

# Martin Kummen

## List of Publications by Year in descending order

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Version: 2024-02-01

29  
papers

2,025  
citations

361413

20  
h-index

454955

30  
g-index

32  
all docs

32  
docs citations

32  
times ranked

3683  
citing authors

#	ARTICLE	IF	CITATIONS
1	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.	21.4	533
2	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.	12.1	308
3	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.	2.8	137
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.	6.0	130
5	The Carnitine-butyrobetaine-trimethylamine-N-oxide pathway and its association with cardiovascular mortality in patients with carotid atherosclerosis. <i>Atherosclerosis</i> , 2016, 247, 64-69.	0.8	116
6	Systemic sclerosis is associated with specific alterations in gastrointestinal microbiota in two independent cohorts. <i>BMJ Open Gastroenterology</i> , 2017, 4, e000134.	2.7	77
7	Altered Gut Microbial Metabolism of Essential Nutrients in Primary Sclerosing Cholangitis. <i>Gastroenterology</i> , 2021, 160, 1784-1798.e0.	1.3	69
8	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.	3.7	67
9	The gut microbiota contributes to a mouse model of spontaneous bile duct inflammation. <i>Journal of Hepatology</i> , 2017, 66, 382-389.	3.7	60
10	Low fibre intake is associated with gut microbiota alterations in chronic heart failure. <i>ESC Heart Failure</i> , 2020, 7, 456-466.	3.1	56
11	Circulating markers of gut barrier function associated with disease severity in primary sclerosing cholangitis. <i>Liver International</i> , 2019, 39, 371-381.	3.9	51
12	The gut microbial influence on cholestatic liver disease. <i>Liver International</i> , 2019, 39, 1186-1196.	3.9	46
13	Gut Microbiota-Dependent Trimethylamine N-Oxide Associates With Inflammation in Common Variable Immunodeficiency. <i>Frontiers in Immunology</i> , 2020, 11, 574500.	4.8	38
14	Impact of HIV and Type 2 diabetes on Gut Microbiota Diversity, Tryptophan Catabolism and Endothelial Dysfunction. <i>Scientific Reports</i> , 2018, 8, 6725.	3.3	35
15	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.	2.9	33
16	Protective and aggressive bacterial subsets and metabolites modify hepatobiliary inflammation and fibrosis in a murine model of PSC. <i>Gut</i> , 2023, 72, 671-685.	12.1	30
17	Liver abnormalities in bowel diseases. <i>Bailliere's Best Practice and Research in Clinical Gastroenterology</i> , 2013, 27, 531-542.	2.4	29
18	Gut mycobiome of primary sclerosing cholangitis patients is characterised by an increase of <i>Trichocladium griseum</i> and <i>Candida</i> species. <i>Gut</i> , 2020, 69, 1890-1892.	12.1	25

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19	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.	3.3	21
20	Intestinal microbiota in primary sclerosing cholangitis. <i>Current Opinion in Gastroenterology</i> , 2017, 33, 85-92.	2.3	20
21	Elevated trimethylamine- <i>N</i> -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.	3.8	20
22	Rosuvastatin alters the genetic composition of the human gut microbiome. <i>Scientific Reports</i> , 2020, 10, 5397.	3.3	20
23	Human Immunodeficiency Virus-Infected Immunological Nonresponders Have Colon-Restricted Gut Mucosal Immune Dysfunction. <i>Journal of Infectious Diseases</i> , 2022, 225, 661-674.	4.0	16
24	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.	1.9	13
25	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.	3.3	9
26	Autotaxin activity predicts transplant-free survival in primary sclerosing cholangitis. <i>Scientific Reports</i> , 2019, 9, 8450.	3.3	8
27	Associations of neopterin and kynurenine-tryptophan ratio with survival in primary sclerosing cholangitis. <i>Scandinavian Journal of Gastroenterology</i> , 2021, 56, 443-452.	1.5	8
28	Response to Faecal microbiota profiles as diagnostic biomarkers in primary sclerosing cholangitis by Lehmann et al. <i>Gut</i> , 2017, 66, 755-756.	12.1	3
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.	2.1	3