Lilan Ling

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/938696/publications.pdf

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24 papers	6,320 citations	304743 22 h-index	610901 24 g-index
24 all docs	24 docs citations	24 times ranked	8924 citing authors

#	Article	IF	Citations
1	Gut microbiota dysbiosis and diarrhea in kidney transplant recipients. American Journal of Transplantation, 2019, 19, 488-500.	4.7	70
2	Genome-Wide Screening for Enteric Colonization Factors in Carbapenem-Resistant ST258 Klebsiella pneumoniae. MBio, 2019, 10, .	4.1	32
3	Minimal residual disease negativity in multiple myeloma is associated with intestinal microbiota composition. Blood Advances, 2019, 3, 2040-2044.	5.2	50
4	Gut uropathogen abundance is a risk factor for development of bacteriuria and urinary tract infection. Nature Communications, 2019, 10, 5521.	12.8	123
5	Impact of gut colonization with butyrate producing microbiota on respiratory viral infection following allo-HCT. Blood, 2018, 131, blood-2018-01-828996.	1.4	155
6	Reconstitution of the gut microbiota of antibiotic-treated patients by autologous fecal microbiota transplant. Science Translational Medicine, 2018, 10, .	12.4	258
7	The effects of amine-modified single-walled carbon nanotubes on the mouse microbiota. International Journal of Nanomedicine, 2018, Volume 13, 5275-5286.	6.7	2
8	Cooperating Commensals Restore Colonization Resistance to Vancomycin-Resistant Enterococcus faecium. Cell Host and Microbe, 2017, 21, 592-602.e4.	11.0	237
9	Commensal microbes provide first line defense against Listeria monocytogenes infection. Journal of Experimental Medicine, 2017, 214, 1973-1989.	8.5	173
10	Protective Factors in the Intestinal Microbiome Against Clostridium difficile Infection in Recipients of Allogeneic Hematopoietic Stem Cell Transplantation. Journal of Infectious Diseases, 2017, 215, 1117-1123.	4.0	81
11	The oral microbiota in patients with pancreatic cancer, patients with IPMNs, and controls: a pilot study. Cancer Causes and Control, 2017, 28, 959-969.	1.8	69
12	Pathogenicity Locus, Core Genome, and Accessory Gene Contributions to Clostridium difficile Virulence. MBio, 2017, 8, .	4.1	51
13	Intestinal Microbiota and Relapse After Hematopoietic-Cell Transplantation. Journal of Clinical Oncology, 2017, 35, 1650-1659.	1.6	252
14	Complete Genome Sequence of Enterococcus faecium ATCC 700221. Genome Announcements, 2016, 4, .	0.8	9
15	Commensal microbiota affects ischemic stroke outcome by regulating intestinal Î³Î´ T cells. Nature Medicine, 2016, 22, 516-523.	30.7	770
16	Distinct but Spatially Overlapping Intestinal Niches for Vancomycin-Resistant Enterococcus faecium and Carbapenem-Resistant Klebsiella pneumoniae. PLoS Pathogens, 2015, 11, e1005132.	4.7	100
17	Innate Immune Defenses Mediated by Two ILC Subsets Are Critical for Protection against Acute Clostridium difficile Infection. Cell Host and Microbe, 2015, 18, 27-37.	11.0	240
18	Intestinal Blautia Is Associated with Reduced Death from Graft-versus-Host Disease. Biology of Blood and Marrow Transplantation, 2015, 21, 1373-1383.	2.0	619

#	Article	IF	Citations
19	Loss of Microbiota-Mediated Colonization Resistance toClostridium difficileInfection With Oral Vancomycin Compared With Metronidazole. Journal of Infectious Diseases, 2015, 212, 1656-1665.	4.0	157
20	Precision microbiome reconstitution restores bile acid mediated resistance to Clostridium difficile. Nature, 2015, 517, 205-208.	27.8	1,506
21	Gut Microbiota and Tacrolimus Dosing in Kidney Transplantation. PLoS ONE, 2015, 10, e0122399.	2.5	133
22	Gut Microbial Community Structure and Complications After Kidney Transplantation. Transplantation, 2014, 98, 697-705.	1.0	131
23	The effects of intestinal tract bacterial diversity on mortality following allogeneic hematopoietic stem cell transplantation. Blood, 2014, 124, 1174-1182.	1.4	711
24	Intestinal Microbiota Containing Barnesiella Species Cures Vancomycin-Resistant Enterococcus faecium Colonization. Infection and Immunity, 2013, 81, 965-973.	2.2	391