

Martin Kummen

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

29
papers

1,219
citations

17
h-index

32
g-index

32
ext. papers

1,675
ext. citations

7.5
avg, IF

4.1
L-index

#	Paper	IF	Citations
29	Probiotics to HIV-Infected Immunological Nonresponders: Altered Mucosal Immunity and Microbial Diversity Restricted to Ileum. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2022 , 89, 77-86	3.1	0
28	Altered Gut Microbial Metabolism of Essential Nutrients in Primary Sclerosing Cholangitis. <i>Gastroenterology</i> , 2021 , 160, 1784-1798.e0	13.3	17
27	Mortality and microbial diversity after allogeneic hematopoietic stem cell transplantation: secondary analysis of a randomized nutritional intervention trial. <i>Scientific Reports</i> , 2021 , 11, 11593	4.9	1
26	Associations of neopterin and kynurenine-tryptophan ratio with survival in primary sclerosing cholangitis. <i>Scandinavian Journal of Gastroenterology</i> , 2021 , 56, 443-452	2.4	4
25	NLRP3 inflammasome deficiency attenuates metabolic disturbances involving alterations in the gut microbial profile in mice exposed to high fat diet. <i>Scientific Reports</i> , 2020 , 10, 21006	4.9	5
24	Low fibre intake is associated with gut microbiota alterations in chronic heart failure. <i>ESC Heart Failure</i> , 2020 , 7, 456-466	3.7	19
23	Gut mycobiome of primary sclerosing cholangitis patients is characterised by an increase of and species. <i>Gut</i> , 2020 , 69, 1890-1892	19.2	6
22	Rosuvastatin alters the genetic composition of the human gut microbiome. <i>Scientific Reports</i> , 2020 , 10, 5397	4.9	8
21	HIV-infected immunological non-responders have colon-restricted gut mucosal immune dysfunction. <i>Journal of Infectious Diseases</i> , 2020 ,	7	5
20	Gut Microbiota-Dependent Trimethylamine N-Oxide Associates With Inflammation in Common Variable Immunodeficiency. <i>Frontiers in Immunology</i> , 2020 , 11, 574500	8.4	16
19	Rifaximin alters gut microbiota profile, but does not affect systemic inflammation - a randomized controlled trial in common variable immunodeficiency. <i>Scientific Reports</i> , 2019 , 9, 167	4.9	23
18	The gut microbial influence on cholestatic liver disease. <i>Liver International</i> , 2019 , 39, 1186-1196	7.9	24
17	Autotaxin activity predicts transplant-free survival in primary sclerosing cholangitis. <i>Scientific Reports</i> , 2019 , 9, 8450	4.9	5
16	Selective IgA deficiency in humans is associated with reduced gut microbial diversity. <i>Journal of Allergy and Clinical Immunology</i> , 2019 , 143, 1969-1971.e11	11.5	22
15	Consistent alterations in faecal microbiomes of patients with primary sclerosing cholangitis independent of associated colitis. <i>Alimentary Pharmacology and Therapeutics</i> , 2019 , 50, 580-589	6.1	36
14	Circulating markers of gut barrier function associated with disease severity in primary sclerosing cholangitis. <i>Liver International</i> , 2019 , 39, 371-381	7.9	22
13	Gut Microbiota Signature in Heart Failure Defined From Profiling of Independent Cohorts. <i>Journal of the American College of Cardiology</i> , 2018 , 71, 1184-1186	15.1	79

12	Impact of HIV and Type 2 diabetes on Gut Microbiota Diversity, Tryptophan Catabolism and Endothelial Dysfunction. <i>Scientific Reports</i> , 2018 , 8, 6725	4.9	23
11	The gut microbial profile in patients with primary sclerosing cholangitis is distinct from patients with ulcerative colitis without biliary disease and healthy controls. <i>Gut</i> , 2017 , 66, 611-619	19.2	216
10	Intestinal microbiota in primary sclerosing cholangitis. <i>Current Opinion in Gastroenterology</i> , 2017 , 33, 85-92	3	17
9	Elevated trimethylamine--oxide (TMAO) is associated with poor prognosis in primary sclerosing cholangitis patients with normal liver function. <i>United European Gastroenterology Journal</i> , 2017 , 5, 532-541	5.3	13
8	Response to Faecal microbiota profiles as diagnostic biomarkers in primary sclerosing cholangitis by R Bleemann. <i>Gut</i> , 2017 , 66, 755-756	19.2	3
7	Guanylate Cyclase C Activation Shapes the Intestinal Microbiota in Patients with Familial Diarrhea and Increased Susceptibility for Crohn's Disease. <i>Inflammatory Bowel Diseases</i> , 2017 , 23, 1752-1761	4.5	7
6	Systemic sclerosis is associated with specific alterations in gastrointestinal microbiota in two independent cohorts. <i>BMJ Open Gastroenterology</i> , 2017 , 4, e000134	3.9	46
5	The gut microbiota contributes to a mouse model of spontaneous bile duct inflammation. <i>Journal of Hepatology</i> , 2017 , 66, 382-389	13.4	44
4	Altered gut microbiota profile in common variable immunodeficiency associates with levels of lipopolysaccharide and markers of systemic immune activation. <i>Mucosal Immunology</i> , 2016 , 9, 1455-1465	9.2	81
3	The Carnitine-butYRObetaine-trimethylamine-N-oxide pathway and its association with cardiovascular mortality in patients with carotid atherosclerosis. <i>Atherosclerosis</i> , 2016 , 247, 64-9	3.1	82
2	Genome-wide association analysis identifies variation in vitamin D receptor and other host factors influencing the gut microbiota. <i>Nature Genetics</i> , 2016 , 48, 1396-1406	36.3	369
1	Liver abnormalities in bowel diseases. <i>Baillieres Best Practice and Research in Clinical Gastroenterology</i> , 2013 , 27, 531-42	2.5	25