome profile. United European Gastroenterology Journal, 2019, 7, 807-814.	1.6	51
74	Amelioration of DSS-induced murine colitis by VSL#3 supplementation is primarily associated with changes in ileal microbiota composition Gut Microbes, 2014, 5, 494-503.	4.3	50
75	Viruses and Microbiome Alterations. Annals of the American Thoracic Society, 2014, 11, S57-S60.	1.5	50
76	New enzyme for reductive cancer chemotherapy, YieF, and its improvement by directed evolution. Molecular Cancer Therapeutics, 2006, 5, 97-103.	1.9	49
77	Influence and effect of the human microbiome in allergy and asthma. Current Opinion in Rheumatology, 2015, 27, 373-380.	2.0	49
78	Airway Microbiota and the Implications of Dysbiosis in Asthma. Current Allergy and Asthma Reports, 2016, 16, 52.	2.4	48
79	The human microbiome in the 21st century. Nature Communications, 2020, 11, 5256.	5.8	48
80	Expression quantitative trait locus fine mapping of the 17q12–21 asthma locus in African American children: a genetic association and gene expression study. Lancet Respiratory Medicine,the, 2020, 8, 482-492.	5.2	47
81	Microbial Manipulation of Immune Function for Asthma Prevention: Inferences from Clinical Trials. Proceedings of the American Thoracic Society, 2007, 4, 277-282.	3.5	46
82	Dog introduction alters the home dust microbiota. Indoor Air, 2018, 28, 539-547.	2.0	46
83	Distinct associations of sputum and oral microbiota with atopic, immunologic, and clinical features in mild asthma. Journal of Allergy and Clinical Immunology, 2020, 146, 1016-1026.	1.5	46
84	Gut Microbiota and Allergic Disease. New Insights. Annals of the American Thoracic Society, 2016, 13, S51-S54.	1.5	44
85	Novel strategies to combat bacterial virulence. Current Opinion in Critical Care, 2008, 14, 593-599.	1.6	43
86	Fecal Microbiota Transplantation in Pouchitis: Clinical, Endoscopic, Histologic, and Microbiota Results from a Pilot Study. Digestive Diseases and Sciences, 2020, 65, 1099-1106.	1,1	41
87	Bacteroides are associated with GALT iNKT cell function and reduction of microbial translocation in HIV-1 infection. Mucosal Immunology, 2017, 10, 69-78.	2.7	40
88	Breast Milk Transforming Growth Factor \hat{I}^2 Is Associated With Neonatal Gut Microbial Composition. Journal of Pediatric Gastroenterology and Nutrition, 2017, 65, e60-e67.	0.9	40
89	Prenatal antimicrobial use and early-childhood body mass index. International Journal of Obesity, 2018, 42, 1-7.	1.6	38
90	Differences in the fecal microbiota of neonates born at home or in the hospital. Scientific Reports, 2018, 8, 15660.	1.6	38

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91	Novel Microbiome-Based Therapeutics for Chronic Rhinosinusitis. Current Allergy and Asthma Reports, 2015, 15, 504.	2.4	36
92	The Lung Microbiome and Airway Disease. Annals of the American Thoracic Society, 2016, 13, S462-S465.	1.5	36
93	Clinical Features, Virus Identification, and Sinusitis as a Complication ofÂUpper Respiratory Tract Illness in Children Ages 4-7ÂYears. Journal of Pediatrics, 2016, 171, 133-139.e1.	0.9	36
94	Polymorphisms in the Pseudomonas aeruginosa type III secretion protein, PcrV – Implications for anti-PcrV immunotherapy. Microbial Pathogenesis, 2010, 48, 197-204.	1.3	35
95	Use of bronchoalveolar lavage to assess the respiratory microbiome: signal in the noise. Lancet Respiratory Medicine, the, 2013, 1, 354-356.	5.2	35
96	Translating the gut microbiome: ready for the clinic?. Nature Reviews Gastroenterology and Hepatology, 2019, 16, 656-661.	8.2	33
97	Secretion of Pseudomonas aeruginosa type III cytotoxins is dependent on pseudomonas quinolone signal concentration. Microbial Pathogenesis, 2010, 49, 196-203.	1.3	31
98	Maternal group B <i>Streptococcus</i> and the infant gut microbiota. Journal of Developmental Origins of Health and Disease, 2016, 7, 45-53.	0.7	31
99	Seasonal airway microbiome and transcriptome interactions promote childhood asthma exacerbations. Journal of Allergy and Clinical Immunology, 2022, 150, 204-213.	1.5	31
100	Cervicovaginal Microbiome Composition Is Associated with Metabolic Profiles in Healthy Pregnancy. MBio, 2020, 11 , .	1.8	30
101	Probiotic strategies for treatment of respiratory diseases. Trends in Microbiology, 2013, 21, 485-492.	3.5	27
102	Gut Microbial Metabolism and Nonalcoholic Fatty Liver Disease. Hepatology Communications, 2019, 3, 29-43.	2.0	27
103	Gut microbiota in HIV–pneumonia patients is related to peripheral CD4 counts, lung microbiota, and in vitro macrophage dysfunction. Microbiome, 2019, 7, 37.	4.9	25
104	Microbiota, Epigenetics, and Trained Immunity. Convergent Drivers and Mediators of the Asthma Trajectory from Pregnancy to Childhood. American Journal of Respiratory and Critical Care Medicine, 2021, 203, 802-808.	2.5	23
105	Maternal gut microbiome regulates immunity to RSV infection in offspring. Journal of Experimental Medicine, 2021, 218, .	4.2	22
106	Probiotic manipulation of the gastrointestinal microbiota. Gut Microbes, 2010, 1, 335-338.	4.3	21
107	Gut microbiome is associated with multiple sclerosis activity in children. Annals of Clinical and Translational Neurology, 2021, 8, 1867-1883.	1.7	21
108	Fetal and early postnatal lead exposure measured in teeth associates with infant gut microbiota. Environment International, 2020, 144, 106062.	4.8	21

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109	Heterogeneity of Microbiota Dysbiosis in Chronic Rhinosinusitis: Potential Clinical Implications and Microbial Community Mechanisms Contributing to Sinonasal Inflammation. Frontiers in Cellular and Infection Microbiology, 2018, 8, 168.	1.8	18
110	Association between cesarean delivery types and obesity in preadolescence. International Journal of Obesity, 2020, 44, 2023-2034.	1.6	17
111	Corroborating evidence refutes batch effect as explanation for fetal bacteria. Microbiome, 2021, 9, 10.	4.9	17
112	Strain-resolved analysis in a randomized trial of antibiotic pretreatment and maintenance dose delivery mode with fecal microbiota transplant for ulcerative colitis. Scientific Reports, 2022, 12, 5517.	1.6	17
113	Fecal Microbiota Transplantation for Recurrent <i>Clostridium difficile</i> Infection in Pediatric Patients. Journal of Pediatric Gastroenterology and Nutrition, 2015, 60, 1-3.	0.9	16
114	Matrix metalloproteases in bronchoalveolar lavage fluid of patients with type III Pseudomonas aeruginosa pneumonia. Journal of Infection, 2009, 59, 49-55.	1.7	15
115	Fecal Microbial Therapy. Journal of Pediatric Gastroenterology and Nutrition, 2014, 59, 157-161.	0.9	15
116	A chronic rhinosinusitis–derived isolate of <i>Pseudomonas aeruginosa</i> induces acute and pervasive effects on the murine upper airway microbiome and host immune response. International Forum of Allergy and Rhinology, 2016, 6, 1229-1237.	1.5	15
117	Synchronous genitourinary lichen sclerosus signals a distinct urinary microbiome profile in men with urethral stricture disease. World Journal of Urology, 2021, 39, 605-611.	1.2	13
118	Microscopic Colitis Patients Possess a Perturbed and Inflammatory Gut Microbiota. Digestive Diseases and Sciences, 2022, 67, 2433-2443.	1.1	13
119	Infant gut bacterial community composition and foodâ€related manifestation of atopy in early childhood. Pediatric Allergy and Immunology, 2022, 33, .	1.1	13
120	Maternal prenatal immunity, neonatal trained immunity, and early airway microbiota shape childhood asthma development. Allergy: European Journal of Allergy and Clinical Immunology, 2022, 77, 3617-3628.	2.7	13
121	Unconjugated bilirubin is associated with protection from early-life wheeze and childhood asthma. Journal of Allergy and Clinical Immunology, 2021, 148, 128-138.	1.5	12
122	Microbiome–Immune Interactions in Allergy and Asthma. Journal of Allergy and Clinical Immunology: in Practice, 2022, 10, 2244-2251.	2.0	12
123	Dual epithelial and immune cell function of Dvl1 regulates gut microbiota composition and intestinal homeostasis. JCI Insight, 2016, 1, .	2.3	11
124	Moraxella-dominated pediatric nasopharyngeal microbiota associate with upper respiratory infection and sinusitis. PLoS ONE, 2021, 16, e0261179.	1.1	11
125	From Microbe to Microbiota: Considering Microbial Community Composition in Infections and Airway Diseases. American Journal of Respiratory and Critical Care Medicine, 2012, 185, 691-692.	2.5	10
126	Does Pet-Keeping Modify the Association of Delivery Mode with Offspring Body Size?. Maternal and Child Health Journal, 2015, 19, 1426-1433.	0.7	10

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127	Nucleic Acid Extraction Efficiency and Bacterial Recovery from Maxillary Sinus Mucosal Samples Obtained by Brushing or Biopsy. American Journal of Rhinology and Allergy, 2010, 24, 263-265.	1.0	9
128	Rearrangement of a Large Novel Pseudomonas aeruginosa Gene Island in Strains Isolated from a Patient Developing Ventilator-Associated Pneumonia. Journal of Clinical Microbiology, 2014, 52, 2430-2438.	1.8	9
129	Role of the lung microbiome in HIV pathogenesis. Current Opinion in HIV and AIDS, 2018, 13, 45-52.	1.5	9
130	Maternal and cord blood vitamin D level and the infant gut microbiota in a birth cohort study. Maternal Health, Neonatology and Perinatology, 2020, 6, 5.	1.0	9
131	Development of a standardized approach for environmental microbiota investigations related to asthma development in children. Journal of Microbiological Methods, 2012, 91, 231-239.	0.7	8
132	Intestinal inflammation alters the antigen-specific immune response to a skin commensal. Cell Reports, 2022, 39, 110891.	2.9	8
133	Motility and biofilm formation of the emerging gastrointestinal pathogen <i>Campylobacter concisus </i> differs under microaerophilic and anaerobic environments. Gut Microbes, 2019, 10, 34-44.	4.3	7
134	Distinct lung microbiota associate with HIV-associated chronic lung disease in children. Scientific Reports, 2020, 10, 16186.	1.6	7
135	Pneumonia surveillance with culture-independent metatranscriptomics in HIV-positive adults in Uganda: a cross-sectional study. Lancet Microbe, The, 2022, 3, e357-e365.	3.4	7
136	A20 in dendritic cells restrains intestinal anti-bacterial peptide expression and preserves commensal homeostasis. PLoS ONE, 2019, 14, e0218999.	1.1	6
137	Associations of physical activity with gut microbiota in pre-adolescent children. Physical Activity and Nutrition, 2021, 25, 24-37.	0.4	6
138	Early-life gut microbiota and attention deficit hyperactivity disorder in preadolescents. Pediatric Research, 2023, 93, 2051-2060.	1.1	5
139	Race-Specific Association of Caesarean-Section Delivery with Body Size at Age 2 Years. Ethnicity and Disease, 2016, 26, 61.	1.0	4
140	Prebiotic to Improve Calcium Absorption in Postmenopausal Women After Gastric Bypass: A Randomized Controlled Trial. Journal of Clinical Endocrinology and Metabolism, 2022, 107, 1053-1064.	1.8	4
141	Forced evolution of <i>Escherichia coli </i> cells with the ability to effectively utilize non-natural amino acids <scp> </scp> -tert-leucine, <scp> </scp> -norleucine and \hat{I}^3 -methyl- <scp> </scp> -leucine. Biocatalysis and Biotransformation, 2010, 28, 293-303.	1.1	3
142	Rules of engagement in the gut microbiome. Nature Medicine, 2018, 24, 1642-1644.	15.2	3
143	Gut Microbial Regulation of Autism Spectrum Disorder Symptoms. Trends in Endocrinology and Metabolism, 2020, 31, 809-811.	3.1	2
144	Fungus fuels mucosal wounds in Crohn's disease. Immunity, 2021, 54, 856-858.	6.6	1

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145	Effect of Early Infant Probiotic Supplementation on Eczema, Asthma, and Rhinitis at 7 Years of Age. Pediatrics, 2022, , .	1.0	1
146	Relationship between Bacterial Colonization of Human Digestive and Respiratory Tract. World Review of Nutrition and Dietetics, 2013, , 64-71.	0.1	0