Laurie E Comstock

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2,078 20 45 g-index

46 2,647 14.2 5.47 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
43	The Host Shapes the Gut Microbiota via Fecal MicroRNA. <i>Cell Host and Microbe</i> , 2016 , 19, 32-43	23.4	394
42	Extensive surface diversity of a commensal microorganism by multiple DNA inversions. <i>Nature</i> , 2001 , 414, 555-8	50.4	251
41	An ecological network of polysaccharide utilization among human intestinal symbionts. <i>Current Biology</i> , 2014 , 24, 40-49	6.3	240
40	Bacterial glycans: key mediators of diverse host immune responses. <i>Cell</i> , 2006 , 126, 847-50	56.2	141
39	Bacterial antagonism in host-associated microbial communities. <i>Science</i> , 2018 , 361,	33.3	128
38	Bacteroides fragilis type VI secretion systems use novel effector and immunity proteins to antagonize human gut Bacteroidales species. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 3627-32	11.5	123
37	Type VI secretion systems of human gut Bacteroidales segregate into three genetic architectures, two of which are contained on mobile genetic elements. <i>BMC Genomics</i> , 2016 , 17, 58	4.5	91
36	Longitudinal analysis of the prevalence, maintenance, and IgA response to species of the order Bacteroidales in the human gut. <i>Infection and Immunity</i> , 2011 , 79, 2012-20	3.7	88
35	Evidence of extensive DNA transfer between bacteroidales species within the human gut. <i>MBio</i> , 2014 , 5, e01305-14	7.8	84
34	Importance of glycans to the host-bacteroides mutualism in the mammalian intestine. <i>Cell Host and Microbe</i> , 2009 , 5, 522-6	23.4	79
33	Bacteroides thetaiotaomicron: a dynamic, niche-adapted human symbiont. <i>BioEssays</i> , 2003 , 25, 926-9	4.1	71
32	An antimicrobial protein of the gut symbiont Bacteroides fragilis with a MACPF domain of host immune proteins. <i>Molecular Microbiology</i> , 2014 , 94, 1361-74	4.1	51
31	Bacteroidales Secreted Antimicrobial Proteins Target Surface Molecules Necessary for Gut Colonization and Mediate Competition In Vivo. <i>MBio</i> , 2016 , 7,	7.8	44
30	Phylum-wide general protein O-glycosylation system of the Bacteroidetes. <i>Molecular Microbiology</i> , 2013 , 88, 772-83	4.1	44
29	Bacteroides fragilis NCTC9343 produces at least three distinct capsular polysaccharides: cloning, characterization, and reassignment of polysaccharide B and C biosynthesis loci. <i>Infection and Immunity</i> , 2000 , 68, 6176-81	3.7	42
28	Streamlined Genetic Manipulation of Diverse and Isolates from the Human Gut Microbiota. <i>MBio</i> , 2019 , 10,	7.8	34
27	Type VI Secretion Systems and the Gut Microbiota. <i>Microbiology Spectrum</i> , 2019 , 7,	8.9	33

(2000-2017)

26	Gut Symbiont Secretes a Eukaryotic-Like Ubiquitin Protein That Mediates Intraspecies Antagonism. <i>MBio</i> , 2017 , 8,	7.8	25
25	The evolution of the type VI secretion system as a disintegration weapon. <i>PLoS Biology</i> , 2020 , 18, e300	0732,0	24
24	A family of anti-Bacteroidales peptide toxins wide-spread in the human gut microbiota. <i>Nature Communications</i> , 2019 , 10, 3460	17.4	22
23	Acquisition of MACPF domain-encoding genes is the main contributor to LPS glycan diversity in gut Bacteroides species. <i>ISME Journal</i> , 2018 , 12, 2919-2928	11.9	11
22	Genetic and Biochemical Analysis of Anaerobic Respiration in Bacteroides fragilis and Its Importance. <i>MBio</i> , 2020 , 11,	7.8	10
21	Identification of a Fifth Antibacterial Toxin Produced by a Single Bacteroides fragilis Strain. <i>Journal of Bacteriology</i> , 2019 , 201,	3.5	9
20	Mobile Type VI secretion system loci of the gut Bacteroidales display extensive intra-ecosystem transfer, multi-species spread and geographical clustering. <i>PLoS Genetics</i> , 2021 , 17, e1009541	6	7
19	The evolution of tit-for-tat in bacteria via the type VI secretion system. <i>Nature Communications</i> , 2020 , 11, 5395	17.4	5
18	Nanaerobic growth enables direct visualization of dynamic cellular processes in human gut symbionts. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 24484-24493	11.5	5
17	Small RNAs Repress Expression of Polysaccharide Utilization Loci of Gut Bacteroides Species. Journal of Bacteriology, 2016 , 198, 2396-8	3.5	4
16	Friend turned foe: a role for bacterial sulfatases in colitis. Cell Host and Microbe, 2015, 17, 540-1	23.4	3
15	Analysis of a phase-variable restriction modification system of the human gut symbiont Bacteroides fragilis. <i>Nucleic Acids Research</i> , 2020 , 48, 11040-11053	20.1	2
14	Genetic Diversity of the Capsular Polysaccharide C Biosynthesis Region of Bacteroides fragilis. <i>Infection and Immunity</i> , 2000 , 68, 6182-6188	3.7	2
13	A New Pillar in Pilus Assembly. <i>Cell</i> , 2016 , 165, 520-1	56.2	2
12	Abscesses397-408		1
11	Utilizing Ribose Compounds: How Bacteroides PUL It Off. Cell Host and Microbe, 2020 , 27, 6-8	23.4	1
10	Type VI Secretion Systems and the Gut Microbiota 2019 , 343-350		1
9	Bacteroides fragilis NCTC9343 Produces at Least Three Distinct Capsular Polysaccharides: Cloning, Characterization, and Reassignment of Polysaccharide B and C Biosynthesis Loci. <i>Infection and Immunity</i> , 2000 , 68, 6176-6181	3.7	О

Bacteroidetocins Target the Essential Outer Membrane Protein BamA of Symbionts and Pathogens. MBio, 2021, 12, e0228521

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The evolution of the type VI secretion system as a disintegration weapon 2020, 18, e3000720

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