

Naga Betrapally

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5908865/publications.pdf>

Version: 2024-02-01

17
papers

1,356
citations

687363

13
h-index

996975

15
g-index

17
all docs

17
docs citations

17
times ranked

2233
citing authors

#	ARTICLE	IF	CITATIONS
1	Salivary microbiota reflects changes in gut microbiota in cirrhosis with hepatic encephalopathy. <i>Hepatology</i> , 2015, 62, 1260-1271.	7.3	272
2	Impaired Gut-Liver-Brain Axis in Patients with Cirrhosis. <i>Scientific Reports</i> , 2016, 6, 26800.	3.3	163
3	Chronic opioid use is associated with altered gut microbiota and predicts readmissions in patients with cirrhosis. <i>Alimentary Pharmacology and Therapeutics</i> , 2017, 45, 319-331.	3.7	149
4	Systems biology analysis of omeprazole therapy in cirrhosis demonstrates significant shifts in gut microbiota composition and function. <i>American Journal of Physiology - Renal Physiology</i> , 2014, 307, G951-G957.	3.4	125
5	Changes in the Intestinal Microbiome and Alcoholic and Nonalcoholic Liver Diseases: Causes or Effects?. <i>Gastroenterology</i> , 2016, 150, 1745-1755.e3.	1.3	104
6	Decompensated cirrhosis and microbiome interpretation. <i>Nature</i> , 2015, 525, E1-E2.	27.8	90
7	Gut microbiota drive the development of neuroinflammatory response in cirrhosis in mice. <i>Hepatology</i> , 2016, 64, 1232-1248.	7.3	83
8	Gut microbiome and liver disease. <i>Translational Research</i> , 2017, 179, 49-59.	5.0	78
9	Rifaximin Exerts Beneficial Effects Independent of its Ability to Alter Microbiota Composition. <i>Clinical and Translational Gastroenterology</i> , 2016, 7, e187.	2.5	75
10	Gut Microbiota Alterations can predict Hospitalizations in Cirrhosis Independent of Diabetes Mellitus. <i>Scientific Reports</i> , 2015, 5, 18559.	3.3	74
11	HCV eradication does not impact gut dysbiosis or systemic inflammation in cirrhotic patients. <i>Alimentary Pharmacology and Therapeutics</i> , 2016, 44, 638-643.	3.7	53
12	Whole-gene analysis of inter-genogroup reassortant rotaviruses from the Dominican Republic: Emergence of equine-like G3 strains and evidence of their reassortment with locally-circulating strains. <i>Virology</i> , 2019, 534, 114-131.	2.4	38
13	Gut microbial composition can differentially regulate bile acid synthesis in humanized mice. <i>Hepatology Communications</i> , 2017, 1, 61-70.	4.3	35
14	Metagenomic sequencing generates the whole genomes of porcine rotavirus A, C, and H from the United States. <i>PLoS ONE</i> , 2020, 15, e0244498.	2.5	14
15	Distribution of rotavirus genotypes in the postvaccine introduction era in Ashaiman, Greater Accra Region, Ghana, 2014-2016. <i>Journal of Medical Virology</i> , 2019, 91, 2025-2028.	5.0	2
16	The Beneficial Impact of Rifaximin on Systemic and Intestinal Inflammation and Ammonia Occurs even without Microbiota: More than an Antibiotic. <i>Journal of Hepatology</i> , 2016, 64, S447.	3.7	1
17	280 The Beneficial Impact of Rifaximin on Systemic and Intestinal Inflammation and Ammonia Occurs Even Without Microbiota: More Than an Antibiotic. <i>Gastroenterology</i> , 2016, 150, S1022-S1023.	1.3	0