Ferdinando Bonfiglio

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8372427/publications.pdf

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45 papers

2,259 citations

279701 23 h-index 254106 43 g-index

50 all docs

50 docs citations

50 times ranked

4009 citing authors

#	Article	IF	CITATIONS
1	Dynamics of the human gut microbiome in inflammatory bowel disease. Nature Microbiology, 2017, 2, 17004.	5.9	830
2	Functional variants in the sucrase–isomaltase gene associate with increased risk of irritable bowel syndrome. Gut, 2018, 67, 263-270.	6.1	120
3	Genome-wide analysis of 53,400 people with irritable bowel syndrome highlights shared genetic pathways with mood and anxiety disorders. Nature Genetics, 2021, 53, 1543-1552.	9.4	96
4	Mitochondrial dysfunction in down syndrome: molecular mechanisms and therapeutic targets. Molecular Medicine, 2018, 24, 2.	1.9	85
5	Chronic pro-oxidative state and mitochondrial dysfunctions are more pronounced in fibroblasts from Down syndrome foeti with congenital heart defects. Human Molecular Genetics, 2013, 22, 1218-1232.	1.4	77
6	Metformin restores the mitochondrial network and reverses mitochondrial dysfunction in Down syndrome cells. Human Molecular Genetics, 2017, 26, ddx016.	1.4	70
7	Increased Prevalence of Rare Sucrase-isomaltase PathogenicÂVariants in Irritable Bowel Syndrome Patients. Clinical Gastroenterology and Hepatology, 2018, 16, 1673-1676.	2.4	64
8	Common variants at 21q22.3 locus influence MX1 and TMPRSS2 gene expression and susceptibility to severe COVID-19. IScience, 2021, 24, 102322.	1.9	60
9	Systemic Inflammation in Preclinical Ulcerative Colitis. Gastroenterology, 2021, 161, 1526-1539.e9.	0.6	58
10	Female-Specific Association Between Variants on Chromosome 9 and Self-Reported Diagnosis of Irritable Bowel Syndrome. Gastroenterology, 2018, 155, 168-179.	0.6	55
11	NRIP1/RIP140 siRNA-mediated attenuation counteracts mitochondrial dysfunction in Down syndrome. Human Molecular Genetics, 2014, 23, 4406-4419.	1.4	53
12	Dense genotyping of immune-related loci identifies HLA variants associated with increased risk of collagenous colitis. Gut, 2017, 66, 421-428.	6.1	50
13	Genetic and phenotypic attributes of splenic marginal zone lymphoma. Blood, 2022, 139, 732-747.	0.6	49
14	Stool frequency is associated with gut microbiota composition. Gut, 2017, 66, 559-560.	6.1	45
15	<i>TRPM8</i> polymorphisms associated with increased risk of IBS-C and IBS-M. Gut, 2017, 66, 1725-1727.	6.1	36
16	Predictors of recurrence following laparoscopic radical hysterectomy for early-stage cervical cancer: A multi-institutional study. Gynecologic Oncology, 2020, 159, 164-170.	0.6	35
17	A <scp>GWAS</scp> metaâ€analysis from 5 populationâ€based cohorts implicates ion channel genes in the pathogenesis of irritable bowel syndrome. Neurogastroenterology and Motility, 2018, 30, e13358.	1.6	34
18	Evaluation of Rapid Library Preparation Protocols for Whole Genome Sequencing Based Outbreak Investigation. Frontiers in Public Health, 2019, 7, 241.	1.3	32

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19	Collagenous Colitis Is Associated With HLA Signature and Shares Genetic Risks With Other Immune-Mediated Diseases. Gastroenterology, 2020, 159, 549-561.e8.	0.6	31
20	Survival implication of lymphadenectomy in patients surgically treated for apparent early-stage uterine serous carcinoma. Journal of Gynecologic Oncology, 2020, 31, e64.	1.0	30
21	Faecal microbiota composition associates with abdominal pain in the general population. Gut, 2018, 67, gutjnl-2017-314792.	6.1	29
22	Interferon-λ Enhances the Differentiation of Naive B Cells into Plasmablasts via the mTORC1 Pathway. Cell Reports, 2020, 33, 108211.	2.9	29
23	Overexpression of Chromosome 21 miRNAs May Affect Mitochondrial Function in the Hearts of Down Syndrome Fetuses. International Journal of Genomics, 2017, 2017, 1-10.	0.8	27
24	LACC1 polymorphisms in inflammatory bowel disease and juvenile idiopathic arthritis. Genes and Immunity, 2016, 17, 261-264.	2.2	25
25	HLA Associations Distinguish Collagenous From Lymphocytic Colitis. American Journal of Gastroenterology, 2016, 111, 1211-1213.	0.2	24
26	Functional Analyses of the Crohn's Disease Risk Gene LACC1. PLoS ONE, 2016, 11, e0168276.	1.1	24
27	A rationally designed oral vaccine induces immunoglobulin A in the murine gut that directs the evolution of attenuated Salmonella variants. Nature Microbiology, 2021, 6, 830-841.	5.9	21
28	A metaâ€analysis of reflux genomeâ€wide association studies in 6750 Northern Europeans from the general population. Neurogastroenterology and Motility, 2017, 29, e12923.	1.6	20
29	Crohn's Disease Is Associated With Activation of Circulating Innate Lymphoid Cells. Inflammatory Bowel Diseases, 2021, 27, 1128-1138.	0.9	18
30	The prevalence and transcriptional activity of the mucosal microbiota of ulcerative colitis patients. Scientific Reports, 2018, 8, 17278.	1.6	17
31	Pioglitazone Improves Mitochondrial Organization and Bioenergetics in Down Syndrome Cells. Frontiers in Genetics, 2019, 10, 606.	1.1	17
32	Targeted Analysis of Serum Proteins Encoded at Known Inflammatory Bowel Disease Risk Loci. Inflammatory Bowel Diseases, 2019, 25, 306-316.	0.9	15
33	GWAS of stool frequency provides insights into gastrointestinal motility and irritable bowel syndrome. Cell Genomics, 2021, 1, 100069.	3.0	15
34	Human Trisomic iPSCs from Down Syndrome Fibroblasts Manifest Mitochondrial Alterations Early during Neuronal Differentiation. Biology, 2021, 10, 609.	1.3	11
35	Rare Hypomorphic Sucrase Isomaltase Variants in Relation to Irritable Bowel Syndrome Risk in UK Biobank. Gastroenterology, 2021, 161, 1712-1714.	0.6	11
36	Circulating tumor DNA for comprehensive noninvasive monitoring of lymphoma treated with ibrutinib plus nivolumab. Blood Advances, 2021, 5, 4674-4685.	2.5	10

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37	Short stature and primary ovarian insufficiency possibly due to chromosomal position effect in a balanced X;1 translocation. Molecular Cytogenetics, 2015, 8, 50.	0.4	8
38	Germline rare variants of lectin pathway genes predispose to asymptomatic SARS-CoV-2 infection in elderly individuals. Genetics in Medicine, 2022, , .	1.1	7
39	Evaluation of the Different Stratification Models for POD24 Prediction in Patients with Follicular Lymphoma. Blood, 2020, 136, 24-25.	0.6	5
40	Overexpression of the Hsa21 Transcription Factor RUNX1 Modulates the Extracellular Matrix in Trisomy 21 Cells. Frontiers in Genetics, 2022, 13, 824922.	1.1	4
41	OC-047â€Epigenetic alterations at diagnosis predict susceptibility, prognosis and treatment escalation in inflammatory bowel disease – ibd character. , 2017, , .		2
42	OUP accepted manuscript. Human Molecular Genetics, 2021, , .	1.4	2
43	Epigenetic Alterations at Diagnosis Predict Susceptibility, Prognosis and Treatment Escalation in Inflammatory Bowel Disease and IBD Character. Gastroenterology, 2017, 152, S565.	0.6	1
44	P061 Epigenetic alterations at diagnosis predict susceptibility, prognosis and treatment escalation in inflammatory bowel disease $\hat{a} \in BD$ Character. Journal of Crohn's and Colitis, 2017, 11, S108-S108.	0.6	0
45	Common Variants at 21q22.3 Locus Influence $\langle i \rangle$ MX1 $\langle i \rangle$ Gene Expression and Susceptibility to Severe COVID-19. SSRN Electronic Journal, 0, , .	0.4	O