

Anna Latiano

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/8961616/anna-latiano-publications-by-year.pdf>

Version: 2024-04-10

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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

100 papers	12,043 citations	37 h-index	109 g-index
114 ext. papers	14,186 ext. citations	8.7 avg, IF	4.41 L-index

#	Paper	IF	Citations
100	Genome-wide analysis of 53,400 people with irritable bowel syndrome highlights shared genetic pathways with mood and anxiety disorders. <i>Nature Genetics</i> , 2021 , 53, 1543-1552	36.3	11
99	False-positive results of SARS-CoV-2 IgM/IgG antibody tests in sera stored before the 2020 pandemic in Italy. <i>International Journal of Infectious Diseases</i> , 2021 , 104, 159-163	10.5	11
98	Impact of the COVID-19 outbreak and the serum prevalence of SARS-CoV-2 antibodies in patients with inflammatory bowel disease treated with biologic drugs. <i>Digestive and Liver Disease</i> , 2021 , 53, 277-282	23.2	12
97	Mapping the human genetic architecture of COVID-19. <i>Nature</i> , 2021 ,	50.4	162
96	Germline Alterations in Patients With IBD-associated Colorectal Cancer. <i>Inflammatory Bowel Diseases</i> , 2021 ,	4.5	1
95	Association of Genetic Variants Affecting microRNAs and Pancreatic Cancer Risk. <i>Frontiers in Genetics</i> , 2021 , 12, 693933	4.5	2
94	Circulating levels of cytokines, chemokines and growth factors in patients with achalasia. <i>Biomedical Reports</i> , 2021 , 15, 92	1.8	
93	Worse impact of second wave COVID-19 pandemic in adults but not in children with inflammatory bowel disease: an Italian single tertiary center experience. <i>European Review for Medical and Pharmacological Sciences</i> , 2021 , 25, 2744-2747	2.9	2
92	Genomewide Association Study of Severe Covid-19 with Respiratory Failure. <i>New England Journal of Medicine</i> , 2020 , 383, 1522-1534	59.2	913
91	microRNA-mRNA network model in patients with achalasia. <i>Neurogastroenterology and Motility</i> , 2020 , 32, e13764	4	2
90	Glycosylation of Immunoglobulin G Associates With Clinical Features of Inflammatory Bowel Diseases. <i>Gastroenterology</i> , 2018 , 154, 1320-1333.e10	13.3	82
89	Female-Specific Association Between Variants on Chromosome 9 and Self-Reported Diagnosis of Irritable Bowel Syndrome. <i>Gastroenterology</i> , 2018 , 155, 168-179	13.3	31
88	Do pancreatic cancer and chronic pancreatitis share the same genetic risk factors? A PANcreatic Disease ReseArch (PANDoRA) consortium investigation. <i>International Journal of Cancer</i> , 2018 , 142, 290-296	7.5	12
87	Plasma N-Glycan Signatures Are Associated With Features of Inflammatory Bowel Diseases. <i>Gastroenterology</i> , 2018 , 155, 829-843	13.3	47
86	Promoter methylation of the and genes correlates with the composition of the immunoglobulin G glycome in inflammatory bowel disease. <i>Clinical Epigenetics</i> , 2018 , 10, 75	7.7	17
85	IBD risk loci are enriched in multigenic regulatory modules encompassing putative causative genes. <i>Nature Communications</i> , 2018 , 9, 2427	17.4	95
84	Crohn's Colitis: Development of a multiplex gene expression assay comparing mRNA levels of susceptibility genes. <i>Clinics and Research in Hepatology and Gastroenterology</i> , 2017 , 41, 435-444	2.4	2

83	Addendum: Palmieri, O. et al. Functional Implications of MicroRNAs in Crohn's Disease Revealed by Integrating MicroRNA and Messenger RNA Expression Profiling. <i>Int. J. Mol. Sci.</i> 2017 , <i>18</i> , 1580. <i>International Journal of Molecular Sciences</i> , 2017 , <i>18</i> , 2113	6.3	78
82	Functional Implications of MicroRNAs in Crohn's Disease Revealed by Integrating MicroRNA and Messenger RNA Expression Profiling. <i>International Journal of Molecular Sciences</i> , 2017 , <i>18</i> ,	6.3	14
81	Crohn's Disease Localization Displays Different Predisposing Genetic Variants. <i>PLoS ONE</i> , 2017 , <i>12</i> , e0168821	3.7	10
80	The HLA-DQ1 insertion is a strong achalasia risk factor and displays a geospatial north-south gradient among Europeans. <i>European Journal of Human Genetics</i> , 2016 , <i>24</i> , 1228-31	5.3	16
79	Metabolomic profile in pancreatic cancer patients: a consensus-based approach to identify highly discriminating metabolites. <i>Oncotarget</i> , 2016 , <i>7</i> , 5815-29	3.3	56
78	Gene expression of muscular and neuronal pathways is cooperatively dysregulated in patients with idiopathic achalasia. <i>Scientific Reports</i> , 2016 , <i>6</i> , 31549	4.9	19
77	Inflammatory Bowel Disease Meets Systems Biology: A Multi-Omics Challenge and Frontier. <i>OMICS A Journal of Integrative Biology</i> , 2016 , <i>20</i> , 692-698	3.8	13
76	Systematic analysis of circadian genes using genome-wide cDNA microarrays in the inflammatory bowel disease transcriptome. <i>Chronobiology International</i> , 2015 , <i>32</i> , 903-16	3.6	31
75	Genetic sharing and heritability of paediatric age of onset autoimmune diseases. <i>Nature Communications</i> , 2015 , <i>6</i> , 8442	17.4	46
74	Meta-analysis of shared genetic architecture across ten pediatric autoimmune diseases. <i>Nature Medicine</i> , 2015 , <i>21</i> , 1018-27	50.5	143
73	Genome-wide Pathway Analysis Using Gene Expression Data of Colonic Mucosa in Patients with Inflammatory Bowel Disease. <i>Inflammatory Bowel Diseases</i> , 2015 , <i>21</i> , 1260-8	4.5	14
72	Impact of genetic polymorphisms on the pathogenesis of idiopathic achalasia: Association with IL33 gene variant. <i>Human Immunology</i> , 2014 , <i>75</i> , 364-9	2.3	8
71	Common variants in the HLA-DQ region confer susceptibility to idiopathic achalasia. <i>Nature Genetics</i> , 2014 , <i>46</i> , 901-4	36.3	75
70	Variation in genes encoding for interferon β and ϵ in the prediction of HCV-1 treatment-induced viral clearance. <i>Liver International</i> , 2014 , <i>34</i> , 1369-77	7.9	7
69	Genetic variation in the lymphotoxin- α (LTA)/tumour necrosis factor- α (TNF) α locus as a risk factor for idiopathic achalasia. <i>Gut</i> , 2014 , <i>63</i> , 1401-9	19.2	16
68	Association between variants of PRDM1 and NDP52 and Crohn's disease, based on exome sequencing and functional studies. <i>Gastroenterology</i> , 2013 , <i>145</i> , 339-47	13.3	125
67	Genetic variants of membrane metalloproteinase genes in inflammatory bowel diseases. <i>Digestive and Liver Disease</i> , 2013 , <i>45</i> , 1003-10	3.3	2
66	Associations between genetic polymorphisms in IL-33, IL1R1 and risk for inflammatory bowel disease. <i>PLoS ONE</i> , 2013 , <i>8</i> , e62144	3.7	68

65	Erythrocytes-mediated delivery of dexamethasone 21-phosphate in steroid-dependent ulcerative colitis: a randomized, double-blind Sham-controlled study. <i>Inflammatory Bowel Diseases</i> , 2013 , 19, 1872-9	4.5	18
64	Association study of a polymorphism in clock gene PERIOD3 and risk of inflammatory bowel disease. <i>Chronobiology International</i> , 2012 , 29, 994-1003	3.6	30
63	The expression of leucine-rich repeat gene family members in colorectal cancer. <i>Experimental Biology and Medicine</i> , 2012 , 237, 1123-8	3.7	16
62	Host-microbe interactions have shaped the genetic architecture of inflammatory bowel disease. <i>Nature</i> , 2012 , 491, 119-24	50.4	3239
61	Neuroimmune interactions in patients with inflammatory bowel diseases: disease activity and clinical behavior based on Substance P serum levels. <i>Journal of Crohn's and Colitis</i> , 2012 , 6, 563-70	1.5	19
60	Glucocorticoid resistance in Crohn's disease and ulcerative colitis: an association study investigating GR and FKBP5 gene polymorphisms. <i>Pharmacogenomics Journal</i> , 2012 , 12, 432-8	3.5	29
59	Discovering genetic variants in Crohn's disease by exploring genomic regions enriched of weak association signals. <i>Digestive and Liver Disease</i> , 2011 , 43, 623-31	3.3	4
58	Deep resequencing of GWAS loci identifies independent rare variants associated with inflammatory bowel disease. <i>Nature Genetics</i> , 2011 , 43, 1066-73	36.3	584
57	Genetics and ulcerative colitis: what are the clinical implications?. <i>Current Drug Targets</i> , 2011 , 12, 1383-93		4
56	Meta-analysis identifies 29 additional ulcerative colitis risk loci, increasing the number of confirmed associations to 47. <i>Nature Genetics</i> , 2011 , 43, 246-52	36.3	1028
55	RS-SNP: a random-set method for genome-wide association studies. <i>BMC Genomics</i> , 2011 , 12, 166	4.5	1
54	IL23R, ATG16L1, IRGM, OCTN1, and OCTN2 mRNA expression in inflamed and noninflamed mucosa of IBD patients. <i>Inflammatory Bowel Diseases</i> , 2011 , 17, 1832-3	4.5	5
53	Dissecting the mucosal expression of human leucine-rich repeat family genes in inflammatory bowel disease patients. <i>Inflammatory Bowel Diseases</i> , 2011 , 17, 1834-5	4.5	1
52	Genome-wide expression profiling identifies an impairment of negative feedback signals in the Crohn's disease-associated NOD2 variant L1007fsinsC. <i>Journal of Immunology</i> , 2011 , 186, 4027-38	5.3	18
51	Investigation of multiple susceptibility loci for inflammatory bowel disease in an Italian cohort of patients. <i>PLoS ONE</i> , 2011 , 6, e22688	3.7	45
50	Genome-wide association identifies multiple ulcerative colitis susceptibility loci. <i>Nature Genetics</i> , 2010 , 42, 332-7	36.3	491
49	Genome-wide meta-analysis increases to 71 the number of confirmed Crohn's disease susceptibility loci. <i>Nature Genetics</i> , 2010 , 42, 1118-25	36.3	1946
48	The -A2518G polymorphism of monocyte chemoattractant protein-1 is associated with Crohn's disease. <i>American Journal of Gastroenterology</i> , 2010 , 105, 1586-94	0.7	21

47	IL-1beta-511 and IL-1RN*2 polymorphisms in inflammatory bowel disease: An Italian population study and meta-analysis of European studies. <i>Digestive and Liver Disease</i> , 2010 , 42, 179-84	3.3	13
46	Variants at the 3p21 locus influence susceptibility and phenotype both in adults and early-onset patients with inflammatory bowel disease. <i>Inflammatory Bowel Diseases</i> , 2010 , 16, 1108-17	4.5	14
45	Polymorphism of the IRGM gene might predispose to fistulizing behavior in Crohn's disease. <i>American Journal of Gastroenterology</i> , 2009 , 104, 110-6	0.7	67
44	Association of genetic profiles to Crohn's disease by linear combinations of single nucleotide polymorphisms. <i>Artificial Intelligence in Medicine</i> , 2009 , 46, 131-8	7.4	3
43	Ulcerative colitis-risk loci on chromosomes 1p36 and 12q15 found by genome-wide association study. <i>Nature Genetics</i> , 2009 , 41, 216-20	36.3	325
42	Common variants at five new loci associated with early-onset inflammatory bowel disease. <i>Nature Genetics</i> , 2009 , 41, 1335-40	36.3	389
41	High resolution melting (HRM) analysis for the detection of ER22/23EK, BclI, and N363S polymorphisms of the glucocorticoid receptor gene. <i>Journal of Steroid Biochemistry and Molecular Biology</i> , 2009 , 113, 269-74	5.1	23
40	Enteropathic spondyloarthropathy: a common genetic background with inflammatory bowel disease?. <i>World Journal of Gastroenterology</i> , 2009 , 15, 2456-62	5.6	17
39	MAST3: a novel IBD risk factor that modulates TLR4 signaling. <i>Genes and Immunity</i> , 2008 , 9, 602-12	4.4	27
38	Erythrocyte-mediated delivery of dexamethasone in patients with mild-to-moderate ulcerative colitis, refractory to mesalamine: a randomized, controlled study. <i>American Journal of Gastroenterology</i> , 2008 , 103, 2509-16	0.7	59
37	Gene-centric association mapping of chromosome 3p implicates MST1 in IBD pathogenesis. <i>Mucosal Immunology</i> , 2008 , 1, 131-8	9.2	67
36	The association of MYO9B gene in Italian patients with inflammatory bowel diseases. <i>Alimentary Pharmacology and Therapeutics</i> , 2008 , 27, 241-8	6.1	27
35	Gender-stratified analysis of DLG5 R30Q in 4707 patients with Crohn disease and 4973 controls from 12 Caucasian cohorts. <i>Journal of Medical Genetics</i> , 2008 , 45, 36-42	5.8	37
34	Replication of interleukin 23 receptor and autophagy-related 16-like 1 association in adult- and pediatric-onset inflammatory bowel disease in Italy. <i>World Journal of Gastroenterology</i> , 2008 , 14, 4643-51	5.6	57
33	DMBT1 confers mucosal protection in vivo and a deletion variant is associated with Crohn's disease. <i>Gastroenterology</i> , 2007 , 133, 1499-509	13.3	81
32	Evaluating the role of the genetic variations of PTPN22, NFKB1, and FcGR11A genes in inflammatory bowel disease: a meta-analysis. <i>Inflammatory Bowel Diseases</i> , 2007 , 13, 1212-9	4.5	32
31	Sequential evaluation of thiopurine methyltransferase, inosine triphosphate pyrophosphatase, and HPRT1 genes polymorphisms to explain thiopurines' toxicity and efficacy. <i>Alimentary Pharmacology and Therapeutics</i> , 2007 , 26, 737-45	6.1	34
30	Regularized least squares classifiers may predict Crohn's disease from profiles of single nucleotide polymorphisms. <i>Annals of Human Genetics</i> , 2007 , 71, 537-49	2.2	6

29	Analysis of candidate genes on chromosomes 5q and 19p in celiac disease. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2007 , 45, 180-6	2.8	14
28	Dissecting genetic predisposition to inflammatory bowel disease: current progress and prospective application. <i>Expert Review of Clinical Immunology</i> , 2007 , 3, 287-98	5.1	4
27	Polymorphisms of tumor necrosis factor-alpha but not MDR1 influence response to medical therapy in pediatric-onset inflammatory bowel disease. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2007 , 44, 171-9	2.8	67
26	Erythrocytes as a controlled drug delivery system: clinical evidences. <i>Journal of Controlled Release</i> , 2006 , 116, e43-5	11.7	11
25	Contribution of IBD5 locus to clinical features of IBD patients. <i>American Journal of Gastroenterology</i> , 2006 , 101, 318-25	0.7	24
24	Genetic variation in myosin IXB is associated with ulcerative colitis. <i>Gastroenterology</i> , 2006 , 131, 1768-74	13.3	82
23	Variants of OCTN1-2 cation transporter genes are associated with both Crohn's disease and ulcerative colitis. <i>Alimentary Pharmacology and Therapeutics</i> , 2006 , 23, 497-506	6.1	47
22	Increased intestinal permeability and NOD2 variants in familial and sporadic Crohn's disease. <i>Alimentary Pharmacology and Therapeutics</i> , 2006 , 23, 1455-61	6.1	78
21	HLA and enteric antineuronal antibodies in patients with achalasia. <i>Neurogastroenterology and Motility</i> , 2006 , 18, 520-5	4	29
20	Haplotype-based association analysis of 56 functional candidate genes in the IBD6 locus on chromosome 19. <i>European Journal of Human Genetics</i> , 2006 , 14, 780-90	5.3	21
19	Evidence of transmission ratio distortion of DLG5 R30Q variant in general and implication of an association with Crohn disease in men. <i>Human Genetics</i> , 2006 , 119, 305-11	6.3	55
18	Erythrocytes-mediated delivery of dexamethasone in steroid-dependent IBD patients-a pilot uncontrolled study. <i>American Journal of Gastroenterology</i> , 2005 , 100, 1370-5	0.7	66
17	Idiopathic achalasia is not allelic to alacrima achalasia adrenal insufficiency syndrome at the ALADIN locus. <i>Digestive and Liver Disease</i> , 2005 , 37, 312-5	3.3	18
16	Multidrug resistance 1 gene polymorphisms are not associated with inflammatory bowel disease and response to therapy in Italian patients. <i>Alimentary Pharmacology and Therapeutics</i> , 2005 , 22, 1129-38	6.1	57
15	Association of DLG5 R30Q variant with inflammatory bowel disease. <i>European Journal of Human Genetics</i> , 2005 , 13, 835-9	5.3	61
14	HLA-DRB1 alleles may influence disease phenotype in patients with inflammatory bowel disease: a critical reappraisal with review of the literature. <i>Diseases of the Colon and Rectum</i> , 2005 , 48, 57-64; discussion 64-5	3.1	22
13	Variants of CARD15 are associated with an aggressive clinical course of Crohn's disease--an IG-IBD study. <i>American Journal of Gastroenterology</i> , 2005 , 100, 84-92	0.7	98
12	Anti-Saccharomyces cerevisiae mannan antibodies in inflammatory bowel disease: comparison of different assays and correlation with clinical features. <i>Alimentary Pharmacology and Therapeutics</i> , 2004 , 20, 1143-52	6.1	14

11	The frame-shift mutation of the NOD2/CARD15 gene is significantly increased in ulcerative colitis: an *IG-IBD study. <i>Gastroenterology</i> , 2004 , 126, 625-7	13.3	24
10	Frequency of NOD2/CARD15 variants in both sporadic and familial cases of Crohn's disease across Italy. An Italian Group for Inflammatory Bowel Disease Study. <i>Digestive and Liver Disease</i> , 2004 , 36, 121-4	3.3	29
9	CARD15 genotyping in inflammatory bowel disease patients by multiplex pyrosequencing. <i>Clinical Chemistry</i> , 2003 , 49, 1675-9	5.5	29
8	Genetics of inflammatory bowel disease: the beginning of the end or the end of the beginning?. <i>Digestive and Liver Disease</i> , 2003 , 35, 442-9	3.3	17
7	Linkage of ulcerative colitis to the pericentromeric region of chromosome 16 in Italian inflammatory bowel disease families is independent of the presence of common CARD15 mutations. <i>Journal of Medical Genetics</i> , 2003 , 40, 837-41	5.8	5
6	Association of Crohn's disease and ulcerative colitis with haplotypes of the MLH1 gene in Italian inflammatory bowel disease patients. <i>Journal of Medical Genetics</i> , 2002 , 39, 332-4	5.8	7
5	Combined segregation and linkage analysis of inflammatory bowel disease in the IBD1 region using severity to characterise Crohn's disease and ulcerative colitis. On behalf of the GISC. <i>European Journal of Human Genetics</i> , 2000 , 8, 846-52	5.3	16
4	Antineutrophil cytoplasmic antibodies in inflammatory bowel disease: clinical role and review of the literature. <i>Diseases of the Colon and Rectum</i> , 2000 , 43, 999-1007	3.1	21
3	Genetic analysis in Italian families with inflammatory bowel disease supports linkage to the IBD1 locus--a GISC study. <i>European Journal of Human Genetics</i> , 1999 , 7, 567-73	5.3	73
2	Helicobacter pylori infection and growth delay in older children. <i>Archives of Disease in Childhood</i> , 1997 , 77, 46-9	2.2	98
1	The ABO blood group locus and a chromosome 3 gene cluster associate with SARS-CoV-2 respiratory failure in an Italian-Spanish genome-wide association analysis		23