

Martin Kummen

List of Publications by Citations

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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 | 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 |
| 28 | 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 |
| 27 | 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 |
| 26 | 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 |
| 25 | 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 |
| 24 | 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 |
| 23 | 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 |
| 22 | 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 |
| 21 | Liver abnormalities in bowel diseases. <i>Baillieres Best Practice and Research in Clinical Gastroenterology</i> , 2013 , 27, 531-42 | 2.5 | 25 |
| 20 | The gut microbial influence on cholestatic liver disease. <i>Liver International</i> , 2019 , 39, 1186-1196 | 7.9 | 24 |
| 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 | 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 |
| 17 | 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 |
| 16 | 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 |
| 15 | 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 |
| 14 | Intestinal microbiota in primary sclerosing cholangitis. <i>Current Opinion in Gastroenterology</i> , 2017 , 33, 85-92 | 3 | 17 |
| 13 | Altered Gut Microbial Metabolism of Essential Nutrients in Primary Sclerosing Cholangitis. <i>Gastroenterology</i> , 2021 , 160, 1784-1798.e0 | 13.3 | 17 |

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|----|---|------|----|
| 12 | Gut Microbiota-Dependent Trimethylamine N-Oxide Associates With Inflammation in Common Variable Immunodeficiency. <i>Frontiers in Immunology</i> , 2020 , 11, 574500 | 8.4 | 16 |
| 11 | 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 |
| 10 | Rosuvastatin alters the genetic composition of the human gut microbiome. <i>Scientific Reports</i> , 2020 , 10, 5397 | 4.9 | 8 |
| 9 | Guanylate Cyclase C Activation Shapes the Intestinal Microbiota in Patients with Familial Diarrhea and Increased Susceptibility for Crohns Disease. <i>Inflammatory Bowel Diseases</i> , 2017 , 23, 1752-1761 | 4.5 | 7 |
| 8 | 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 |
| 7 | Autotaxin activity predicts transplant-free survival in primary sclerosing cholangitis. <i>Scientific Reports</i> , 2019 , 9, 8450 | 4.9 | 5 |
| 6 | 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 |
| 5 | HIV-infected immunological non-responders have colon-restricted gut mucosal immune dysfunction. <i>Journal of Infectious Diseases</i> , 2020 , | 7 | 5 |
| 4 | 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 |
| 3 | Response to Faecal microbiota profiles as diagnostic biomarkers in primary sclerosing cholangitis by RBlmann. <i>Gut</i> , 2017 , 66, 755-756 | 19.2 | 3 |
| 2 | 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 |
| 1 | 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 |