

Ramnik J Xavier

List of Publications by Year in descending order

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Version: 2024-02-01

270
papers

82,810
citations

807

118
h-index

481

270
g-index

302
all docs

302
docs citations

302
times ranked

84871
citing authors

#	ARTICLE	IF	CITATIONS
1	Host-microbe interactions have shaped the genetic architecture of inflammatory bowel disease. <i>Nature</i> , 2012, 491, 119-124.	13.7	4,038
2	Guidelines for the use and interpretation of assays for monitoring autophagy. <i>Autophagy</i> , 2012, 8, 445-544.	4.3	3,122
3	The Treatment-Naive Microbiome in New-Onset Crohn's Disease. <i>Cell Host and Microbe</i> , 2014, 15, 382-392.	5.1	2,582
4	Genome-wide association defines more than 30 distinct susceptibility loci for Crohn's disease. <i>Nature Genetics</i> , 2008, 40, 955-962.	9.4	2,422
5	Dysfunction of the intestinal microbiome in inflammatory bowel disease and treatment. <i>Genome Biology</i> , 2012, 13, R79.	13.9	2,258
6	Genetics and pathogenesis of inflammatory bowel disease. <i>Nature</i> , 2011, 474, 307-317.	13.7	2,040
7	Trained immunity: A program of innate immune memory in health and disease. <i>Science</i> , 2016, 352, aaf1098.	6.0	1,809
8	Multi-omics of the gut microbial ecosystem in inflammatory bowel diseases. <i>Nature</i> , 2019, 569, 655-662.	13.7	1,638
9	Genome-wide association study identifies new susceptibility loci for Crohn disease and implicates autophagy in disease pathogenesis. <i>Nature Genetics</i> , 2007, 39, 596-604.	9.4	1,633
10	mTOR- and HIF-1 α -mediated aerobic glycolysis as metabolic basis for trained immunity. <i>Science</i> , 2014, 345, 1250684.	6.0	1,517
11	Population-based metagenomics analysis reveals markers for gut microbiome composition and diversity. <i>Science</i> , 2016, 352, 565-569.	6.0	1,398
12	Succinate Dehydrogenase Supports Metabolic Repurposing of Mitochondria to Drive Inflammatory Macrophages. <i>Cell</i> , 2016, 167, 457-470.e13.	13.5	1,396
13	The Microbiome in Inflammatory Bowel Disease: Current Status and the Future Ahead. <i>Gastroenterology</i> , 2014, 146, 1489-1499.	0.6	1,374
14	Rational design of highly active sgRNAs for CRISPR-Cas9-mediated gene inactivation. <i>Nature Biotechnology</i> , 2014, 32, 1262-1267.	9.4	1,351
15	Defining trained immunity and its role in health and disease. <i>Nature Reviews Immunology</i> , 2020, 20, 375-388.	10.6	1,345
16	A key role for autophagy and the autophagy gene Atg16l1 in mouse and human intestinal Paneth cells. <i>Nature</i> , 2008, 456, 259-263.	13.7	1,341
17	Epigenetic programming of monocyte-to-macrophage differentiation and trained innate immunity. <i>Science</i> , 2014, 345, 1251086.	6.0	1,338
18	Bacille Calmette-Guérin induces NOD2-dependent nonspecific protection from reinfection via epigenetic reprogramming of monocytes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 17537-17542.	3.3	1,294

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19	Meta-analysis identifies 29 additional ulcerative colitis risk loci, increasing the number of confirmed associations to 47. <i>Nature Genetics</i> , 2011, 43, 246-252.	9.4	1,201
20	A single-cell survey of the small intestinal epithelium. <i>Nature</i> , 2017, 551, 333-339.	13.7	1,197
21	Gut microbiome structure and metabolic activity in inflammatory bowel disease. <i>Nature Microbiology</i> , 2019, 4, 293-305.	5.9	1,094
22	Disease-Specific Alterations in the Enteric Virome in Inflammatory Bowel Disease. <i>Cell</i> , 2015, 160, 447-460.	13.5	1,036
23	The Dynamics of the Human Infant Gut Microbiome in Development and in Progression toward Type 1 Diabetes. <i>Cell Host and Microbe</i> , 2015, 17, 260-273.	5.1	1,008
24	CARD9 impacts colitis by altering gut microbiota metabolism of tryptophan into aryl hydrocarbon receptor ligands. <i>Nature Medicine</i> , 2016, 22, 598-605.	15.2	1,001
25	Variation in Microbiome LPS Immunogenicity Contributes to Autoimmunity in Humans. <i>Cell</i> , 2016, 165, 842-853.	13.5	968
26	<i>Candida albicans</i> Infection Affords Protection against Reinfection via Functional Reprogramming of Monocytes. <i>Cell Host and Microbe</i> , 2012, 12, 223-232.	5.1	926
27	Pyruvate Kinase M2 Regulates Hif-1 α Activity and IL-1 β Induction and Is a Critical Determinant of the Warburg Effect in LPS-Activated Macrophages. <i>Cell Metabolism</i> , 2015, 21, 65-80.	7.2	887
28	Proton pump inhibitors affect the gut microbiome. <i>Gut</i> , 2016, 65, 740-748.	6.1	885
29	Identification of a Molecular Signaling Network that Regulates a Cellular Necrotic Cell Death Pathway. <i>Cell</i> , 2008, 135, 1311-1323.	13.5	878
30	BCG Vaccination Protects against Experimental Viral Infection in Humans through the Induction of Cytokines Associated with Trained Immunity. <i>Cell Host and Microbe</i> , 2018, 23, 89-100.e5.	5.1	860
31	Mother-to-Infant Microbial Transmission from Different Body Sites Shapes the Developing Infant Gut Microbiome. <i>Cell Host and Microbe</i> , 2018, 24, 133-145.e5.	5.1	822
32	Virus-Plus-Susceptibility Gene Interaction Determines Crohn's Disease Gene Atg16L1 Phenotypes in Intestine. <i>Cell</i> , 2010, 141, 1135-1145.	13.5	809
33	Linking the Human Gut Microbiome to Inflammatory Cytokine Production Capacity. <i>Cell</i> , 2016, 167, 1125-1136.e8.	13.5	806
34	Intra- and Inter-cellular Rewiring of the Human Colon during Ulcerative Colitis. <i>Cell</i> , 2019, 178, 714-730.e22.	13.5	806
35	Natural history of the infant gut microbiome and impact of antibiotic treatment on bacterial strain diversity and stability. <i>Science Translational Medicine</i> , 2016, 8, 343ra81.	5.8	763
36	A defined commensal consortium elicits CD8 T cells and anti-cancer immunity. <i>Nature</i> , 2019, 565, 600-605.	13.7	741

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37	Deep resequencing of GWAS loci identifies independent rare variants associated with inflammatory bowel disease. <i>Nature Genetics</i> , 2011, 43, 1066-1073.	9.4	698
38	Identification of a candidate therapeutic autophagy-inducing peptide. <i>Nature</i> , 2013, 494, 201-206.	13.7	669
39	Ectopic colonization of oral bacteria in the intestine drives T _H 1 cell induction and inflammation. <i>Science</i> , 2017, 358, 359-365.	6.0	612
40	Deletion polymorphism upstream of IRGM associated with altered IRGM expression and Crohn's disease. <i>Nature Genetics</i> , 2008, 40, 1107-1112.	9.4	604
41	Interplay of host genetics and gut microbiota underlying the onset and clinical presentation of inflammatory bowel disease. <i>Gut</i> , 2018, 67, 108-119.	6.1	590
42	Glutaminolysis and Fumarate Accumulation Integrate Immunometabolic and Epigenetic Programs in Trained Immunity. <i>Cell Metabolism</i> , 2016, 24, 807-819.	7.2	584
43	Genome-wide association identifies multiple ulcerative colitis susceptibility loci. <i>Nature Genetics</i> , 2010, 42, 332-337.	9.4	572
44	The Gut Microbiome Contributes to a Substantial Proportion of the Variation in Blood Lipids. <i>Circulation Research</i> , 2015, 117, 817-824.	2.0	534
45	Indoleacrylic Acid Produced by Commensal Peptostreptococcus Species Suppresses Inflammation. <i>Cell Host and Microbe</i> , 2017, 22, 25-37.e6.	5.1	523
46	Gene-microbiota interactions contribute to the pathogenesis of inflammatory bowel disease. <i>Science</i> , 2016, 352, 1116-1120.	6.0	498
47	Prediction of complicated disease course for children newly diagnosed with Crohn's disease: a multicentre inception cohort study. <i>Lancet, The</i> , 2017, 389, 1710-1718.	6.3	482
48	Long-Lasting Effects of BCG Vaccination on Both Heterologous Th1/Th17 Responses and Innate Trained Immunity. <i>Journal of Innate Immunity</i> , 2014, 6, 152-158.	1.8	478
49	A novel <i>Ruminococcus gnavus</i> clade enriched in inflammatory bowel disease patients. <i>Genome Medicine</i> , 2017, 9, 103.	3.6	478
50	Fine-mapping inflammatory bowel disease loci to single-variant resolution. <i>Nature</i> , 2017, 547, 173-178.	13.7	473
51	Common variants at five new loci associated with early-onset inflammatory bowel disease. <i>Nature Genetics</i> , 2009, 41, 1335-1340.	9.4	459
52	Microbial genes and pathways in inflammatory bowel disease. <i>Nature Reviews Microbiology</i> , 2019, 17, 497-511.	13.6	447
53	Pediatric Crohn disease patients exhibit specific ileal transcriptome and microbiome signature. <i>Journal of Clinical Investigation</i> , 2014, 124, 3617-3633.	3.9	431
54	Dynamics of metatranscription in the inflammatory bowel disease gut microbiome. <i>Nature Microbiology</i> , 2018, 3, 337-346.	5.9	408

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55	Pathway paradigms revealed from the genetics of inflammatory bowel disease. <i>Nature</i> , 2020, 578, 527-539.	13.7	408
56	T Helper Cell Cytokines Modulate Intestinal Stem Cell Renewal and Differentiation. <i>Cell</i> , 2018, 175, 1307-1320.e22.	13.5	388
57	Advances in inflammatory bowel disease pathogenesis: linking host genetics and the microbiome. <i>Gut</i> , 2013, 62, 1505-1510.	6.1	387
58	Human genetic variation and the gut microbiome in disease. <i>Nature Reviews Genetics</i> , 2017, 18, 690-699.	7.7	383
59	Growth dynamics of gut microbiota in health and disease inferred from single metagenomic samples. <i>Science</i> , 2015, 349, 1101-1106.	6.0	382
60	Reovirus infection triggers inflammatory responses to dietary antigens and development of celiac disease. <i>Science</i> , 2017, 356, 44-50.	6.0	367
61	Host and Environmental Factors Influencing Individual Human Cytokine Responses. <i>Cell</i> , 2016, 167, 1111-1124.e13.	13.5	364
62	BCG-induced trained immunity in NK cells: Role for non-specific protection to infection. <i>Clinical Immunology</i> , 2014, 155, 213-219.	1.4	359
63	Impaired Aryl Hydrocarbon Receptor Ligand Production by the Gut Microbiota Is a Key Factor in Metabolic Syndrome. <i>Cell Metabolism</i> , 2018, 28, 737-749.e4.	7.2	356
64	Gut microbiota composition and functional changes in inflammatory bowel disease and irritable bowel syndrome. <i>Science Translational Medicine</i> , 2018, 10, .	5.8	351
65	<i>Akkermansia muciniphila</i> induces intestinal adaptive immune responses during homeostasis. <i>Science</i> , 2019, 364, 1179-1184.	6.0	347
66	Genome-wide enhancer maps link risk variants to disease genes. <i>Nature</i> , 2021, 593, 238-243.	13.7	332
67	Complex host genetics influence the microbiome in inflammatory bowel disease. <i>Genome Medicine</i> , 2014, 6, 107.	3.6	322
68	Global chemical effects of the microbiome include new bile-acid conjugations. <i>Nature</i> , 2020, 579, 123-129.	13.7	316
69	Strain-Level Analysis of Mother-to-Child Bacterial Transmission during the First Few Months of Life. <i>Cell Host and Microbe</i> , 2018, 24, 146-154.e4.	5.1	311
70	Gut Microbiome Function Predicts Response to Anti-integrin Biologic Therapy in Inflammatory Bowel Diseases. <i>Cell Host and Microbe</i> , 2017, 21, 603-610.e3.	5.1	306
71	Structure-based protein function prediction using graph convolutional networks. <i>Nature Communications</i> , 2021, 12, 3168.	5.8	300
72	Impaired Autophagy of an Intracellular Pathogen Induced by a Crohn's Disease Associated ATG16L1 Variant. <i>PLoS ONE</i> , 2008, 3, e3391.	1.1	299

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73	Atg16L1 T300A variant decreases selective autophagy resulting in altered cytokine signaling and decreased antibacterial defense. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 7741-7746.	3.3	298
74	Strain Tracking Reveals the Determinants of Bacterial Engraftment in the Human Gut Following Fecal Microbiota Transplantation. Cell Host and Microbe, 2018, 23, 229-240.e5.	5.1	292
75	Adaptive Evolution within Gut Microbiomes of Healthy People. Cell Host and Microbe, 2019, 25, 656-667.e8.	5.1	289
76	The Human and Mouse Enteric Nervous System at Single-Cell Resolution. Cell, 2020, 182, 1606-1622.e23.	13.5	287
77	Inflammatory Bowel Disease as a Model for Translating the Microbiome. Immunity, 2014, 40, 843-854.	6.6	284
78	Autophagy and the Immune System. Annual Review of Immunology, 2012, 30, 611-646.	9.5	282
79	A Functional Genomics Approach to Understand Variation in Cytokine Production in Humans. Cell, 2016, 167, 1099-1110.e14.	13.5	275
80	Bacteroides-Derived Sphingolipids Are Critical for Maintaining Intestinal Homeostasis and Symbiosis. Cell Host and Microbe, 2019, 25, 668-680.e7.	5.1	274
81	Trained immunity, tolerance, priming and differentiation: distinct immunological processes. Nature Immunology, 2021, 22, 2-6.	7.0	274
82	Paneth cells secrete lysozyme via secretory autophagy during bacterial infection of the intestine. Science, 2017, 357, 1047-1052.	6.0	267
83	Pathogen Cell-to-Cell Variability Drives Heterogeneity in Host Immune Responses. Cell, 2015, 162, 1309-1321.	13.5	255
84	Novel bile acid biosynthetic pathways are enriched in the microbiome of centenarians. Nature, 2021, 599, 458-464.	13.7	251
85	Mutations causing medullary cystic kidney disease type 1 lie in a large VNTR in MUC1 missed by massively parallel sequencing. Nature Genetics, 2013, 45, 299-303.	9.4	237
86	An Integrative View of Microbiome-Host Interactions in Inflammatory Bowel Diseases. Cell Host and Microbe, 2015, 17, 577-591.	5.1	235
87	Circadian clock protein BMAL1 regulates IL-1 β in macrophages via NRF2. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E8460-E8468.	3.3	230
88	Intestinal virome changes precede autoimmunity in type I diabetes-susceptible children. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E6166-E6175.	3.3	227
89	Ulcerative colitis mucosal transcriptomes reveal mitochondriopathy and personalized mechanisms underlying disease severity and treatment response. Nature Communications, 2019, 10, 38.	5.8	215
90	Gut Microbiota Regulation of T Cells During Inflammation and Autoimmunity. Annual Review of Immunology, 2019, 37, 599-624.	9.5	214

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91	Atg16L1 is Required for Autophagy in Intestinal Epithelial Cells and Protection of Mice From Salmonella Infection. <i>Gastroenterology</i> , 2013, 145, 1347-1357.	0.6	211
92	Dysbiosis, inflammation, and response to treatment: a longitudinal study of pediatric subjects with newly diagnosed inflammatory bowel disease. <i>Genome Medicine</i> , 2016, 8, 75.	3.6	211
93	Human gut bacteria produce β -17-modulating bile acid metabolites. <i>Nature</i> , 2022, 603, 907-912.	13.7	210
94	Compositional and Temporal Changes in the Gut Microbiome of Pediatric Ulcerative Colitis Patients Are Linked to Disease Course. <i>Cell Host and Microbe</i> , 2018, 24, 600-610.e4.	5.1	193
95	Identification of <i>Atg5</i> -dependent transcriptional changes and increases in mitochondrial mass in <i>Atg5</i> -deficient T lymphocytes. <i>Autophagy</i> , 2009, 5, 625-635.	4.3	187
96	The Ubiquitin Ligase Smurf1 Functions in Selective Autophagy of Mycobacterium tuberculosis and Anti-tuberculous Host Defense. <i>Cell Host and Microbe</i> , 2017, 21, 59-72.	5.1	184
97	Predictive metabolomic profiling of microbial communities using amplicon or metagenomic sequences. <i>Nature Communications</i> , 2019, 10, 3136.	5.8	176
98	Increased Intestinal Microbial Diversity Following Fecal Microbiota Transplant for Active Crohn's Disease. <i>Inflammatory Bowel Diseases</i> , 2016, 22, 2182-2190.	0.9	175
99	A role for bacterial urease in gut dysbiosis and Crohn's disease. <i>Science Translational Medicine</i> , 2017, 9, .	5.8	171
100	Autophagy Controls BCG-Induced Trained Immunity and the Response to Intravesical BCG Therapy for Bladder Cancer. <i>PLoS Pathogens</i> , 2014, 10, e1004485.	2.1	167
101	Elevated rates of horizontal gene transfer in the industrialized human microbiome. <i>Cell</i> , 2021, 184, 2053-2067.e18.	13.5	167
102	Associations between host gene expression, the mucosal microbiome, and clinical outcome in the pelvic pouch of patients with inflammatory bowel disease. <i>Genome Biology</i> , 2015, 16, 67.	3.8	166
103	Genomic variation and strain-specific functional adaptation in the human gut microbiome during early life. <i>Nature Microbiology</i> , 2019, 4, 470-479.	5.9	164
104	Functional genomics identifies type I interferon pathway as central for host defense against <i>Candida albicans</i> . <i>Nature Communications</i> , 2013, 4, 1342.	5.8	157
105	Therapeutic Opportunities in Inflammatory Bowel Disease: Mechanistic Dissection of Host-Microbiome Relationships. <i>Cell</i> , 2019, 178, 1041-1056.	13.5	156
106	Genetic Variants Synthesize to Produce Paneth Cell Phenotypes That Define Subtypes of Crohn's Disease. <i>Gastroenterology</i> , 2014, 146, 200-209.	0.6	155
107	Differential Effects of Environmental and Genetic Factors on T and B Cell Immune Traits. <i>Cell Reports</i> , 2016, 17, 2474-2487.	2.9	154
108	Transcriptional Atlas of Intestinal Immune Cells Reveals that Neuropeptide \pm -CGRP Modulates Group 2 Innate Lymphoid Cell Responses. <i>Immunity</i> , 2019, 51, 696-708.e9.	6.6	154

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109	Cholesterol Metabolism by Uncultured Human Gut Bacteria Influences Host Cholesterol Level. <i>Cell Host and Microbe</i> , 2020, 28, 245-257.e6.	5.1	151
110	Inter-individual variability and genetic influences on cytokine responses to bacteria and fungi. <i>Nature Medicine</i> , 2016, 22, 952-960.	15.2	148
111	Calcitonin Gene-Related Peptide Negatively Regulates Alarmin-Driven Type 2 Innate Lymphoid Cell Responses. <i>Immunity</i> , 2019, 51, 709-723.e6.	6.6	144
112	GSDMD is critical for autoinflammatory pathology in a mouse model of Familial Mediterranean Fever. <i>Journal of Experimental Medicine</i> , 2018, 215, 1519-1529.	4.2	143
113	Spatial and Temporal Mapping of Human Innate Lymphoid Cells Reveals Elements of Tissue Specificity. <i>Immunity</i> , 2019, 50, 505-519.e4.	6.6	139
114	A Noncanonical Autophagy Pathway Restricts <i>Toxoplasma gondii</i> Growth in a Strain-Specific Manner in IFN- β -Activated Human Cells. <i>MBio</i> , 2015, 6, e01157-15.	1.8	137
115	Gut Microbial Associations to Plasma Metabolites Linked to Cardiovascular Phenotypes and Risk. <i>Circulation Research</i> , 2019, 124, 1808-1820.	2.0	137
116	Role of dietary fiber in the recovery of the human gut microbiome and its metabolome. <i>Cell Host and Microbe</i> , 2021, 29, 394-407.e5.	5.1	137
117	ATG5 regulates plasma cell differentiation. <i>Autophagy</i> , 2013, 9, 528-537.	4.3	134
118	Host genetic variation and its microbiome interactions within the Human Microbiome Project. <i>Genome Medicine</i> , 2018, 10, 6.	3.6	134
119	<sc>TMEM</sc> 41B is a novel regulator of autophagy and lipid mobilization. <i>EMBO Reports</i> , 2018, 19, .	2.0	134
120	Genome-wide association studies: a new window into immune-mediated diseases. <i>Nature Reviews Immunology</i> , 2008, 8, 631-643.	10.6	130
121	Beclin 2 Functions in Autophagy, Degradation of G Protein-Coupled Receptors, and Metabolism. <i>Cell</i> , 2013, 154, 1085-1099.	13.5	130
122	Autophagy-Independent Lysosomal Targeting Regulated by ULK1/2-FIP200 and ATG9. <i>Cell Reports</i> , 2017, 20, 2341-2356.	2.9	126
123	Mechanisms of Pediatric Inflammatory Bowel Disease. <i>Annual Review of Immunology</i> , 2016, 34, 31-64.	9.5	124
124	SIKs control osteocyte responses to parathyroid hormone. <i>Nature Communications</i> , 2016, 7, 13176.	5.8	124
125	Functional CRISPR screening identifies the ufmylation pathway as a regulator of SQSTM1/p62. <i>ELife</i> , 2016, 5, .	2.8	122
126	Clinical and biological predictors of response to standardised paediatric colitis therapy (PROTECT): a multicentre inception cohort study. <i>Lancet, The</i> , 2019, 393, 1708-1720.	6.3	121

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127	Autophagy is essential for cardiac morphogenesis during vertebrate development. <i>Autophagy</i> , 2014, 10, 572-587.	4.3	117
128	Epitope-specific immunotherapy targeting CD4-positive T cells in coeliac disease: two randomised, double-blind, placebo-controlled phase 1 studies. <i>The Lancet Gastroenterology and Hepatology</i> , 2017, 2, 479-493.	3.7	113
129	A Pleiotropic Missense Variant in SLC39A8 Is Associated With Crohn's Disease and Human Gut Microbiome Composition. <i>Gastroenterology</i> , 2016, 151, 724-732.	0.6	109
130	Genetic Coding Variant in GPR65 Alters Lysosomal pH and Links Lysosomal Dysfunction with Colitis Risk. <i>Immunity</i> , 2016, 44, 1392-1405.	6.6	106
131	Integration of multi-omics data and deep phenotyping enables prediction of cytokine responses. <i>Nature Immunology</i> , 2018, 19, 776-786.	7.0	103
132	Ubiquitin Ligase TRIM62 Regulates CARD9-Mediated Anti-fungal Immunity and Intestinal Inflammation. <i>Immunity</i> , 2015, 43, 715-726.	6.6	102
133	Single-cell analyses of Crohn's disease tissues reveal intestinal intraepithelial T cells heterogeneity and altered subset distributions. <i>Nature Communications</i> , 2021, 12, 1921.	5.8	96
134	<i>Clorf106</i> is a colitis risk gene that regulates stability of epithelial adherens junctions. <i>Science</i> , 2018, 359, 1161-1166.	6.0	95
135	Targeting innate immunity for tuberculosis vaccination. <i>Journal of Clinical Investigation</i> , 2019, 129, 3482-3491.	3.9	95
136	Discs large (Dlg1) complexes in lymphocyte activation. <i>Journal of Cell Biology</i> , 2004, 166, 173-178.	2.3	92
137	Salt-Inducible Kinases: Physiology, Regulation by cAMP, and Therapeutic Potential. <i>Trends in Endocrinology and Metabolism</i> , 2018, 29, 723-735.	3.1	92
138	Dectin-1-Dependent LC3 Recruitment to Phagosomes Enhances Fungicidal Activity in Macrophages. <i>Journal of Infectious Diseases</i> , 2014, 210, 1844-1854.	1.9	90
139	Invertible promoters mediate bacterial phase variation, antibiotic resistance, and host adaptation in the gut. <i>Science</i> , 2019, 363, 181-187.	6.0	85
140	Multi-omics reveal microbial determinants impacting responses to biologic therapies in inflammatory bowel disease. <i>Cell Host and Microbe</i> , 2021, 29, 1294-1304.e4.	5.1	85
141	Cytokine release and gastrointestinal symptoms after gluten challenge in celiac disease. <i>Science Advances</i> , 2019, 5, eaaw7756.	4.7	84
142	The Crohn's disease polymorphism, ATG16L1 T300A, alters the gut microbiota and enhances the local Th1/Th17 response. <i>ELife</i> , 2019, 8, .	2.8	84
143	From genetics of inflammatory bowel disease towards mechanistic insights. <i>Trends in Immunology</i> , 2013, 34, 371-378.	2.9	82
144	Gut microbial co-abundance networks show specificity in inflammatory bowel disease and obesity. <i>Nature Communications</i> , 2020, 11, 4018.	5.8	80

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145	Differential Effect of Genetic Burden on Disease Phenotypes in Crohn's Disease and Ulcerative Colitis: Analysis of a North American Cohort. <i>American Journal of Gastroenterology</i> , 2014, 109, 395-400.	0.2	77
146	Sex-Specific Regulation of Inflammation and Metabolic Syndrome in Obesity. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020, 40, 1787-1800.	1.1	77
147	Role of Autophagy in the Maintenance of Intestinal Homeostasis. <i>Gastroenterology</i> , 2015, 149, 553-562.	0.6	76
148	QRICH1 dictates the outcome of ER stress through transcriptional control of proteostasis. <i>Science</i> , 2021, 371, .	6.0	73
149	B cell genomics behind cross-neutralization of SARS-CoV-2 variants and SARS-CoV. <i>Cell</i> , 2021, 184, 3205-3221.e24.	13.5	73
150	RNF166 Determines Recruitment of Adaptor Proteins during Antibacterial Autophagy. <i>Cell Reports</i> , 2016, 17, 2183-2194.	2.9	72
151	The RIG-I-like helicase receptor MDA5 (IFIH1) is involved in the host defense against <i>Candida</i> infections. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2015, 34, 963-974.	1.3	69
152	Human Genome-Wide RNAi Screen Identifies an Essential Role for Inositol Pyrophosphates in Type-I Interferon Response. <i>PLoS Pathogens</i> , 2014, 10, e1003981.	2.1	68
153	Small-molecule screening identifies inhibition of salt-inducible kinases as a therapeutic strategy to enhance immunoregulatory functions of dendritic cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 12468-12473.	3.3	68
154	Insights into the genetic epidemiology of Crohn's and rare diseases in the Ashkenazi Jewish population. <i>PLoS Genetics</i> , 2018, 14, e1007329.	1.5	66
155	Capsular polysaccharide correlates with immune response to the human gut microbe <i>Ruminococcus gnavus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	66
156	Mechanisms and function of autophagy in intestinal disease. <i>Autophagy</i> , 2018, 14, 216-220.	4.3	64
157	Antigen discovery and specification of immunodominance hierarchies for MHCII-restricted epitopes. <i>Nature Medicine</i> , 2018, 24, 1762-1772.	15.2	64
158	Understanding human immune function using the resources from the Human Functional Genomics Project. <i>Nature Medicine</i> , 2016, 22, 831-833.	15.2	63
159	Functional profiling of the gut microbiome in disease-associated inflammation. <i>Genome Medicine</i> , 2013, 5, 65.	3.6	61
160	Sorting nexin 5 mediates virus-induced autophagy and immunity. <i>Nature</i> , 2021, 589, 456-461.	18.7	61
161	The T300A Crohn's disease risk polymorphism impairs function of the WD40 domain of ATG16L1. <i>Nature Communications</i> , 2016, 7, 11821.	5.8	59
162	The Role of the Histone Methyltransferase Enhancer of Zeste Homolog 2 (EZH2) in the Pathobiological Mechanisms Underlying Inflammatory Bowel Disease (IBD). <i>Journal of Biological Chemistry</i> , 2017, 292, 706-722.	1.6	59

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