CITATION REPORT List of articles citing

Meta-analysis identifies 29 additional ulcerative colitis risk loci, increasing the number of confirmed associations to 47

DOI: 10.1038/ng.764 Nature Genetics, 2011, 43, 246-52.

Source: https://exaly.com/paper-pdf/51025163/citation-report.pdf

Version: 2024-04-20

This report has been generated based on the citations recorded by exaly.com for the above article. 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.

#	Paper	IF	Citations
1131			
1130	Genetic risk prediction in complex disease. 2011 , 20, R182-8		130
1129	Recent insights into the genetics of inflammatory bowel disease. 2011 , 140, 1704-12		312
1128	Single nucleotide polymorphisms that increase expression of the guanosine triphosphatase RAC1 are associated with ulcerative colitis. 2011 , 141, 633-41		58
1127	TLE1 modifies the effects of NOD2 in the pathogenesis of Crohn's disease. 2011 , 141, 972-981.e1-2		26
1126	Genetic profiling in inflammatory bowel disease: from association to bedside. 2011 , 141, 1566-71.e1		14
1125	Genetic risks and childhood-onset asthma. 2011 , 128, 266-70; quiz 271-2		22
1124	Association of FCGR2A, JAK2 or HNF4A variants with ulcerative colitis in Koreans. 2011 , 43, 856-61		22
1123	Ulcerative colitis. 2011 , 365, 1713-25		785
1122	Genetic polymorphisms of IL-23R and IL-17A and novel insights into their associations with inflammatory bowel disease. 2011 , 60, 1527-36		98
1121	Candidate mucosal and surrogate biomarkers of inflammatory bowel disease in the era of new technology. 2011 , 46, 1407-17		15
1120	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
1119	Genomics and the multifactorial nature of human autoimmune disease. 2011 , 365, 1612-23		243
1118	Metagenomics and personalized medicine. 2011 , 147, 44-56		164
1117	TH1 and TH17 interactions in untreated inflamed mucosa of inflammatory bowel disease, and their potential to mediate the inflammation. 2011 , 56, 633-40		96
1116	[New knowledge in genetics and inflammatory bowel disease. Are there any practical applications?]. 2011 , 34, 591-8		2
1115	T helper 17 cell heterogeneity and pathogenicity in autoimmune disease. 2011 , 32, 395-401		162

1114	Replication of genetic variation in the MYO9B gene in Crohn's disease. 2011 , 72, 592-7	15
1113	Genetic variations in matrix metalloproteinases may be associated with increased risk of ulcerative colitis. 2011 , 72, 1117-27	16
1112	Genetics and pathogenesis of inflammatory bowel disease. 2011 , 474, 307-17	1613
1111	Comparative Genetic Analysis of Type 1 Diabetes and Inflammatory Bowel Disease. 2011 ,	
1110	Ulcerative Colitis in Children and Adolescents. 2011,	
1109	Farnesoid X receptor (FXR) activation and FXR genetic variation in inflammatory bowel disease. 2011 , 6, e23745	89
1108	Protein characterization of a candidate mechanism SNP for Crohn's disease: the macrophage stimulating protein R689C substitution. 2011 , 6, e27269	22
1107	Host genetic susceptibility, dysbiosis, and viral triggers in inflammatory bowel disease. 2011 , 27, 321-7	56
1106	The prognostic impact of germline 46/1 haplotype of Janus kinase 2 in cytogenetically normal acute myeloid leukemia. 2011 , 96, 1613-8	11
1105	Bacteria in the pathogenesis of inflammatory bowel disease. 2011 , 39, 1067-72	35
1104	Celiac disease: moving from genetic associations to causal variants. 2011 , 80, 203-313	13
1103	Genome-wide analysis of extended pedigrees confirms IL2-IL21 linkage and shows additional regions of interest potentially influencing coeliac disease risk. 2011 , 78, 428-37	11
1102	New insight in the pathogenesis of functional gastrointestinal disorders: association between genetics and colonic transit. 2011 , 23, 893-7	6
1101	The kinase LRRK2 is a regulator of the transcription factor NFAT that modulates the severity of inflammatory bowel disease. 2011 , 12, 1063-70	268
1100	Blood pressure loci identified with a gene-centric array. 2011 , 89, 688-700	137
1099	Dissecting the genetics of complex inheritance: linkage disequilibrium mapping provides insight into Crohn disease. 2011 , 89, 798-805	27
1098	Expression and functional importance of innate immune receptors by intestinal epithelial cells. 2011 , 68, 3661-73	40
1097	Assessment of heterogeneity between European Populations: a Baltic and Danish replication case-control study of SNPs from a recent European ulcerative colitis genome wide association study. 2011 , 12, 139	6

Evaluation of 22 genetic variants with Crohn's disease risk in the Ashkenazi Jewish popula case-control study. 2011 , 12, 63	ation: a
1095 Th17-related cytokines: new players in the control of chronic intestinal inflammation. 201	1 , 9, 122 61
Reviving pegylated interferon as a therapeutic agent for hepatitis D: no more room for nucleos(t)ides?. 2011 , 53, 2131-3	3
Genome-wide association studies in primary sclerosing cholangitis: still more questions the answers?. 2011 , 53, 2133-5	nan 3
1092 Deciphering the genetic predisposition to primary sclerosing cholangitis. 2011 , 31, 188-20	21
Comment on: Familial Mediterranean fever caused by homozygous E148Q mutation comp Budd-Chiari syndrome and polyarteritis nodosa: reply. 2011 , 50, 1349-1350	olicated by
1090 Exploring the hidden heritability of inflammatory bowel disease. 2011 , 60, 1447-8	5
Commonalities and differences between Crohn's disease and ulcerative colitis: the genet their interpretation. 2011 , 10, 447-54	ic clues to
Severity of innate immune-mediated colitis is controlled by the cytokine deficiency-inductions susceptibility-1 (Cdcs1) locus. 2011 , 108, 7137-41	ed colitis 26
Correction: Corrigendum: Meta-analysis identifies 29 additional ulcerative colitis risk loci, the number of confirmed associations to 47. <i>Nature Genetics</i> , 2011 , 43, 919-919	increasing 36.3 8
Comment on: Familial Mediterranean fever caused by homozygous E148Q mutation comp Budd-Chiari syndrome and polyarteritis nodosa. 2011 , 50, 1348-9; author reply 1349	olicated by
Canine breeds at high risk of developing inflammatory bowel disease in the south-eastern , 169, 635 $$	n UK. 2011 40
1084 Viruses, autophagy genes, and Crohn's disease. 2011 , 3, 1281-311	30
$_{1083}$ Comparison of family history and SNPs for predicting risk of complex disease. 2012 , 8, e1	002973 78
1082 Knockout of Mkp-1 exacerbates colitis in Il-10-deficient mice. 2012 , 302, G1322-35	19
Interferon regulatory factor 5 gene polymorphism confers risk to several rheumatic disea correlates with expression of alternative thymic transcripts. 2012 , 51, 619-26	ises and 29
IL-26 is overexpressed in rheumatoid arthritis and induces proinflammatory cytokine produced and Th17 cell generation. 2012 , 10, e1001395	duction 98
	ind

(2012-2012)

1078	Replication study of 10 genes showing evidence for association with multiple sclerosis: validation of TMEM39A, IL12B and CBLB [correction of CLBL] genes. 2012 , 18, 959-65	24
1077	Unraveling the regulatory mechanisms underlying tissue-dependent genetic variation of gene expression. 2012 , 8, e1002431	163
1076	Brain expression genome-wide association study (eGWAS) identifies human disease-associated variants. 2012 , 8, e1002707	174
1075	Macrophage-stimulating protein polymorphism rs3197999 is associated with a gain of function: implications for inflammatory bowel disease. 2012 , 13, 321-7	16
1074	Immunochip analyses identify a novel risk locus for primary biliary cirrhosis at 13q14, multiple independent associations at four established risk loci and epistasis between 1p31 and 7q32 risk variants. 2012 , 21, 5209-21	122
1073	A genome-wide association scan on the levels of markers of inflammation in Sardinians reveals associations that underpin its complex regulation. 2012 , 8, e1002480	112
1072	PPARlin Inflammatory Bowel Disease. 2012 , 2012, 620839	47
1071	Epigenetics in inflammatory bowel disease. 2012 , 28, 577-84	34
1070	Inflammatory bowel disease pathogenesis: what is new?. 2012 , 28, 301-9	64
1069	The impact of genomics on pediatric research and medicine. 2012 , 129, 1150-60	9
1068	Identification of a genetic locus controlling bacteria-driven colitis and associated cancer through effects on innate inflammation. 2012 , 209, 1309-24	37
1067	The role of LRRK2 in inflammatory bowel disease. 2012 , 22, 1092-4	32
1066	A functional methylome map of ulcerative colitis. 2012 , 22, 2130-7	96
1065	Bias in high-tier medical journals concerning physician-academic relationships with industry. 2012 , 30, 320-2	11
1064	Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. <i>Nature Genetics</i> , 2012 , 44, 991-1005	621
1063	Association of celiac disease genes with inflammatory bowel disease in Finnish and Swedish patients. 2012 , 13, 474-80	21
1062	Autophagy in immunity: implications in etiology of autoimmune/autoinflammatory diseases. 2012 , 8, 1286-99	100
1061	Protective and aggravating effects of Nlrp3 inflammasome activation in IBD models: influence of genetic and environmental factors. 2012 , 30 Suppl 1, 82-90	100

1060	Drosophila melanogaster Selection for Survival of Bacillus cereus Infection: Life History Trait Indirect Responses. 2012 , 2012, 935970	15
1059	The chbG gene of the chitobiose (chb) operon of Escherichia coli encodes a chitooligosaccharide deacetylase. 2012 , 194, 4959-71	22
1058	A lecture on the genetics of primary sclerosing cholangitis. 2012 , 30 Suppl 1, 32-8	12
1057	Markers that differentiate early from late IBD. 2012 , 30, 380-2	1
1056	Why and where to look in the environment with regard to the etiology of inflammatory bowel disease. 2012 , 30 Suppl 3, 28-32	10
1055	Nutrigenomics and nutrigenetics in inflammatory bowel diseases. 2012 , 46, 735-47	21
1054	Protein tyrosine phosphatases and type 1 diabetes: genetic and functional implications of PTPN2 and PTPN22. 2012 , 9, 188-200	14
1053	Current advantages in the application of proteomics in inflammatory bowel disease. 2012 , 57, 2755-64	14
1052	Genetics of spondyloarthritisbeyond the MHC. 2012 , 8, 296-304	142
1051	Endoplasmic reticulum stress and inflammation. 2012 , 30, 341-6	41
1050	IL-7 in human health and disease. 2012 , 24, 218-24	112
1049	Geographical variation and incidence of inflammatory bowel disease among US women. 2012 , 61, 1686-92	150
1048	Association study of a polymorphism in clock gene PERIOD3 and risk of inflammatory bowel disease. 2012 , 29, 994-1003	30
1047	Abnormal genetic and epigenetic changes in signal transducer and activator of transcription 4 in the pathogenesis of inflammatory bowel diseases. 2012 , 57, 2600-7	20
1046	Crohn's disease. 2012 , 380, 1590-605	1304
1045	Ulcerative colitis. 2012 , 380, 1606-19	1036
1044	Nematode modulation of inflammatory bowel disease. 2012 , 249, 871-86	23
1043	Foxp3 exploits a pre-existent enhancer landscape for regulatory T cell lineage specification. 2012 , 151, 153-66	342

(2012-2012)

1042	receptor+ innate lymphoid cells. 2012 , 37, 674-84	244
1041	Genetics in diagnosing and managing inflammatory bowel disease. 2012 , 41, 513-22	9
1040	Cytokine gene polymorphisms and human autoimmune disease in the era of genome-wide association studies. 2012 , 32, 139-51	46
1039	Epigenetics: filling in the 'heritability gap' and identifying gene-environment interactions in ulcerative colitis. 2012 , 4, 72	10
1038	Regulation of intestinal homeostasis by innate and adaptive immunity. 2012 , 24, 673-80	70
1037	Modifying T-cell trafficking to the intestinal as a potential management for inflammatory bowel disease. 2012 , 21, 975-84	5
1036	Surgical systems biology and personalized longitudinal phenotyping in critical care. 2012 , 9, 593-608	5
1035	Pregnane X receptor as a target for treatment of inflammatory bowel disorders. 2012 , 33, 323-30	111
1034	Inflammatory bowel disease and pregnancy: overlapping pathways. 2012 , 160, 65-83	19
1033	The IL-23/IL-17 pathway in inflammatory bowel disease. 2012 , 6, 223-37	59
1032	Twin studies in autoimmune disease: genetics, gender and environment. 2012 , 38, J156-69	189
1031	The impact of the gut microbiota on human health: an integrative view. 2012 , 148, 1258-70	2117
1030	Lactocepin secreted by Lactobacillus exerts anti-inflammatory effects by selectively degrading proinflammatory chemokines. 2012 , 11, 387-96	149
1029	Meta-analysis of genetic polymorphisms in granulomatosis with polyangiitis (Wegener's) reveals shared susceptibility loci with rheumatoid arthritis. 2012 , 64, 3463-71	27
1028	Network-based SNP meta-analysis identifies joint and disjoint genetic features across common human diseases. 2012 , 13, 490	1
1027	Comparative analysis of inflamed and non-inflamed colon biopsies reveals strong proteomic inflammation profile in patients with ulcerative colitis. 2012 , 12, 76	42
1026	Systems analysis of inflammatory bowel disease based on comprehensive gene information. 2012 , 13, 25	5
1025	Interleukin-7 receptor blockade suppresses adaptive and innate inflammatory responses in experimental colitis. 2012 , 9, 39	31

Month of birth, vitamin D and risk of immune-mediated disease: a case control study. 2012 , 10, 69	100
Clinical review: Genome-wide association studies of skeletal phenotypes: what we have learned and where we are headed. 2012 , 97, E1958-77	80
1022 Targeting T-cell migration in inflammatory bowel disease. 2012 , 272, 411-29	45
Familial autoimmunity in systemic sclerosis results of a French-based case-control family study. 2012 , 39, 532-8	20
1020 The human IL-7 receptor gene: deletions, polymorphisms and mutations. 2012 , 24, 225-30	50
Host-microbe interactions have shaped the genetic architecture of inflammatory bowel disease. 2012, 491, 119-24	3239
1018 ANKRD55 and DHCR7 are novel multiple sclerosis risk loci. 2012 , 13, 253-7	37
Efficacy of hepatitis B vaccination and revaccination and factors impacting on response in patients with inflammatory bowel disease. 2012 , 107, 1460-6	98
1016 Towards a novel molecular classification of IBD. 2012 , 30, 425-7	13
1015 Higher predicted vitamin D status is associated with reduced risk of Crohn's disease. 2012 , 142, 482-9	297
1014 Within you, without you: is gastroenterology ready to embrace the "exposome"?. 2012 , 142, 1403-4	
	8
1013 The age of gene discovery in very early onset inflammatory bowel disease. 2012 , 143, 285-8	69
The age of gene discovery in very early onset inflammatory bowel disease. 2012 , 143, 285-8 Hormone therapy increases risk of ulcerative colitis but not Crohn's disease. 2012 , 143, 1199-1206	
	69
Hormone therapy increases risk of ulcerative colitis but not Crohn's disease. 2012 , 143, 1199-1206 Genome-wide association studies of asthma indicate opposite immunopathogenesis direction from	69 77
Hormone therapy increases risk of ulcerative colitis but not Crohn's disease. 2012, 143, 1199-1206 Genome-wide association studies of asthma indicate opposite immunopathogenesis direction from autoimmune diseases. 2012, 130, 861-8.e7 Identification of IL18RAP/IL18R1 and IL12B as leprosy risk genes demonstrates shared	69 77 109
Hormone therapy increases risk of ulcerative colitis but not Crohn's disease. 2012, 143, 1199-1206 Genome-wide association studies of asthma indicate opposite immunopathogenesis direction from autoimmune diseases. 2012, 130, 861-8.e7 Identification of IL18RAP/IL18R1 and IL12B as leprosy risk genes demonstrates shared pathogenesis between inflammation and infectious diseases. 2012, 91, 935-41	69 77 109 58

1006	Autoimmunity: insights from human genomics. 2012 , 24, 513-5	1
1005	Results of the 2nd scientific workshop of the ECCO (III): basic mechanisms of intestinal healing. 2012 , 6, 373-85	38
1004	Second European evidence-based consensus on the diagnosis and management of ulcerative colitis part 1: definitions and diagnosis. 2012 , 6, 965-90	592
1003	LRRK2 as a negative regulator of NFAT: implications for the pathogenesis of inflammatory bowel disease. 2012 , 8, 227-9	7
1002	Rectocolite hfhorragique : pidfhiologie, physiopathologie, diagnostic, histoire naturelle et stratgie thfapeutique. 2012 , 7, 1-19	1
1001	Animal models of inflammatory bowel disease. 2012 , 105, 263-320	162
1000	Revisiting the old link between infection and autoimmune disease with commensals and T helper 17 cells. 2012 , 54, 50-68	20
999	Evidence from genetics for a role of autophagy and innate immunity in IBD pathogenesis. 2012 , 30, 330-3	39
998	A genome-wide association study identifies two new susceptibility loci for lung adenocarcinoma in the Japanese population. <i>Nature Genetics</i> , 2012 , 44, 900-3	137
997	The microbiome and inflammatory bowel disease: is there a therapeutic role for fecal microbiota transplantation?. 2012 , 107, 1452-9	146
996	Markers of intestinal inflammation in patients with ankylosing spondylitis: a pilot study. 2012 , 14, R261	34
995	Inflammatory bowel diseases phenotype, C. difficile and NOD2 genotype are associated with shifts in human ileum associated microbial composition. 2012 , 7, e26284	178
994	Multiple sclerosis risk variant HLA-DRB1*1501 associates with high expression of DRB1 gene in different human populations. 2012 , 7, e29819	67
993	Analysis of IL12B gene variants in inflammatory bowel disease. 2012 , 7, e34349	43
992	Effector CD4+ T cell expression signatures and immune-mediated disease associated genes. 2012 , 7, e38510	13
991	Genome-wide association study of African and European Americans implicates multiple shared and ethnic specific loci in sarcoidosis susceptibility. 2012 , 7, e43907	76
990	Limited evidence for parent-of-origin effects in inflammatory bowel disease associated loci. 2012 , 7, e45287	9
989	PTGER4 expression-modulating polymorphisms in the 5p13.1 region predispose to Crohn's disease and affect NF-B and XBP1 binding sites. 2012 , 7, e52873	33

988	Epigenetics and the developmental origins of inflammatory bowel diseases. 2012 , 26, 909-15	46
987	Non-MHC genes linked to autoimmune disease. 2012 , 32, 193-285	8
986	Evolutionary Insights into the P opulation-Specificity (b) the Genetic Factors Associated with Inflammatory Bowel Diseases. 2012 ,	1
985	The Role of the Microbiota in Gastrointestinal Health and Disease. 2012 ,	1
984	Innate and adaptive immunity in host-microbiota mutualism. 2012, 4, 685-98	11
983	Use of genome-wide association studies for drug repositioning. 2012 , 30, 317-20	275
982	HVEM signalling at mucosal barriers provides host defence against pathogenic bacteria. 2012, 488, 222-5	94
981	Amino acid position 11 of HLA-DRI is a major determinant of chromosome 6p association with ulcerative colitis. 2012 , 13, 245-52	30
980	Identification of cis-regulatory variation influencing protein abundance levels in human plasma. 2012 , 21, 3719-26	71
979	Role of nutrition and microbiota in susceptibility to inflammatory bowel diseases. 2012 , 56, 524-35	96
978	Digesting the genetics of inflammatory bowel disease: insights from studies of autophagy risk genes. 2012 , 18, 782-92	32
977	Genome-wide expression profiling implicates a MAST3-regulated gene set in colonic mucosal inflammation of ulcerative colitis patients. 2012 , 18, 1072-80	16
976	Etiology of pouchitis. 2012 , 18, 1146-55	45
975	Genome-wide methylation profiling in Crohn's disease identifies altered epigenetic regulation of key host defense mechanisms including the Th17 pathway. 2012 , 18, 889-99	121
974	Alterations in the gut microbiome of children with severe ulcerative colitis. 2012, 18, 1799-808	193
973	Serum anti-glycan antibody biomarkers for inflammatory bowel disease diagnosis and progression: a systematic review and meta-analysis. 2012 , 18, 1872-84	50
972	Mucosal genome-wide methylation changes in inflammatory bowel disease. 2012 , 18, 2128-37	103
971	Genome-wide peripheral blood leukocyte DNA methylation microarrays identified a single association with inflammatory bowel diseases. 2012 , 18, 2334-41	70

970	The immunogenetic architecture of autoimmune disease. 2012 , 4,	58
969	What is personalized medicine and what should it replace?. 2012 , 9, 418-24	60
968	Heparanase enzyme in chronic inflammatory bowel disease and colon cancer. 2012 , 69, 2501-13	27
967	Recent advances in understanding ulcerative colitis. 2012 , 7, 103-11	28
966	How does knowledge from translational research impact our clinical care of pediatric inflammatory bowel disease patients?. 2012 , 14, 275-81	3
965	Interaction between genetic and epigenetic variation defines gene expression patterns at the asthma-associated locus 17q12-q21 in lymphoblastoid cell lines. 2012 , 131, 1161-71	47
964	Five years of GWAS discovery. 2012 , 90, 7-24	1635
963	Use of a multiethnic approach to identify rheumatoid- arthritis-susceptibility loci, 1p36 and 17q12. 2012 , 90, 524-32	60
962	Identification of IRF8, TMEM39A, and IKZF3-ZPBP2 as susceptibility loci for systemic lupus erythematosus in a large-scale multiracial replication study. 2012 , 90, 648-60	134
961	A general framework for two-stage analysis of genome-wide association studies and its application to case-control studies. 2012 , 90, 760-73	21
960	A genome-wide association study reveals that variants within the HLA region are associated with risk for nonobstructive azoospermia. 2012 , 90, 900-6	54
959	Epithelial barrier: an interface for the cross-communication between gut flora and immune system. 2012 , 245, 147-63	127
958	Investigation of JAK2, STAT3 and CCR6 polymorphisms and their gene-gene interactions in inflammatory bowel disease. 2012 , 39, 247-52	25
957	Enterocyte STAT5 promotes mucosal wound healing via suppression of myosin light chain kinase-mediated loss of barrier function and inflammation. 2012 , 4, 109-24	54
956	Dominant disease-causing effect of NOD2 mutations in a family with all family members affected by Crohn's disease. 2012 , 18, 395-6	5
955	Associations between genetic variants in the IRGM gene and inflammatory bowel diseases in the Korean population. 2013 , 19, 106-14	37
954	IL-10R polymorphisms are associated with very-early-onset ulcerative colitis. 2013 , 19, 115-23	178
953	Drug-induced inflammatory bowel disease and IBD-like conditions. 2013 , 19, 445-56	38

952	Meta-analysis of genome-wide association studies identifies ten loci influencing allergic sensitization. <i>Nature Genetics</i> , 2013 , 45, 902-906	36.3	191
951	Expression quantitative trait loci analysis identifies associations between genotype and gene expression in human intestine. 2013 , 144, 1488-96, 1496.e1-3		51
950	Genome-wide association analysis identifies new susceptibility loci for Behæt's disease and epistasis between HLA-B*51 and ERAP1. <i>Nature Genetics</i> , 2013 , 45, 202-7	36.3	375
949	Mapping of immune-mediated disease genes. 2013 , 14, 325-53		92
948	Th1 not Th17 cells drive spontaneous MS-like disease despite a functional regulatory T cell response. 2013 , 126, 501-15		27
947	The microbiota and inflammatory bowel disease: insights from animal models. 2013 , 24, 102-6		49
946	What can we learn from inflammatory bowel disease in developing countries?. 2013, 15, 313		18
945	Antimicrobial Peptides and Innate Immunity. 2013,		8
944	The ubiquitin ligase parkin mediates resistance to intracellular pathogens. 2013 , 501, 512-6		387
943	Update on primary sclerosing cholangitis. 2013 , 59, 571-82		92
942	The genetics of complex cholestatic disorders. 2013 , 144, 1357-74		111
941	Refinement in localization and identification of gene regions associated with Crohn disease. 2013 , 92, 107-13		21
940	Identification of multiple genetic susceptibility loci in Takayasu arteritis. 2013, 93, 298-305		115
939	Oral contraceptives, reproductive factors and risk of inflammatory bowel disease. 2013 , 62, 1153-9		127
938	Novel insights into autoimmune liver diseases provided by genome-wide association studies. 2013 , 46, 41-54		82
937	Association between variants of PRDM1 and NDP52 and Crohn's disease, based on exome sequencing and functional studies. 2013 , 145, 339-47		125
936	Molecular Genetics of Inflammatory Bowel Disease. 2013,		
935	La mBecine gBomique, une rBlitlen pleine Dolution. PremiBe partie. 2013 , 28, 93-108		

934	Beyond gene discovery in inflammatory bowel disease: the emerging role of epigenetics. 2013 , 145, 293-308	206
933	The role of Haptoglobin and its related protein, Zonulin, in inflammatory bowel disease. 2013 , 1, e27321	81
932	Involvement of JAK/STAT signaling in the pathogenesis of inflammatory bowel disease. 2013, 76, 1-8	197
931	Keine spezifischen Einflussfaktoren. 2013 , 15, 14-15	
930	IRF5, but not TLR4, DEFB1, or VDR, is associated with the risk of ulcerative colitis in a Han Chinese population. 2013 , 48, 1145-51	10
929	Candidate genes for type 1 diabetes modulate pancreatic islet inflammation and Etell apoptosis. 2013 , 15 Suppl 3, 71-81	96
928	Mucosal CXCR4+ IgG plasma cells contribute to the pathogenesis of human ulcerative colitis through FcR-mediated CD14 macrophage activation. 2013 , 62, 1734-44	68
927	Influence of HLA-C expression level on HIV control. 2013 , 340, 87-91	277
926	Glycomics meets genomics, epigenomics and other high throughput omics for system biology studies. 2013 , 17, 34-40	45
925	Autophagy and intestinal homeostasis. 2013 , 75, 241-62	59
924	Immune aspects of the pathogenesis of inflammatory bowel disease. 2013 , 137, 283-97	64
923	Two susceptibility loci to Takayasu arteritis reveal a synergistic role of the IL12B and HLA-B regions in a Japanese population. 2013 , 93, 289-97	103
922	Examination of the relationship between variation at 17q21 and childhood wheeze phenotypes. 2013 , 131, 685-94	51
921	The GETECCU clinical guideline for the treatment of ulcerative colitis: A guideline created using GRADE methodology. 2013 , 36, 483.e1-483.e46	
920	[Therapeutic guidelines on ulcerative colitis: a GRADE methodology based effort of GETECCU]. 2013 , 36, e1-47	21
919	Genetic variants of membrane metallopeptidase genes in inflammatory bowel diseases. 2013 , 45, 1003-10	2
918	Immune-mediated disease genetics: the shared basis of pathogenesis. 2013 , 34, 22-6	71
917	Animal models of chemically induced intestinal inflammation: predictivity and ethical issues. 2013 , 139, 71-86	33

916	Effect of barrier microbes on organ-based inflammation. 2013 , 131, 1465-78	45
915	Autophagy and inflammatory diseases. 2013 , 91, 250-8	88
914	The cell biology of the intestinal epithelium and its relation to inflammatory bowel disease. 2013 , 45, 798-806	17
913	Genome-wide association analysis in primary sclerosing cholangitis and ulcerative colitis identifies risk loci at GPR35 and TCF4. 2013 , 58, 1074-83	118
912	The genetics of innate immunity sensors and human disease. 2013 , 32, 157-208	17
911	IL-1 receptor 2 (IL-1R2) and its role in immune regulation. 2013 , 32, 1-8	123
910	Association of ZAP70 and PTPN6, but Not BANK1 or CLEC2D, with inflammatory bowel disease in the Tunisian population. 2013 , 17, 321-6	13
909	Genome-wide linkage scan for psoriasis susceptibility loci in multiplex Tunisian families. 2013 , 168, 583-7	17
908	Dysplastic lesions in inflammatory bowel disease: molecular pathogenesis to morphology. 2013 , 137, 338-50	50
907	Fate determination of mature autoreactive B cells. 2013 , 118, 1-36	7
907	Fate determination of mature autoreactive B cells. 2013 , 118, 1-36 Darmgesundheit und Mikrobiota. 2013 , 67-83	7
		7 563
906	Darmgesundheit und Mikrobiota. 2013 , 67-83 GWAS of 126,559 individuals identifies genetic variants associated with educational attainment.	
906	Darmgesundheit und Mikrobiota. 2013 , 67-83 GWAS of 126,559 individuals identifies genetic variants associated with educational attainment. 2013 , 340, 1467-71	563
906 905 904	Darmgesundheit und Mikrobiota. 2013, 67-83 GWAS of 126,559 individuals identifies genetic variants associated with educational attainment. 2013, 340, 1467-71 Ulcerative colitis. 2013, 346, f432	563
906 905 904 903	Darmgesundheit und Mikrobiota. 2013, 67-83 GWAS of 126,559 individuals identifies genetic variants associated with educational attainment. 2013, 340, 1467-71 Ulcerative colitis. 2013, 346, f432 Host genotype, intestinal microbiota and inflammatory disorders. 2013, 109 Suppl 2, S76-80 Global chromatin state analysis reveals lineage-specific enhancers during the initiation of human T	563 99 25
906 905 904 903 902	Darmgesundheit und Mikrobiota. 2013, 67-83 GWAS of 126,559 individuals identifies genetic variants associated with educational attainment. 2013, 340, 1467-71 Ulcerative colitis. 2013, 346, f432 Host genotype, intestinal microbiota and inflammatory disorders. 2013, 109 Suppl 2, S76-80 Global chromatin state analysis reveals lineage-specific enhancers during the initiation of human T helper 1 and T helper 2 cell polarization. 2013, 38, 1271-84	563 99 25 70

898	Effector and memory T cell responses to commensal bacteria. 2013 , 34, 299-306	52
897	Large sample size, wide variant spectrum, and advanced machine-learning technique boost risk prediction for inflammatory bowel disease. 2013 , 92, 1008-12	114
896	Protective and pathological properties of IL-22 in liver disease: implications for viral hepatitis. 2013 , 182, 21-8	50
895	Moving the genetics of inflammatory bowel diseases from bench to bedside: first steps towards personalised medicine. 2013 , 62, 1531-3	8
894	R-spondin 2 signalling mediates susceptibility to fatal infectious diarrhoea. 2013 , 4, 1898	52
893	Innate immunity and inflammatory bowel disease: a review of clinical evidence and future application. 2013 , 6, 415-9	5
892	Inflammatory bowel disease: mechanisms, redox considerations, and therapeutic targets. 2013 , 19, 1711-47	143
891	New insights on the pathomechanisms of inflammatory bowel disease. 2013 , 14, 455-62	4
890	A genome-wide association study on a southern European population identifies a new Crohn's disease susceptibility locus at RBX1-EP300. 2013 , 62, 1440-5	35
889	Cleavage and polyadenylation specificity factor 1 (CPSF1) regulates alternative splicing of interleukin 7 receptor (IL7R) exon 6. 2013 , 19, 103-15	25
888	Protein therapeutics targeted at the TNF superfamily. 2013 , 66, 51-80	14
887	Inflammasome in intestinal inflammation and cancer. 2013 , 2013, 654963	36
886	Loci associated with N-glycosylation of human immunoglobulin G show pleiotropy with autoimmune diseases and haematological cancers. 2013 , 9, e1003225	242
885	All SNPs are not created equal: genome-wide association studies reveal a consistent pattern of enrichment among functionally annotated SNPs. 2013 , 9, e1003449	209
884	Deep resequencing of GWAS loci identifies rare variants in CARD9, IL23R and RNF186 that are associated with ulcerative colitis. 2013 , 9, e1003723	149
883	A genome-wide association study reveals ARL15, a novel non-HLA susceptibility gene for rheumatoid arthritis in North Indians. 2013 , 65, 3026-35	22
882	Phenotype and function of B cells and dendritic cells from interferon regulatory factor 5-deficient mice with and without a mutation in DOCK2. 2013 , 25, 295-306	39
881	[Inflammatory bowel disease]. 2013, 102, 1046-52	

880	Susceptibility loci associated with specific and shared subtypes of lymphoid malignancies. 2013 , 9, e1003220	38
879	PUMA: a unified framework for penalized multiple regression analysis of GWAS data. 2013 , 9, e1003101	26
878	Preferential binding to Elk-1 by SLE-associated IL10 risk allele upregulates IL10 expression. 2013 , 9, e1003870	33
877	The role of TL1A and DR3 in autoimmune and inflammatory diseases. 2013 , 2013, 258164	48
876	Irf8-regulated genomic responses drive pathological inflammation during cerebral malaria. 2013 , 9, e1003491	38
875	High intestinal and systemic levels of interleukin-23/T-helper 17 pathway in Chinese patients with inflammatory bowel disease. 2013 , 2013, 425915	25
874	Autophagy proteins control goblet cell function by potentiating reactive oxygen species production. 2013 , 32, 3130-44	165
873	Dysregulated phosphatidylinositol signaling promotes endoplasmic-reticulum-stress-mediated intestinal mucosal injury and inflammation in zebrafish. 2014 , 7, 93-106	38
872	Ethnicity differences in genetic susceptibility to ulcerative colitis: a comparison of Indian asians and white northern Europeans. 2013 , 19, 2888-94	9
871	Stories of love and hate: innate immunity and host-microbe crosstalk in the intestine. 2013 , 29, 125-32	29
870	Gene expression profiles of ileal inflammatory bowel disease correlate with disease phenotype and advance understanding of its immunopathogenesis. 2013 , 19, 2509-21	29
869	Replication study of ulcerative colitis risk loci in a Lithuanian-Latvian case-control sample. 2013 , 19, 2349-55	10
868	Genetic architecture of human fibrotic diseases: disease risk and disease progression. 2013 , 4, 159	10
867	Common genetic variation and antidepressant efficacy in major depressive disorder: a meta-analysis of three genome-wide pharmacogenetic studies. 2013 , 170, 207-17	181
866	Genetic variants in the IL12B gene are associated with inflammatory bowel diseases in the Korean population. 2013 , 28, 1588-94	14
865	Taking Crohn's disease personally. 2013 , 4, e0011	1
864	CC Chemokine Ligand 20 and Its Cognate Receptor CCR6 in Mucosal T Cell Immunology and Inflammatory Bowel Disease: Odd Couple or Axis of Evil?. 2013 , 4, 194	72
863	Associations between genetic polymorphisms in IL-33, IL1R1 and risk for inflammatory bowel disease. 2013 , 8, e62144	68

(2013-2013)

862	Validation of reported genetic risk factors for periodontitis in a large-scale replication study. 2013 , 40, 563-72	68
861	HVEM: An unusual TNF receptor family member important for mucosal innate immune responses to microbes. 2013 , 4, 146-51	19
860	Targeted resequencing identifies defective variants of decoy receptor 3 in pediatric-onset inflammatory bowel disease. 2013 , 14, 447-52	15
859	Association of ANXA11 genetic variation with sarcoidosis in African Americans and European Americans. 2013 , 14, 13-8	39
858	Phenotypic manifestations of inflammatory bowel disease differ between Hispanics and non-Hispanic whites: results of a large cohort study. 2013 , 108, 231-9	42
857	A PheWAS approach in studying HLA-DRB1*1501. 2013 , 14, 187-91	75
856	Next generation exome sequencing of paediatric inflammatory bowel disease patients identifies rare and novel variants in candidate genes. 2013 , 62, 977-84	92
855	The microbiome in inflammatory bowel disease and beyond. 2013 , 13 Suppl 6, s29-31	
854	Systematic functional regulatory assessment of disease-associated variants. 2013, 110, 9607-12	75
853	Gene-centric analysis identifies variants associated with interleukin-6 levels and shared pathways with other inflammation markers. 2013 , 6, 163-70	34
852	Inherited predisposition to myeloproliferative neoplasms. 2013 , 4, 237-53	40
851	Phosphatase regulation of intercellular junctions. 2013 , 1, e26713	20
850	Role of Autophagy-Related Genes in the Pathology of Inflammatory Bowel Disease. 2013,	1
849	Genome-wide methylation analyses of primary human leukocyte subsets identifies functionally important cell-type-specific hypomethylated regions. 2013 , 122, e52-60	56
848	Genetic and functional identification of the likely causative variant for cholesterol gallstone disease at the ABCG5/8 lithogenic locus. 2013 , 57, 2407-17	61
847	Genome-wide association study of ulcerative colitis in Koreans suggests extensive overlapping of genetic susceptibility with Caucasians. 2013 , 19, 954-66	62
846	Genetic susceptibility in IBD: overlap between ulcerative colitis and Crohn's disease. 2013, 19, 240-5	32
845	Impact of allele copy number of polymorphisms in FCGR3A and FCGR3B genes on susceptibility to ulcerative colitis. 2013 , 19, 2061-8	9

844	Association of the MEFV gene variations with inflammatory bowel disease in Turkey. 2013 , 47, e23-7		27
843	Spontaneous regression of colonic lymphoma following infliximab and azathioprine withdrawal in patients with Crohn's disease. 2013 , 19, E69-70		9
842	Ulcerative colitis with rare chromosomal abnormalities: report of 2 patients. 2013 , 19, E68-9		
841	Idiopathic myelofibrosis and pyoderma gangrenosum involving a mutation of Janus kinase 2 (JAK2V617F), showing poor prognosis. 2013 , 23, 256-7		7
840	The role of the innate and adaptive immune system in pediatric inflammatory bowel disease. 2013 , 19, 2011-20		7
839	Inflammatory Bowel Disease. 2013 , 863-878		
838	DNA methylation in inflammatory bowel disease and beyond. <i>World Journal of Gastroenterology</i> , 2013 , 19, 5238-49	5.6	26
837	Rac2-deficiency leads to exacerbated and protracted colitis in response to Citrobacter rodentium infection. 2013 , 8, e61629		19
836	Characterization of the gut-associated microbiome in inflammatory pouch complications following ileal pouch-anal anastomosis. 2013 , 8, e66934		63
835	No evidence of association between common autoimmunity STAT4 and IL23R risk polymorphisms and non-anterior uveitis. 2013 , 8, e72892		2
834	Gene expression analysis of peripheral cells for subclassification of pediatric inflammatory bowel disease in remission. 2013 , 8, e79549		10
833	Functional consequences of the macrophage stimulating protein 689C inflammatory bowel disease risk allele. 2013 , 8, e83958		13
832	Animal models of ulcerative colitis and their application in drug research. 2013, 7, 1341-57		97
831	Why does melanoma metastasize into the brain? Genes with pleiotropic effects might be the key. 2013 , 4, 75		3
830	Regulation of intestinal homeostasis by innate immune cells. 2013 , 13, 227-34		23
829	miR-20b, miR-98, miR-125b-1*, and let-7e* as new potential diagnostic biomarkers in ulcerative colitis. <i>World Journal of Gastroenterology</i> , 2013 , 19, 4289-99	5.6	67
828	Structure-function analysis of the C-clamp of TCF/Pangolin in Wnt/Etatenin signaling. 2014 , 9, e86180		13
827	Allele-specific methylation occurs at genetic variants associated with complex disease. 2014 , 9, e98464		29

(2014-2014)

826	Polymorphisms in the inflammatory pathway genes TLR2, TLR4, TLR9, LY96, NFKBIA, NFKB1, TNFA, TNFRSF1A, IL6R, IL10, IL23R, PTPN22, and PPARG are associated with susceptibility of inflammatory bowel disease in a Danish cohort. 2014 , 9, e98815	75
825	The NOD2 p.Leu1007fsX1008 mutation (rs2066847) is a stronger predictor of the clinical course of Crohn's disease than the FOXO3A intron variant rs12212067. 2014 , 9, e108503	20
824	An image-based genetic assay identifies genes in T1D susceptibility loci controlling cellular antiviral immunity in mouse. 2014 , 9, e108777	4
823	Genetics of eosinophilic esophagitis. 2014 , 32, 22-9	4
822	Intestinal epithelium in inflammatory bowel disease. 2014 , 1, 24	99
821	Identifying human disease genes: advances in molecular genetics and computational approaches. 2014 , 13, 5073-87	7
820	Secretory IgA is Concentrated in the Outer Layer of Colonic Mucus along with Gut Bacteria. 2014 , 3, 390-403	90
819	Designing biologic selectivity for inflammatory bowel diseaserole of vedolizumab. 2015 , 9, 147-54	6
818	Association among genetic predisposition, gut microbiota, and host immune response in the etiopathogenesis of inflammatory bowel disease. 2014 , 47, 727-37	37
817	Performance of the Montreal classification for inflammatory bowel diseases. <i>World Journal of Gastroenterology</i> , 2014 , 20, 15374-81	35
816	HLA in Gastrointestinal Inflammatory Disorders. 2014,	O
815	Estimation and partitioning of (co)heritability of inflammatory bowel disease from GWAS and immunochip data. 2014 , 23, 4710-20	73
814	Association of the interleukin-22 genetic polymorphisms with ulcerative colitis. 2014 , 9, 183	7
813	Haemolysin of Escherichia coli in IBD: a potentiator of inflammatory activity in the colon. 2014 , 63, 1893-901	43
812	Associations between TNFSF15 polymorphisms and susceptibility to ulcerative colitis and Crohn's disease: A meta-analysis. 2014 , 47, 512-8	9
811	Anti-TNF antibody-induced psoriasiform skin lesions in patients with inflammatory bowel disease are characterised by interferon-æxpressing Th1 cells and IL-17A/IL-22-expressing Th17 cells and respond to anti-IL-12/IL-23 antibody treatment. 2014 , 63, 567-77	210
810	Higher activity of the inducible nitric oxide synthase contributes to very early onset inflammatory bowel disease. 2014 , 5, e46	55
809	Multi-omics analysis of inflammatory bowel disease. 2014 , 162, 62-8	28

808	Pleiotropic associations of risk variants identified for other cancers with lung cancer risk: the PAGE and TRICL consortia. 2014 , 106, dju061	28
807	Methylation QTLs are associated with coordinated changes in transcription factor binding, histone modifications, and gene expression levels. 2014 , 10, e1004663	187
806	Crohn's disease susceptibility variants in Colombian tuberculosis patients. 2014 , 18, 89-94	1
805	[Second European evidence-based Consensus on the diagnosis and management of ulcerative colitis Part 1: Definitions and diagnosis (Spanish version)]. 2014 , 79, 263-89	11
804	Immune deficiency-related enteropathy-lymphocytopenia-alopecia syndrome results from tetratricopeptide repeat domain 7A deficiency. 2014 , 134, 1354-1364.e6	48
803	Towards personalized care in IBD. 2014 , 11, 287-99	46
802	Genetic polymorphisms of tumour necrosis factor alpha (TNF-#promoter gene and response to TNF-#nhibitors in Spanish patients with inflammatory bowel disease. 2014 , 41, 63-8	21
801	The microbiome in inflammatory bowel disease and its modulation as a therapeutic manoeuvre. 2014 , 73, 452-6	7
800	A genome-wide association study identifies a functional ERAP2 haplotype associated with birdshot chorioretinopathy. 2014 , 23, 6081-7	82
799	IL-17A alone weakly affects the transcriptome of intestinal epithelial cells but strongly modulates the TNF-Induced expression of inflammatory mediators and inflammatory bowel disease susceptibility genes. 2014 , 20, 1502-15	10
798	IBD candidate genes and intestinal barrier regulation. 2014 , 20, 1829-49	95
797	Genetics of Behat's disease: lessons learned from genomewide association studies. 2014 , 26, 56-63	55
796	Mucosal transcriptomics implicates under expression of BRINP3 in the pathogenesis of ulcerative colitis. 2014 , 20, 1802-12	22
795	Overexpression of ATP-activated P2X7 receptors in the intestinal mucosa is implicated in the pathogenesis of Crohn's disease. 2014 , 20, 444-57	62
794	Clinical, serologic, and genetic factors associated with pyoderma gangrenosum and erythema nodosum in inflammatory bowel disease patients. 2014 , 20, 525-33	42
793	HVEM is a TNF Receptor with Multiple Regulatory Roles in the Mucosal Immune System. 2014 , 14, 67-72	15
792	Genetics and innate and adaptive immunity in IBD. 2014 , 79, 41-55	14
791	Inflammatory bowel diseases: from pathogenesis to laboratory testing. 2014 , 52, 471-81	23

(2014-2014)

790	NFIL3-deficient mice develop microbiota-dependent, IL-12/23-driven spontaneous colitis. 2014 , 192, 1918-27	31
789	Cell death and inflammatory bowel diseases: apoptosis, necrosis, and autophagy in the intestinal epithelium. 2014 , 2014, 218493	70
788	Targeting interleukins for the treatment of inflammatory bowel disease-what lies beyond anti-TNF therapy?. 2014 , 20, 389-97	26
787	Genetic literacy and patient perceptions of IBD testing utility and disease control: a randomized vignette study of genetic testing. 2014 , 20, 901-8	26
786	Copper metabolism domain-containing 1 represses genes that promote inflammation and protects mice from colitis and colitis-associated cancer. 2014 , 147, 184-195.e3	24
7 ⁸ 5	Association of CTLA-4 variants with susceptibility to inflammatory bowel disease: a meta-analysis. 2014 , 75, 227-33	13
784	Associations between PTPN2 polymorphisms and susceptibility to ulcerative colitis and Crohn's disease: a meta-analysis. 2014 , 63, 71-9	11
783	IgA measurements in over 12 000 Swedish twins reveal sex differential heritability and regulatory locus near CD30L. 2014 , 23, 4177-84	9
782	Beneficial modulation of the gut microbiota. 2014 , 588, 4120-30	166
781	Novel rheumatoid arthritis susceptibility locus at 22q12 identified in an extended UK genome-wide association study. 2014 , 66, 24-30	36
780	Genome-wide association study identifies multiple loci associated with bladder cancer risk. 2014 , 23, 1387-98	101
779	Recent advances in Takayasu arteritis. 2014 , 17, 238-47	71
778	Genetic Epidemiology of Psoriasis. 2014 , 3, 61-78	55
777	A genome-wide association study follow-up suggests a possible role for PPARG in systemic sclerosis susceptibility. 2014 , 16, R6	35
776	Oral caffeine administration ameliorates acute colitis by suppressing chitinase 3-like 1 expression in intestinal epithelial cells. 2014 , 49, 1206-16	32
775	Innate immune activity conditions the effect of regulatory variants upon monocyte gene expression. 2014 , 343, 1246949	507
774	Genetic Predisposition, Humans. 2014 , 341-364	1
773	Inflammatory Bowel Diseases. 2014 , 873-888	

772	Next generation sequencing techniques in neurological diseases: redefining clinical and molecular associations. 2014 , 23, R47-53	48
771	Genetic comorbidities in Parkinson's disease. 2014 , 23, 831-41	49
770	Recent insights on the putative role of autophagy in autoimmune diseases. 2014 , 13, 231-41	32
769	Do inhibitory immune receptors play a role in the etiology of autoimmune disease?. 2014 , 150, 31-42	18
768	Impact of genetic polymorphisms on the pathogenesis of idiopathic achalasia: Association with IL33 gene variant. 2014 , 75, 364-9	8
767	Genetic and Environmental Risk Factors for Multiple Sclerosis Role for Interaction Analysis. 2014 , 101-114	1
766	GWAS identifies four novel eosinophilic esophagitis loci. 2014 , 5, 5593	135
765	Genetic variation at the CELF1 (CUGBP, elav-like family member 1 gene) locus is genome-wide associated with Alzheimer's disease and obesity. 2014 , 165B, 283-93	28
764	RNA sequencing identifies dysregulation of the human pancreatic islet transcriptome by the saturated fatty acid palmitate. 2014 , 63, 1978-93	174
763	Covariate-modulated local false discovery rate for genome-wide association studies. 2014 , 30, 2098-104	34
762	Secretory antibodies in breast milk promote long-term intestinal homeostasis by regulating the gut microbiota and host gene expression. 2014 , 111, 3074-9	275
761	Family history of inflammatory bowel disease among patients with ulcerative colitis: a systematic review and meta-analysis. 2014 , 8, 1480-97	39
760	HLA-DQA1-HLA-DRB1 variants confer susceptibility to pancreatitis induced by thiopurine immunosuppressants. <i>Nature Genetics</i> , 2014 , 46, 1131-4	130
759	Sequencing-based approach identified three new susceptibility loci for psoriasis. 2014 , 5, 4331	56
758	A large candidate-gene association study suggests genetic variants at IRF5 and PRDM1 to be associated with aggressive periodontitis. 2014 , 41, 1122-31	20
757	Genetic and microbial factors modulating the ubiquitin proteasome system in inflammatory bowel disease. 2014 , 63, 1265-74	58
756	Peptidoglycan recognition protein 3 and Nod2 synergistically protect mice from dextran sodium sulfate-induced colitis. 2014 , 193, 3055-69	23
755	Variants in nicotinamide adenine dinucleotide phosphate oxidase complex components determine susceptibility to very early onset inflammatory bowel disease. 2014 , 147, 680-689.e2	88

(2014-2014)

754	Human NK cells licensed by killer Ig receptor genes have an altered cytokine program that modifies CD4+ T cell function. 2014 , 193, 940-9	26
753	Genome-wide association study of Crohn's disease in Koreans revealed three new susceptibility loci and common attributes of genetic susceptibility across ethnic populations. 2014 , 63, 80-7	130
75 ²	A genome-wide association study identifies a novel locus at 6q22.1 associated with ulcerative colitis. 2014 , 23, 6927-34	31
751	The role of glycosylation in IBD. 2014 , 11, 588-600	93
750	A transgenic probiotic secreting a parasite immunomodulator for site-directed treatment of gut inflammation. 2014 , 22, 1730-40	47
749	Genetics of canine anal furunculosis in the German shepherd dog. 2014 , 66, 311-24	12
748	Clinical presentation and disease course of inflammatory bowel disease differs by race in a large tertiary care hospital. 2014 , 59, 2228-35	26
747	Microbiota activation and regulation of innate and adaptive immunity. 2014 , 260, 206-20	81
746	Evidence for non-neutralizing autoantibodies against IL-10 signalling components in patients with inflammatory bowel disease. 2014 , 15, 10	7
745	Diet, gut microbes, and genetics in immune function: can we leverage our current knowledge to achieve better outcomes in inflammatory bowel diseases?. 2014 , 31, 16-23	19
744	An excess of risk-increasing low-frequency variants can be a signal of polygenic inheritance in complex diseases. 2014 , 94, 437-52	37
743	Many inflammatory bowel disease risk loci include regions that regulate gene expression in immune cells and the intestinal epithelium. 2014 , 146, 1040-7	82
742	Genetic susceptibility and genotype-phenotype association in 588 Danish children with inflammatory bowel disease. 2014 , 8, 678-85	19
741	Nod/Ripk2 signaling in dendritic cells activates IL-17A-secreting innate lymphoid cells and drives colitis in T-bet-/Rag2-/- (TRUC) mice. 2014 , 111, E2559-66	48
740	The role of type I interferons in intestinal infection, homeostasis, and inflammation. 2014 , 260, 145-67	45
739	Cooperativity among secretory IgA, the polymeric immunoglobulin receptor, and the gut microbiota promotes host-microbial mutualism. 2014 , 162, 10-21	100
738	Epithelial gp130/Stat3 functions: an intestinal signaling node in health and disease. 2014 , 26, 29-37	40
737	Innate and adaptive immunity in inflammatory bowel disease. 2014 , 13, 3-10	467

736	The microbial basis of inflammatory bowel diseases. 2014 , 124, 4190-6	129
735	Inflammatory bowel disease: pathogenesis. <i>World Journal of Gastroenterology</i> , 2014 , 20, 91-9 5.6	543
734	Functional characterization of the human dendritic cell immunodeficiency associated with the IRF8(K108E) mutation. 2014 , 124, 1894-904	51
733	Role of genetics in pediatric inflammatory bowel disease. 2014 , 20, 1878-84	12
732	Stat3: friend or foe in colitis and colitis-associated cancer?. 2014 , 20, 2405-11	28
731	Genome-Wide Association Studies in Primary Biliary Cirrhosis. 2015 , 35, 392-401	45
730	Emerging leadership lecture: Inflammatory bowel disease in Asia: emergence of a "Western" disease. 2015 , 30, 440-5	64
729	Rare and low-frequency variants and their association with plasma levels of fibrinogen, FVII, FVIII, and vWF. 2015 , 126, e19-29	45
728	Estimating heritability of drug-induced liver injury from common variants and implications for future study designs. 2014 , 4, 5762	4
727	Differential diagnosis in inflammatory bowel disease colitis: state of the art and future perspectives. World Journal of Gastroenterology, 2015 , 21, 21-46	108
727 726		108
	perspectives. World Journal of Gastroenterology, 2015 , 21, 21-46	
726	perspectives. World Journal of Gastroenterology, 2015 , 21, 21-46 G Protein-coupled pH-sensing Receptor OGR1 Is a Regulator of Intestinal Inflammation. 2015 , 21, 1269-81	46
726 725	perspectives. World Journal of Gastroenterology, 2015, 21, 21-46 G Protein-coupled pH-sensing Receptor OGR1 Is a Regulator of Intestinal Inflammation. 2015, 21, 1269-81 From genetics to treatment of eosinophilic esophagitis. 2015, 15, 417-25 PTGER4 gene variant rs76523431 is a candidate risk factor for radiological joint damage in	46 19
726 725 724	perspectives. World Journal of Gastroenterology, 2015, 21, 21-46 G Protein-coupled pH-sensing Receptor OGR1 Is a Regulator of Intestinal Inflammation. 2015, 21, 1269-81 From genetics to treatment of eosinophilic esophagitis. 2015, 15, 417-25 PTGER4 gene variant rs76523431 is a candidate risk factor for radiological joint damage in rheumatoid arthritis patients: a genetic study of six cohorts. 2015, 17, 306 Technetium-99 m labeling and evaluation of olsalazine: a novel agent for ulcerative colitis imaging.	46 19 10
726 725 724 723	perspectives. World Journal of Gastroenterology, 2015, 21, 21-46 G Protein-coupled pH-sensing Receptor OGR1 Is a Regulator of Intestinal Inflammation. 2015, 21, 1269-81 From genetics to treatment of eosinophilic esophagitis. 2015, 15, 417-25 PTGER4 gene variant rs76523431 is a candidate risk factor for radiological joint damage in rheumatoid arthritis patients: a genetic study of six cohorts. 2015, 17, 306 Technetium-99 m labeling and evaluation of olsalazine: a novel agent for ulcerative colitis imaging. 2015, 58, 336-41	46 19 10
726 725 724 723 722	perspectives. World Journal of Gastroenterology, 2015, 21, 21-46 G Protein-coupled pH-sensing Receptor OGR1 Is a Regulator of Intestinal Inflammation. 2015, 21, 1269-81 From genetics to treatment of eosinophilic esophagitis. 2015, 15, 417-25 PTGER4 gene variant rs76523431 is a candidate risk factor for radiological joint damage in rheumatoid arthritis patients: a genetic study of six cohorts. 2015, 17, 306 Technetium-99 m labeling and evaluation of olsalazine: a novel agent for ulcerative colitis imaging. 2015, 58, 336-41 Sirtuin/uncoupling protein gene variants and carotid plaque area and morphology. 2015, 10, 1247-52 Genome-wide Pathway Analysis Using Gene Expression Data of Colonic Mucosa in Patients with	46 19 10 12

718	Serum Proteome Profiles in Stricturing Crohn's Disease: A Pilot Study. 2015 , 21, 1935-41	20
717	Paradoxical psoriasis in a large cohort of patients with inflammatory bowel disease receiving treatment with anti-TNF alpha: 5-year follow-up study. 2015 , 42, 880-8	60
716	Very early-onset inflammatory bowel disease: gaining insight through focused discovery. 2015 , 21, 1166-75	67
715	The genetic architecture of inflammatory bowel disease: past, present and future. 2015 , 31, 456-63	21
714	Regulation of intestinal inflammation through interaction of intestinal environmental factors and innate immune cells. 2015 , 35, 028-041	
713	Autophagy in colorectal cancer: An important switch from physiology to pathology. 2015 , 7, 271-84	93
712	Association of ATG5 Gene Polymorphisms With Behlet's Disease and ATG10 Gene Polymorphisms With VKH Syndrome in a Chinese Han Population. 2015 , 56, 8280-7	19
711	Association of Genetic Variations in TNFSF15 With Acute Anterior Uveitis in Chinese Han. 2015 , 56, 4605-10	15
710	Genetics of inflammatory bowel disease from multifactorial to monogenic forms. World Journal of Gastroenterology, 2015 , 21, 12296-310	87
709	Gene Expression-Genotype Analysis Implicates GSDMA, GSDMB, and LRRC3C as Contributors to Inflammatory Bowel Disease Susceptibility. 2015 , 2015, 834805	25
708	Pathology of Inflammatory Bowel Disease. 2015 , 5, 756-765	
707	Application of Comparative Transcriptional Genomics to Identify Molecular Targets for Pediatric IBD. 2015 , 6, 165	13
706	Mechanisms of Microbe-Host Interaction in Crohn's Disease: Dysbiosis vs. Pathobiont Selection. 2015 , 6, 555	51
705	Interleukin(IL)-36 and IL-36 Induce Proinflammatory Mediators from Human Colonic Subepithelial Myofibroblasts. 2015 , 2, 69	17
704	Genetic relationships between internal diseases diagnosed at slaughter and carcass traits in Japanese Black cattle. 2015 , 93, 2714-21	7
703	Endogenous levels of circulating androgens and risk of Crohn's disease and ulcerative colitis among women: a nested case-control study from the nurses' health study cohorts. 2015 , 21, 1378-85	13
702	Exome analysis of patients with concurrent pediatric inflammatory bowel disease and autoimmune disease. 2015 , 21, 1229-36	14
701	Dissecting Allele Architecture of Early Onset IBD Using High-Density Genotyping. 2015 , 10, e0128074	33

700	Association Study of IL-12B Polymorphisms Susceptibility with Ankylosing Spondylitis in Mainland Han Population. 2015 , 10, e0130982	18
699	Type 1 Diabetes Prone NOD Mice Have Diminished Cxcr1 mRNA Expression in Polymorphonuclear Neutrophils and CD4+ T Lymphocytes. 2015 , 10, e0134365	7
698	Allele-Selective Transcriptome Recruitment to Polysomes Primed for Translation: Protein-Coding and Noncoding RNAs, and RNA Isoforms. 2015 , 10, e0136798	9
697	Polymorphisms in PRKCDBP, a Transcriptional Target of TNF-BAre Associated With Inflammatory Bowel Disease in Korean. 2015 , 13, 242-9	3
696	Cyclic AMP-Responsive Element Modulator Polymorphisms Are Potential Genetic Risks for Systemic Lupus Erythematosus. 2015 , 2015, 906086	4
695	The role of innate immunity receptors in the pathogenesis of inflammatory bowel disease. 2015 , 2015, 936193	36
694	Role of TLR4 rs4986790A>G and rs4986791C>T Polymorphisms in the Risk of Inflammatory Bowel Disease. 2015 , 2015, 141070	7
693	Fine mapping of type 1 diabetes susceptibility loci and evidence for colocalization of causal variants with lymphoid gene enhancers. <i>Nature Genetics</i> , 2015 , 47, 381-6	414
692	Loss of n-6 fatty acid induced pediatric obesity protects against acute murine colitis. 2015 , 29, 3151-9	14
691	Inflammatory bowel disease associates with proinflammatory potential of the immunoglobulin G glycome. 2015 , 21, 1237-47	108
690	Programmed cell death and its role in inflammation. 2015 , 2, 12	109
689	Life Extension. 2015,	O
688	Enhanced meta-analysis and replication studies identify five new psoriasis susceptibility loci. 2015 , 6, 7001	122
687	B Cells Producing Pathogenic Autoantibodies. 2015 , 417-439	
686	Circulating microRNAs in Disease Diagnostics and their Potential Biological Relevance. 2015,	7
685	The genetic basis of eosinophilic esophagitis. 2015 , 29, 701-707	12
684	IL7R gene expression network associates with human healthy ageing. 2015 , 12, 21	29
683	The genetics of Crohn's disease and ulcerative colitisstatus quo and beyond. 2015 , 50, 13-23	59

(2015-2015)

682	NOD-like receptors: versatile cytosolic sentinels. 2015 , 95, 149-78	191
681	Intestinal steroidogenesis. 2015 , 103, 64-71	22
680	The allele T of rs10852936 confers risk for early-onset psoriasis. 2015 , 77, 129-31	1
679	Association of a mutation in LACC1 with a monogenic form of systemic juvenile idiopathic arthritis. 2015 , 67, 288-95	85
678	Runx3 at the interface of immunity, inflammation and cancer. 2015 , 1855, 131-43	40
677	Tumor necrosis factor superfamily in innate immunity and inflammation. 2014 , 7, a016279	56
676	Identification of Susceptibility Loci in IL6, RPS9/LILRB3, and an Intergenic Locus on Chromosome 21q22 in Takayasu Arteritis in a Genome-Wide Association Study. 2015 , 67, 1361-8	60
675	Association between CARD8 rs2043211 polymorphism and inflammatory bowel disease: a meta-analysis. 2015 , 44, 253-64	10
674	[Medical therapy of ulcerative colitis]. 2015 , 157, 45-8	1
673	The genetic architecture of the human immune system: a bioresource for autoimmunity and disease pathogenesis. 2015 , 161, 387-403	179
672	Genetic polymorphism in ATG16L1 gene influences the response to adalimumab in Crohn's disease patients. 2015 , 16, 191-204	36
671	Early investigational TNF receptor antagonists for the treatment of ulcerative colitis. 2015 , 24, 761-8	5
670	Contribution of NKX2-3 polymorphisms to inflammatory bowel diseases: a meta-analysis of 35358 subjects. 2014 , 4, 3924	6
669	Commensal bacteria direct selective cargo sorting to promote symbiosis. 2015 , 16, 918-26	133
668	Complex host genetic susceptibility to Staphylococcus aureus infections. 2015 , 23, 529-36	22
667	Insight in genome-wide association of metabolite quantitative traits by exome sequence analyses. 2015 , 11, e1004835	49
666	Autophagy and inflammatory bowel disease: Association between variants of the autophagy-related IRGM gene and susceptibility to Crohn's disease. 2015 , 47, 744-50	25
665	Decreased serum level of IL-7 in patients with active Graves' disease. 2015 , 75, 373-9	4

664	Systematic analysis of circadian genes using genome-wide cDNA microarrays in the inflammatory bowel disease transcriptome. 2015 , 32, 903-16	31
663	Evaluation of European coeliac disease risk variants in a north Indian population. 2015 , 23, 530-5	11
662	Parallels Between Mammals and Flies in Inflammatory Bowel Disease. 2015, 151-189	1
661	High-resolution gene expression profiling using RNA sequencing in patients with inflammatory bowel disease and in mouse models of colitis. 2015 , 9, 492-506	42
660	IL-12 and IL-23 cytokines: from discovery to targeted therapies for immune-mediated inflammatory diseases. 2015 , 21, 719-29	488
659	Takayasu arteritis and ulcerative colitis: high rate of co-occurrence and genetic overlap. 2015 , 67, 2226-32	79
658	Association of vitamin D receptor gene polymorphisms with the susceptibility to ulcerative colitis in patients from Southeast China. 2015 , 35, 530-5	9
657	Differences between adults and children: genetics and beyond. 2015 , 9, 191-6	6
656	Re-annotation of presumed noncoding disease/trait-associated genetic variants by integrative analyses. 2015 , 5, 9453	12
655	Genetic architectures of seropositive and seronegative rheumatic diseases. 2015 , 11, 401-14	30
654	Down the line from genome-wide association studies in inflammatory bowel disease: the resulting clinical benefits and the outlook for the future. 2015 , 11, 33-44	13
653	Management der chronisch-entzfidlichen Darmerkrankungen wfirend der Schwangerschaft. 2015 , 48, 131-138	
652	Contribution of susceptibility variants at FCGR2A and 13q12 to the risk of relapse among Japanese patients with ulcerative colitis. 2015 , 50, 1094-102	2
651	Enrichment of inflammatory bowel disease and colorectal cancer risk variants in colon expression quantitative trait loci. 2015 , 16, 138	29
650	Intestinal steroidogenesis controls PPARI expression in the colon and is impaired during ulcerative colitis. 2015 , 64, 901-10	31
649	Ulcerative Colitis. 2015 , 1573-1612	1
648	Assessment of the genetic basis of rosacea by genome-wide association study. 2015 , 135, 1548-1555	97
647	Can inflammatory bowel disease really be solved by the multiple -omics and meta-omics analyses?. 2015 , 165, 107-8	4

(2015-2015)

646	Genome wide identification of new genes and pathways in patients with both autoimmune thyroiditis and type 1 diabetes. 2015 , 60, 32-9	52
645	Association of IL-1R2 genetic polymorphisms with the susceptibility of ankylosing spondylitis in Northern Chinese Han population. 2015 , 25, 908-12	13
644	Mendelian Randomization: New Applications in the Coming Age of Hypothesis-Free Causality. 2015 , 16, 327-50	162
643	Human genomics. The Genotype-Tissue Expression (GTEx) pilot analysis: multitissue gene regulation in humans. 2015 , 348, 648-60	3242
642	A large-scale genetic analysis reveals a strong contribution of the HLA class II region to giant cell arteritis susceptibility. 2015 , 96, 565-80	96
641	IL12p40 regulates functional development of human CD4+ T cells: enlightenment by the elevated expressions of IL12p40 in patients with inflammatory bowel diseases. 2015 , 94, e613	5
640	Whole-genome sequencing to understand the genetic architecture of common gene expression and biomarker phenotypes. 2015 , 24, 1504-12	7
639	Role of protein tyrosine phosphatases in regulating the immune system: implications for chronic intestinal inflammation. 2015 , 21, 645-55	20
638	Type 3 innate lymphoid cells maintain intestinal epithelial stem cells after tissue damage. 2015 , 212, 1783-91	111
637	MicroRNAs enrichment in GWAS of complex human phenotypes. 2015 , 16, 304	19
636	Identification of a common variant with potential pleiotropic effect on risk of inflammatory bowel disease and colorectal cancer. 2015 , 36, 999-1007	21
635	Type I IFN induces protein ISGylation to enhance cytokine expression and augments colonic inflammation. 2015 , 112, 14313-8	30
634	Understanding inflammatory bowel disease via immunogenetics. 2015 , 64, 91-100	105
633	RIP3 Regulates Autophagy and Promotes Coxsackievirus B3 Infection of Intestinal Epithelial Cells. 2015 , 18, 221-32	38
632	Meta-analysis of shared genetic architecture across ten pediatric autoimmune diseases. 2015 , 21, 1018-27	143
631	The immunogenetics of Beh⊟t's disease: A comprehensive review. 2015 , 64, 137-48	131
630	Reduced sodium/proton exchanger NHE3 activity causes congenital sodium diarrhea. 2015 , 24, 6614-23	80
629	Pervasive pleiotropy between psychiatric disorders and immune disorders revealed by integrative analysis of multiple GWAS. 2015 , 134, 1195-209	57

628	Gallic acid suppresses inflammation in dextran sodium sulfate-induced colitis in mice: Possible mechanisms. <i>International Immunopharmacology</i> , 2015 , 28, 1034-43	94
627	Circulating microRNAs in Inflammatory Bowel Diseases. 2015 , 106, 197-214	2
626	MicroRNAs: new players in IBD. 2015 , 64, 504-17	174
625	Intestinal neuroendocrine cells and goblet cells are mediators of IL-17A-amplified epithelial IL-17C production in human inflammatory bowel disease. 2015 , 8, 943-58	43
624	The IRF5-TNPO3 association with systemic lupus erythematosus has two components that other autoimmune disorders variably share. 2015 , 24, 582-96	57
623	Interleukin-22 and CD160 play additive roles in the host mucosal response to Clostridium difficile infection in mice. 2015 , 144, 587-97	19
622	Leukocyte chemoattractant receptors in human disease pathogenesis. 2015, 10, 51-81	55
621	Pro- and anti-inflammatory cytokine gene single-nucleotide polymorphisms in inflammatory bowel disease. 2015 , 42, 38-45	21
620	Genetic and epigenetic fine mapping of causal autoimmune disease variants. 2015, 518, 337-43	1199
619	Genome-wide association scan in north Indians reveals three novel HLA-independent risk loci for ulcerative colitis. 2015 , 64, 571-9	42
618	Proteomic and genomic evidence implicates the postsynaptic density in schizophrenia. 2015 , 20, 424-32	105
617	Epigenomics of Intestinal Disease. 2016 , 257-273	
616	Application of computational methods in genetic study of inflammatory bowel disease. World Journal of Gastroenterology, 2016 , 22, 949-60	2
615	Emerging Role of Interleukin 22 in Hepatitis B Virus Infection: a Double-edged Sword. 2013 , 1, 103-8	9
614	Polymorphisms Are Associated with Disease Susceptibility and Phenotype in Japanese Patients with Inflammatory Bowel Disease. 2016 , 2016, 6485343	7
613	The Role of the C-Clamp in Wnt-Related Colorectal Cancers. 2016 , 8,	13
612	Functional CRISPR screening identifies the ufmylation pathway as a regulator of SQSTM1/p62. 2016 , 5,	93
611	Association Studies of the GPR103 and BCL2L15 Genes in Autoimmune Thyroid Disease in the Japanese Population. 2016 , 7, 92	9

(2016-2016)

610	Identification and Characterization of a Novel Association between Dietary Potassium and Risk of Crohn's Disease and Ulcerative Colitis. 2016 , 7, 554		28	
609	Validation of Type 2 Diabetes Risk Variants Identified by Genome-Wide Association Studies in Northern Han Chinese. 2016 , 13,		7	
608	A Genome-Wide Methylation Approach Identifies a New Hypermethylated Gene Panel in Ulcerative Colitis. <i>International Journal of Molecular Sciences</i> , 2016 , 17,	6.3	23	
607	Microbiome, Metabolome and Inflammatory Bowel Disease. 2016 , 4,		95	
606	Predicting gene targets from integrative analyses of summary data from GWAS and eQTL studies for 28 human complex traits. 2016 , 8, 84		59	
605	. 2016,		8	
604	Microbial Neuro-Immune Interactions and the Pathophysiology of IBD. 2016 ,		1	
603	Tight junctions in inflammatory bowel diseases and inflammatory bowel disease associated colorectal cancer. <i>World Journal of Gastroenterology</i> , 2016 , 22, 3117-26	5.6	224	
602	Genome-Wide Copy Number Variation Scan Identifies Complement Component C4 as Novel Susceptibility Gene for Crohn's Disease. 2016 , 22, 505-15		8	
601	Increased Expression of Interleukin-36, a Member of the Interleukin-1 Cytokine Family, in Inflammatory Bowel Disease. 2016 , 22, 303-14		78	
600	TNFRSF1B Is Associated with ANCA in IBD. 2016 , 22, 1346-52		4	
599	pANCA and ASCA in Children with IBD-Unclassified, Crohn's Colitis, and Ulcerative Colitis-A Longitudinal Report from the IBD Porto Group of ESPGHAN. 2016 , 22, 1908-14		28	
598	Pathway-based Genome-wide Association Studies Reveal the Association Between Growth Factor Activity and Inflammatory Bowel Disease. 2016 , 22, 1540-51		6	
597	Receptor occupancy and blocking of STAT5 signaling by an anti-IL-7 receptor hantibody in cynomolgus monkeys. 2016 , 90, 191-8		12	
596	Risk factors for primary sclerosing cholangitis. 2016 , 36, 84-91		15	
595	Microbiota at the crossroads of autoimmunity. 2016 , 15, 859-69		82	
594	Congenital Sodium Diarrhea: A Form of Intractable Diarrhea, With a Link to Inflammatory Bowel Disease. 2016 , 63, 170-6		36	
593	The molecular biology of matrix metalloproteinases and tissue inhibitors of metalloproteinases in inflammatory bowel diseases. 2016 , 51, 295-358		44	

592	Association between psoriasis and inflammatory bowel disease: a Danish nationwide cohort study. 2016 , 175, 487-92	83
591	Killer immunoglobulin-like receptor repertoire analysis in a Caucasian Spanish cohort with inflammatory bowel disease. 2016 , 60, 787-792	8
590	Care of Adults with Chronic Childhood Conditions. 2016,	4
589	Local genotype influences DNA methylation at two asthma-associated regions, 5q31 and 17q21, in a founder effect population. 2016 , 53, 232-41	14
588	The serine protease-mediated increase in intestinal epithelial barrier function is dependent on occludin and requires an intact tight junction. 2016 , 311, G466-79	17
587	Meta-analysis of associations between DLG5 R30Q and P1371Q polymorphisms and susceptibility to inflammatory bowel disease. 2016 , 6, 33550	4
586	Systematic analysis of chromatin interactions at disease associated loci links novel candidate genes to inflammatory bowel disease. 2016 , 17, 247	28
585	Comprehensive Assessment of the Association between FCGRs polymorphisms and the risk of systemic lupus erythematosus: Evidence from a Meta-Analysis. 2016 , 6, 31617	24
584	Expanding the Immunology Toolbox: Embracing Public-Data Reuse and Crowdsourcing. 2016 , 45, 1191-1204	13
583	HSPA6 is an ulcerative colitis susceptibility factor that is induced by cigarette smoke and protects intestinal epithelial cells by stabilizing anti-apoptotic Bcl-XL. 2016 , 1862, 788-796	12
582	Stem cell-based therapies in inflammatory bowel disease: promises and pitfalls. 2016 , 9, 533-47	26
581	The TNF Receptor Superfamily in Co-stimulating and Co-inhibitory Responses. 2016 , 44, 1005-19	202
580	MicroRNAs: how many in inflammatory bowel disease?. 2016 , 32, 258-66	33
579	Association Between Genetic Traits for Immune-Mediated Diseases and Alzheimer Disease. 2016 , 73, 691-7	100
578	Koch's postulates, microbial dysbiosis and inflammatory bowel disease. 2016 , 22, 594-9	28
577	Genome-wide Association Studies of Posttraumatic Stress Disorder in 2 Cohorts of US Army Soldiers. 2016 , 73, 695-704	114
576	Genetic variation in IBD: progress, clues to pathogenesis and possible clinical utility. 2016 , 12, 1091-107	56
575	Focusing the Spotlight on the Zebrafish Intestine to Illuminate Mechanisms of Colorectal Cancer. 2016 , 916, 411-37	8

574	Tofacitinib for the treatment of ulcerative colitis. 2016 , 25, 991-7	13
573	Ameliorating Active Ulcerative Colitis via an Orally Available Toll-Like Receptor-9 Modifier: A Prospective Open-Label, Multicenter Phase II Trial. 2016 , 61, 3246-3254	9
572	Targeted Resequencing and Functional Testing Identifies Low-Frequency Missense Variants in the Gene Encoding GARP as Significant Contributors to Atopic Dermatitis Risk. 2016 , 136, 2380-2386	23
57 ¹	Genetic dysbiosis. 2016 , 431-441	
570	Genetically engineered mouse models for studying inflammatory bowel disease. 2016 , 238, 205-19	30
569	The role of ORMDL proteins, guardians of cellular sphingolipids, in asthma. 2016 , 71, 918-30	24
568	EoE-like, EoE-light, or EoE-not? Pursuing variations of a new disease. 2016 , 71, 739-40	
567	Population-specific genome-wide mapping of expression quantitative trait loci in the colon of Han Chinese. 2016 , 17, 600-609	6
566	Review article: the pathogenesis of pouchitis. 2016 , 44, 817-35	53
565	Identification of a human intestinal myeloid cell subset that regulates gut homeostasis. 2016 , 28, 533-545	17
564	Gene and Network Analysis of Common Variants Reveals Novel Associations in Multiple Complex Diseases. 2016 , 204, 783-798	32
563	The 100 classic papers in ulcerative colitis: a bibliometric analysis. 2016 , 10, 1187-1195	7
562	Cardiometabolic risk loci share downstream cis- and trans-gene regulation across tissues and diseases. 2016 , 353, 827-30	166
561	The Polymorphism rs3024505 (C/T) Downstream of the IL10 Gene Is Associated with Crohn's Disease in Serbian Patients with Inflammatory Bowel Disease. 2016 , 240, 15-24	11
560	Classic IL-6R signalling is dispensable for intestinal epithelial proliferation and repair. 2016 , 5, e270	21
559	A protein-truncating R179X variant in RNF186 confers protection against ulcerative colitis. 2016 , 7, 12342	41
558	Tumour necrosis factor-alpha (-308G/A) promoter polymorphism is associated with ulcerative colitis in Brazilian patients. 2016 , 43, 376-382	5
557	Paneth cells: the hub for sensing and regulating intestinal flora. 2016 , 59, 463-7	13

556	Systematic review with meta-analysis: the effect of tobacco smoking on the natural history of ulcerative colitis. 2016 , 44, 117-26	42
555	A meta-analysis of the relationship between MYO9B gene polymorphisms and susceptibility to Crohn's disease and ulcerative colitis. 2016 , 77, 990-996	9
554	Roles of the Epithelial Autophagy in the Intestinal Mucosal Barrier. 2016 , 603-616	1
553	Lineage-Specific Genome Architecture Links Enhancers and Non-coding Disease Variants to Target Gene Promoters. 2016 , 167, 1369-1384.e19	556
552	17q21 asthma-risk variants switch CTCF binding and regulate IL-2 production by T cells. 2016 , 7, 13426	62
551	Genetic Drivers of Epigenetic and Transcriptional Variation in Human Immune Cells. 2016 , 167, 1398-1414.e24	339
550	Paternal chronic colitis causes epigenetic inheritance of susceptibility to colitis. 2016 , 6, 31640	12
549	An Ancient Fecundability-Associated Polymorphism Switches a Repressor into an Enhancer of Endometrial TAP2 Expression. 2016 , 99, 1059-1071	10
548	Human Intestinal Barrier Function in Health and Disease. 2016 , 7, e196	396
547	Increased Prevalence of Inflammatory Bowel Disease in Patients with Mutations in Genes Encoding the Receptor Subunits for TGF[] 2016 , 22, 2058-2062	8
546	Explaining the disease phenotype of intergenic SNP through predicted long range regulation. 2016 , 44, 8641-8654	26
545	Caenorhabditis elegans susceptibility to gut Enterococcus faecalis infection is associated with fat metabolism and epithelial junction integrity. 2016 , 16, 6	24
544	Genome-wide rare copy number variation screening in ulcerative colitis identifies potential susceptibility loci. 2016 , 17, 26	10
543	Mapping hyper-susceptibility to colitis-associated colorectal cancer in FVB/NJ mice. 2016 , 27, 213-24	
542	Genome-Wide Association Studies Suggest Limited Immune Gene Enrichment in Schizophrenia Compared to 5 Autoimmune Diseases. 2016 , 42, 1176-84	42
54 ¹	NUR77 exerts a protective effect against inflammatory bowel disease by negatively regulating the TRAF6/TLR-IL-1R signalling axis. 2016 , 238, 457-69	46
540	Whole-exome Sequence Analysis Implicates Rare Il17REL Variants in Familial and Sporadic Inflammatory Bowel Disease. 2016 , 22, 20-7	12
539	Impact of a multidisciplinary approach in enteropathic spondyloarthritis patients. 2016 , 15, 184-90	30

538	Reparative inflammation takes charge of tissue regeneration. 2016 , 529, 307-15	411
537	Identification of critical variants within SLC44A4, an ulcerative colitis susceptibility gene identified in a GWAS in north Indians. 2016 , 17, 105-9	9
536	IBD Genetic Risk Profile in Healthy First-Degree Relatives of Crohn's Disease Patients. 2016 , 10, 209-15	22
535	Primary Immunodeficiencies and Inflammatory Disease: A Growing Genetic Intersection. 2016 , 37, 126-140	39
534	Genetic characteristics of inflammatory bowel disease in a Japanese population. 2016 , 51, 672-81	38
533	Risk of Inflammatory Bowel Disease with Oral Contraceptives and Menopausal Hormone Therapy: Current Evidence and Future Directions. 2016 , 39, 193-7	21
532	Clinical Features and HLA Association of 5-Aminosalicylate (5-ASA)-induced Nephrotoxicity in Inflammatory Bowel Disease. 2016 , 10, 149-58	57
531	NOD2 gene variant is a risk factor for postoperative complications in patients with Crohn's disease: A genetic association study. 2016 , 160, 74-80	11
530	IBD and Environment: Are There Differences between East and West. 2016 , 34, 84-9	26
529	eQTL analysis links inflammatory bowel disease associated 1q21 locus to ECM1 gene. 2016 , 57, 363-72	7
529 528	eQTL analysis links inflammatory bowel disease associated 1q21 locus to ECM1 gene. 2016 , 57, 363-72 Genetics and Pathogenesis of Inflammatory Bowel Disease. 2016 , 11, 127-48	7
528	Genetics and Pathogenesis of Inflammatory Bowel Disease. 2016 , 11, 127-48 Analysis of five chronic inflammatory diseases identifies 27 new associations and highlights	149
528 527	Genetics and Pathogenesis of Inflammatory Bowel Disease. 2016, 11, 127-48 Analysis of five chronic inflammatory diseases identifies 27 new associations and highlights disease-specific patterns at shared loci. <i>Nature Genetics</i> , 2016, 48, 510-8 Tissue-specific regulatory circuits reveal variable modular perturbations across complex diseases.	149 404
528 527 526	Genetics and Pathogenesis of Inflammatory Bowel Disease. 2016, 11, 127-48 Analysis of five chronic inflammatory diseases identifies 27 new associations and highlights disease-specific patterns at shared loci. <i>Nature Genetics</i> , 2016, 48, 510-8 Tissue-specific regulatory circuits reveal variable modular perturbations across complex diseases. 2016, 13, 366-70	149 404
528 527 526 525	Genetics and Pathogenesis of Inflammatory Bowel Disease. 2016, 11, 127-48 Analysis of five chronic inflammatory diseases identifies 27 new associations and highlights disease-specific patterns at shared loci. <i>Nature Genetics</i> , 2016, 48, 510-8 Tissue-specific regulatory circuits reveal variable modular perturbations across complex diseases. 2016, 13, 366-70 Colitis ulcerosa. 2016, 12, 227-241	149 404 201
528 527 526 525 524	Genetics and Pathogenesis of Inflammatory Bowel Disease. 2016, 11, 127-48 Analysis of five chronic inflammatory diseases identifies 27 new associations and highlights disease-specific patterns at shared loci. Nature Genetics, 2016, 48, 510-8 Tissue-specific regulatory circuits reveal variable modular perturbations across complex diseases. 2016, 13, 366-70 Colitis ulcerosa. 2016, 12, 227-241 Gut inflammation and microbiome in spondyloarthritis. 2016, 36, 457-68	149 404 201

520	Ocular Manifestations in Inflammatory Bowel Disease Are Associated with Other Extra-intestinal Manifestations, Gender, and Genes Implicated in Other Immune-related Traits. 2016 , 10, 43-9		27
519	Tissue-infiltrating neutrophils represent the main source of IL-23 in the colon of patients with IBD. 2016 , 65, 1632-41		62
518	Interferon regulatory factor 5 in human autoimmunity and murine models of autoimmune disease. 2016 , 167, 167-82		45
517	Siblings of patients with Crohn's disease exhibit a biologically relevant dysbiosis in mucosal microbial metacommunities. 2016 , 65, 944-53		49
516	Association analysis of copy numbers of FC-gamma receptor genes for rheumatoid arthritis and other immune-mediated phenotypes. 2016 , 24, 263-70		14
515	The gut microbiota and inflammatory bowel diseases. 2017 , 179, 38-48		84
514	Evidence for three genetic loci involved in both anorexia nervosa risk and variation of body mass index. 2017 , 22, 192-201		31
513	Redox signaling in the gastrointestinal tract. 2017 , 104, 75-103		136
512	Genome-wide association study implicates immune activation of multiple integrin genes in inflammatory bowel disease. <i>Nature Genetics</i> , 2017 , 49, 256-261	36.3	462
511	Exploring the genetic architecture of inflammatory bowel disease by whole-genome sequencing identifies association at ADCY7. <i>Nature Genetics</i> , 2017 , 49, 186-192	36.3	104
510	Inhibition of matrix metalloproteinase-9 by a barbiturate-nitrate hybrid ameliorates dextran sulphate sodium-induced colitis: effect on inflammation-related genes. 2017 , 174, 512-524		10
509	Genetics of inflammatory bowel disease: beyond NOD2. 2017 , 2, 224-234		68
508	Analysis of the common genetic component of large-vessel vasculitides through a meta-Immunochip strategy. 2017 , 7, 43953		34
507	De novo and rare mutations in the HSPA1L heat shock gene associated with inflammatory bowel disease. 2017 , 9, 8		22
506	Modeling intestinal disorders using zebrafish. 2017 , 138, 241-270		29
505	Single nucleotide polymorphisms of IL12B are associated with Takayasu arteritis in Chinese Han population. 2017 , 37, 547-555		6
504	Opportunities and challenges of whole-genome and -exome sequencing. 2017 , 18, 14		110
503	Genetic Characterization and Influence on Inflammatory Bowel Disease Expression in a Diverse Hispanic South Florida Cohort. 2017 , 8, e87		8

502	Type 1 Interferon in the Human Intestine-A Co-ordinator of the Immune Response to the Microbiota. 2017 , 23, 524-533	17
501	Third European Evidence-based Consensus on Diagnosis and Management of Ulcerative Colitis. Part 1: Definitions, Diagnosis, Extra-intestinal Manifestations, Pregnancy, Cancer Surveillance, Surgery, and Ileo-anal Pouch Disorders. 2017 , 11, 649-670	837
500	History of Human Genetics. 2017,	5
499	A rising incidence and poorer male outcomes characterise early onset paediatric inflammatory bowel disease. 2017 , 45, 1534-1541	19
498	The many ways tissue phagocytes respond to dying cells. 2017 , 277, 158-173	46
497	Endoscopic features and genetic background of inflammatory bowel disease complicated with Takayasu arteritis. 2017 , 32, 1011-1017	10
496	GWAB: a web server for the network-based boosting of human genome-wide association data. 2017 , 45, W154-W161	19
495	Genetics of Infectious and Inflammatory Diseases: Overlapping Discoveries from Association and Exome-Sequencing Studies. 2017 , 35, 1-30	28
494	Genome-wide Pleiotropy Between Parkinson Disease and Autoimmune Diseases. 2017 , 74, 780-792	150
493	Roles of intestinal epithelial cells in the maintenance of gut homeostasis. 2017 , 49, e338	2 60
492	Zinc treatment is efficient against Escherichia coli Haemolysin-induced intestinal leakage in mice. 2017 , 7, 45649	19
491	Human Epistatic Interaction Controls IL7R Splicing and Increases Multiple Sclerosis Risk. 2017 , 169, 72-84.e13	55
490	Rip2 Is Required for Nod2-Mediated Lysozyme Sorting in Paneth Cells. 2017 , 198, 3729-3736	22
489	Niacin ameliorates ulcerative colitis via prostaglandin D-mediated D prostanoid receptor 1 activation. 2017 , 9, 571-588	32
488	Runx3 in Immunity, Inflammation and Cancer. 2017, 962, 369-393	21
487	Modeling month-season of birth as a risk factor in mouse models of chronic disease: from multiple sclerosis to autoimmune encephalomyelitis. 2017 , 31, 2709-2719	7
486	Transethnic meta-analysis identifies and as susceptibility genes to systemic sclerosis. 2017 , 76, 1150-1158	49
485	Genome-wide Association Study Identifies 27 Loci Influencing Concentrations of Circulating Cytokines and Growth Factors. 2017 , 100, 40-50	133

484	Atg16 promotes enteroendocrine cell differentiation via regulation of intestinal Slit/Robo signaling. 2017 , 144, 3990-4001	19
483	Small-molecule inhibitors directly target CARD9 and mimic its protective variant in inflammatory bowel disease. 2017 , 114, 11392-11397	28
482	LRRK2 promotes the activation of NLRC4 inflammasome during Typhimurium infection. 2017, 214, 3051-3066	83
481	The contribution of long non-coding RNAs in Inflammatory Bowel Diseases. 2017 , 49, 1067-1072	35
480	Fourteen sequence variants that associate with multiple sclerosis discovered by meta-analysis informed by genetic correlations. 2017 , 2, 24	8
479	Development of autoantibodies precedes clinical manifestations of autoimmune diseases: A comprehensive review. 2017 , 83, 95-112	81
478	The genetics of Takayasu arteritis. 2017 , 46, e179-e187	30
477	Evaluating the Association of Common Variants of the SLC44A4 Gene with Ulcerative Colitis Susceptibility in the Han Chinese Population. 2017 , 21, 555-559	3
476	Parkinson disease-associated transgene disrupts marrow myelopoiesis and peripheral Th17 response. 2017 , 102, 1093-1102	17
475	Proteomic analysis of ascending colon biopsies from a paediatric inflammatory bowel disease inception cohort identifies protein biomarkers that differentiate Crohn's disease from UC. 2017 , 66, 1573-158	3 ⁵⁰
474	Genome-wide association study of classical Hodgkin lymphoma identifies key regulators of disease susceptibility. 2017 , 8, 1892	24
473	Cohort profile: design and first results of the Dutch IBD Biobank: a prospective, nationwide biobank of patients with inflammatory bowel disease. 2017 , 7, e016695	20
472	Peripartum Antibiotics Promote Gut Dysbiosis, Loss of Immune Tolerance, and Inflammatory Bowel Disease in Genetically Prone Offspring. 2017 , 20, 491-504	69
471	Genetik des Morbus Crohn und der Colitis ulcerosa. 2017 , 12, 38-48	4
470	A novel approach to genome-wide association analysis identifies genetic associations with primary biliary cholangitis and primary sclerosing cholangitis in Polish patients. 2017 , 10, 2	10
469	E3 Ubiquitin ligase ZNRF4 negatively regulates NOD2 signalling and induces tolerance to MDP. 2017 , 8, 15865	12
468	SUMOylation pathway alteration coupled with downregulation of SUMO E2 enzyme at mucosal epithelium modulates inflammation in inflammatory bowel disease. 2017 , 7,	16
467	Gleaning Insights from Fecal Microbiota Transplantation and Probiotic Studies for the Rational Design of Combination Microbial Therapies. 2017 , 30, 191-231	41

466	A cross-ethnic survey of CFB and SLC44A4, Indian ulcerative colitis GWAS hits, underscores their potential role in disease susceptibility. 2016 , 25, 111-122	5
465	The genetic background of inflammatory bowel disease: from correlation to causality. 2017 , 241, 146-158	63
464	Disruption of the Hedgehog signaling pathway in inflammatory bowel disease fosters chronic intestinal inflammation. 2017 , 17, 351-369	7
463	Insight into the role of TSLP in inflammatory bowel diseases. 2017 , 16, 55-63	26
462	Inflammatory Bowel Disease: A Global Disease That Needs a Broader Ensemble of Populations. 2017 , 152, 14-16	6
461	Integrating embeddings of multiple gene networks to prioritize complex disease-associated genes. 2017 ,	5
460	Single-Nucleotide Polymorphisms and Inflammation. 2017 , 1329-1346	
459	Simultaneous inference of phenotype-associated genes and relevant tissues from GWAS data via Bayesian integration of multiple tissue-specific gene networks. 2017 , 9, 436-452	9
458	Chromosome 17q21 Genes ORMDL3 and GSDMB in Asthma and Immune Diseases. 2017 , 135, 1-52	61
457	The new NHGRI-EBI Catalog of published genome-wide association studies (GWAS Catalog). 2017 , 45, D896-D901	1321
456	Chitin protects the gut epithelial barrier in a protochordate model of DSS-induced colitis. 2018, 7,	8
455	Lack of Adrenomedullin Aggravates Acute TNBS-Induced Colitis Symptoms in Mice, Especially in Females. 2017 , 8, 1058	7
454	Identification of candidate protective variants for common diseases and evaluation of their protective potential. 2017 , 18, 575	11
453	Diet, Gut Microbiome and Epigenetics: Emerging Links with Inflammatory Bowel Diseases and Prospects for Management and Prevention. 2017 , 9,	75
452	A Potential Role of Infection in the Onset of Inflammatory Bowel Diseases. 2017, 8, 191	34
451	Interactions between Intestinal Microbiota and Host Immune Response in Inflammatory Bowel Disease. 2017 , 8, 942	147
450	The Expression of the Short Isoform of Thymic Stromal Lymphopoietin in the Colon Is Regulated by the Nuclear Receptor Peroxisome Proliferator Activated Receptor-Gamma and Is Impaired during Ulcerative Colitis. 2017 , 8, 1052	8
449	Geography, Ethnicity or Subsistence-Specific Variations in Human Microbiome Composition and Diversity. 2017 , 8, 1162	400

448	Pathomechanisms of Oxidative Stress in Inflammatory Bowel Disease and Potential Antioxidant Therapies. 2017 , 2017, 4535194	223
447	Interleukin-7 and Immunosenescence. 2017 , 2017, 4807853	39
446	HLA and non-HLA genes and familial predisposition to autoimmune diseases in families with a child affected by type 1 diabetes. 2017 , 12, e0188402	11
445	graph-GPA: A graphical model for prioritizing GWAS results and investigating pleiotropic architecture. 2017 , 13, e1005388	8
444	p53 expression in patients with ulcerative colitis - associated with dysplasia and carcinoma: a systematic meta-analysis. 2017 , 17, 111	13
443	Performance of risk prediction for inflammatory bowel disease based on genotyping platform and genomic risk score method. 2017 , 18, 94	20
442	Genome-wide association studies in Crohn's disease: Past, present and future. 2018 , 7, e1001	45
441	Crohn's Disease Candidate Gene Alleles Predict Time to Progression from Inflammatory B1 to Stricturing B2, or Penetrating B3 Phenotype. 2018 , 22, 143-151	8
440	IL-1R2 deficiency suppresses dextran sodium sulfate-induced colitis in mice via regulation of microbiota. 2018 , 496, 934-940	6
439	is a colitis risk gene that regulates stability of epithelial adherens junctions. 2018 , 359, 1161-1166	65
438	Changes in macrophage transcriptome associate with systemic sclerosis and mediate contribution to disease risk. 2018 , 77, 596-601	35
437	Genetic Predisposition to Rosacea. 2018 , 36, 87-92	14
436	Modulatory activities of Chrysophyllum albidum and its fractions on microflora and colonic pump activities during inflammatory phase of colitis healing in experimental mice. 2018 , 22, 38-49	2
435	Genetic profile of patients with early onset inflammatory bowel disease. 2018 , 645, 18-29	19
434	A decade of research on the 17q12-21 asthma locus: Piecing together the puzzle. 2018, 142, 749-764.e3	90
433	Genetic Variation and Gene Expression Levels of Tight Junction Genes Indicates Relationships Between PTEN as well as MAGI1 and Microscopic Colitis. 2018 , 63, 105-112	11
432	Maintenance of intestinal homeostasis by mucosal barriers. 2018 , 38, 5	126
431	The Emerging Immunogenetic Architecture of Schizophrenia. 2018 , 44, 993-1004	31

430	The use of tofacitinib in the treatment of inflammatory bowel disease. 2018 , 10, 837-849	4
429	Evidence for association of STAT4 and IL12RB2 variants with Myasthenia gravis susceptibility: What is the effect on gene expression in thymus?. 2018 , 319, 93-99	3
428	Ameliorating effect of TI-1-162, a hydroxyindenone derivative, against TNBS-induced rat colitis is mediated through suppression of RIP/ASK-1/MAPK signaling. 2018 , 827, 94-102	8
427	Immunopathogenesis of inflammatory bowel disease and mechanisms of biological therapies. 2018 , 53, 379-389	82
426	The role of IL-12/23 in T cell-related chronic inflammation: implications of immunodeficiency and therapeutic blockade. 2018 , 57, 246-254	20
425	Transcriptomic Landscape of Treatment-Nalle Ulcerative Colitis. 2018 , 12, 327-336	31
424	Evidence-based update on rosacea comorbidities and their common physiologic pathways. 2018 , 78, 156-166	48
423	The role of the intestinal microbiota in the pathogenesis and treatment of inflammatory bowel diseases. 2018 , 29, 21-27	
422	Pathogenesis of Periodontal Diseases. 2018,	3
421	Bayesian analysis of genome-wide inflammatory bowel disease data sets reveals new risk loci. 2018 , 26, 265-274	12
420	A meta-analysis of coastal wetland ecosystem services in Liaoning Province, China. 2018 , 200, 349-358	18
419	Genetics of ulcerative colitis: putting into perspective the incremental gains from Indian studies. 2018 , 97, 1493-1507	2
418	Recent Advances in Probiotics as Live Biotherapeutics Against Gastrointestinal Diseases. 2018 , 24, 3162-3171	12
417	ATF3 Sustains IL-22-Induced STAT3 Phosphorylation to Maintain Mucosal Immunity Through Inhibiting Phosphatases. 2018 , 9, 2522	19
416	Differential Intestinal Mucosa Transcriptomic Biomarkers for Crohn's Disease and Ulcerative Colitis. 2018 , 2018, 9208274	16
415	Modulation of faecal metagenome in Crohn's disease: Role of microRNAs as biomarkers. <i>World Journal of Gastroenterology</i> , 2018 , 24, 5223-5233	19
414	Dynamics of Colon Monocyte and Macrophage Activation During Colitis. 2018, 9, 2764	63
413	Uncovering new disease indications for G-protein coupled receptors and their endogenous ligands. 2018 , 19, 345	4

412	Inflammatory Bowel Disease: What Very Early Onset Disease Teaches Us. 2018, 47, 755-772	25
411	Genome-Wide Association Study Reveals Genetic Link between Diarrhea-Associated Entamoeba histolytica Infection and Inflammatory Bowel Disease. 2018 , 9,	16
410	An Ancient Fecundability-Associated Polymorphism Creates a GATA2 Binding Site in a Distal Enhancer of HLA-F. 2018 , 103, 509-521	13
409	IL-7 receptor blockade blunts antigen-specific memory T cell responses and chronic inflammation in primates. 2018 , 9, 4483	26
408	Single-Nucleotide Polymorphism of the MLX Gene Is Associated With Takayasu Arteritis. 2018 , 11, e002296	7
407	Genetic and transcriptional analysis of inflammatory bowel disease-associated pathways in patients with GUCY2C-linked familial diarrhea. 2018 , 53, 1264-1273	4
406	Should we target TNF receptors in the intestinal epithelium with glucocorticoids during systemic inflammation?. 2018 , 22, 1029-1037	3
405	Genetic correlations among psychiatric and immune-related phenotypes based on genome-wide association data. 2018 , 177, 641-657	75
404	Vitamin D and Inflammatory Bowel Disease: Mendelian Randomization Analyses in the Copenhagen Studies and UK Biobank. 2018 , 103, 3267-3277	14
403	Evidence for PTGER4, PSCA, and MBOAT7 as risk genes for gastric cancer on the genome and transcriptome level. 2018 , 7, 5057-5065	16
402	Microbiota-derived short-chain fatty acids promote Th1 cell IL-10 production to maintain intestinal homeostasis. 2018 , 9, 3555	199
401	Rare coding variant analysis in a large cohort of Ashkenazi Jewish families with inflammatory bowel disease. 2018 , 137, 723-734	4
400	The Genetics and Genomics of Asthma. 2018 , 19, 223-246	20
399	Viewpoint: Toward the Genetic Architecture of Disease Severity in Inflammatory Bowel Diseases. 2018 , 24, 1428-1439	3
398	Common genetic variation in the autoimmune regulator (AIRE) locus is associated with autoimmune Addison's disease in Sweden. 2018 , 8, 8395	14
397	Interferon regulatory factor 5 and nuclear factor kappa-B exhibit cooperating but also divergent roles in the regulation of pro-inflammatory cytokines important for the development of TH1 and TH17 responses. 2018 , 285, 3097-3113	14
396	Mucosal Restitution and Repair. 2018, 683-708	
395	Association of Long Noncoding RNAs Polymorphisms With Ankylosing Spondylitis, Vogt-Koyanagi-Harada Disease, and Behcet's Disease. 2018 , 59, 1158-1166	9

(2018-2018)

394	Genes regulated by SATB2 during neurodevelopment contribute to schizophrenia and educational attainment. 2018 , 14, e1007515	17
393	Nitric Oxide Engages an Anti-inflammatory Feedback Loop Mediated by Peroxiredoxin 5 in Phagocytes. 2018 , 24, 838-850	21
392	Leucine-rich repeat kinase 2 aggravates secondary brain injury induced by intracerebral hemorrhage in rats by regulating the P38 MAPK/Drosha pathway. 2018 , 119, 53-64	17
391	Enhancer histone-QTLs are enriched on autoimmune risk haplotypes and influence gene expression within chromatin networks. 2018 , 9, 2905	36
390	Integrative Analysis Identifies Genetic Variants Associated With Autoimmune Diseases Affecting Putative MicroRNA Binding Sites. 2018 , 9, 139	11
389	Donor Genotype in the Interleukin-7 Receptor Echain Predicts Risk of Graft-versus-Host Disease and Cytomegalovirus Infection after Allogeneic Hematopoietic Stem Cell Transplantation. 2018 , 9, 109	8
388	Effector T Helper Cell Subsets in Inflammatory Bowel Diseases. 2018 , 9, 1212	97
387	Promoter methylation of the and genes correlates with the composition of the immunoglobulin G glycome in inflammatory bowel disease. 2018 , 10, 75	17
386	Revisiting IL-2: Biology and therapeutic prospects. 2018 , 3,	200
385	An integrative network-based approach to identify novel disease genes and pathways: a case study in the context of inflammatory bowel disease. 2018 , 19, 264	15
384	Systems Biology in Immunotoxicology. 2018 , 559-581	1
383	Meta-analysis of GWAS on both Chinese and European populations identifies GPR173 as a novel X chromosome susceptibility gene for SLE. 2018 , 20, 92	12
382	Genetik des Morbus Crohn und der Colitis ulcerosa. 2018 , 21, 4-13	
381	Deep Resequencing of Ulcerative Colitis-Associated Genes Identifies Novel Variants in Candidate Genes in the Korean Population. 2018 , 24, 1706-1717	4
380	Pathophysiology of IBD associated diarrhea. 2018 , 6, e1463897	48
379	Genome-wide DNA Methylation in Treatment-nalle Ulcerative Colitis. 2018 , 12, 1338-1347	16
378	Evidence for a potential role of miR-1908-5p and miR-3614-5p in autoimmune disease risk using integrative bioinformatics. 2018 , 94, 83-89	9
377	Histopathological Image QTL Discovery of Immune Infiltration Variants. 2018 , 5, 80-89	8

376	Contribution of STAT3 to Inflammatory and Fibrotic Diseases and Prospects for its Targeting for Treatment. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	60
375	Leveraging multiple gene networks to prioritize GWAS candidate genes via network representation learning. 2018 , 145, 41-50	7
374	Genetic variants in cellular transport do not affect mesalamine response in ulcerative colitis. 2018 , 13, e0192806	5
373	Immune-related genetic enrichment in frontotemporal dementia: An analysis of genome-wide association studies. 2018 , 15, e1002487	77
372	Identification of six novel susceptibility loci for dyslipidemia using longitudinal exome-wide association studies in a Japanese population. 2019 , 111, 520-533	2
371	Allergic and Immunologic Perspectives of Inflammatory Bowel Disease. 2019 , 57, 179-193	17
370	T cell-intrinsic prostaglandin E-EP2/EP4 signaling is critical in pathogenic T17 cell-driven inflammation. 2019 , 143, 631-643	43
369	Association of T Helper 1 Cytokine Gene Single Nucleotide Polymorphisms with Ulcerative Colitis and Crohn's Disease. 2019 , 37, 21-32	1
368	N-Terminomics/TAILS Profiling of Proteases and Their Substrates in Ulcerative Colitis. 2019 , 14, 2471-2483	11
367	Rare and common variant discovery in complex disease: the IBD case study. 2019 , 28, R162-R169	5
366	Inflammatory cytokines: from discoveries to therapies in IBD. 2019 , 19, 1207-1217	47
365	Role of DNA Methylation in the Development and Differentiation of Intestinal Epithelial Cells and Smooth Muscle Cells. 2019 , 25, 377-386	6
364	Interleukin-18 as a drug repositioning opportunity for inflammatory bowel disease: A Mendelian randomization study. 2019 , 9, 9386	11
363	Differential genetic and functional background in inflammatory bowel disease phenotypes of a Greek population: a systems bioinformatics approach. 2019 , 11, 31	5
362	Inflammatory Bowel Disease. 2019 , 373-424	
361	IBD BioResource: an open-access platform of 25 000 patients to accelerate research in Crohn's and Colitis. 2019 , 68, 1537-1540	9
360	Concordance between gene expression in peripheral whole blood and colonic tissue in children with inflammatory bowel disease. 2019 , 14, e0222952	9
359	Functions and regulation of T cell-derived interleukin-10. 2019 , 44, 101344	48

(2019-2019)

358	Inflammatory Bowel Disease (IBD)-A Textbook Case for Multi-Centric Banking of Human Biological Materials. 2019 , 6, 230	1
357	Proteomics-based functional studies reveal that galectin-3 plays a protective role in the pathogenesis of intestinal Behat's disease. 2019 , 9, 11716	5
356	Sustained suppression of IL-18 by employing a vaccine ameliorates intestinal inflammation in TNBS-induced murine colitis. 2019 , 5, FSO405	2
355	Loss of PTPN22 abrogates the beneficial effect of cohousing-mediated fecal microbiota transfer in murine colitis. 2019 , 12, 1336-1347	9
354	Protective Effect of Extract against Dextran-Sulfate-Sodium-Induced Ulcerative Colitis in Rats. 2019 , 11,	13
353	TFF3 interacts with LINGO2 to regulate EGFR activation for protection against colitis and gastrointestinal helminths. 2019 , 10, 4408	27
352	Metabolic Functions of Gut Microbes Associate With Efficacy of Tumor Necrosis Factor Antagonists in Patients With Inflammatory Bowel Diseases. 2019 , 157, 1279-1292.e11	101
351	Functional SNPs in the Human Autoimmunity-Associated Locus 17q12-21. 2019 , 10,	4
350	Non-coding DNA in IBD: from sequence variation in DNA regulatory elements to novel therapeutic potential. 2019 , 68, 928-941	11
349	Degradation of the extracellular matrix is part of the pathology of ulcerative colitis. 2019, 15, 67-76	6
348	Roles of Autophagy-Related Genes in the Pathogenesis of Inflammatory Bowel Disease. 2019, 8,	39
347	Evaluating tofacitinib citrate in the treatment of moderate-to-severe active ulcerative colitis: design, development and positioning of therapy. 2019 , 12, 179-191	11
346	Genetic and Epigenetic Fine Mapping of Complex Trait Associated Loci in the Human Liver. 2019 , 105, 89-107	20
345	Integrative Analysis of Transcriptomic and Proteomic Profiling in Inflammatory Bowel Disease Colon Biopsies. 2019 , 25, 1906-1918	11
344	Common Genetic Component in Autoimmunity. 2019 , 221-236	
343	Genetics of Rare Autoimmune Diseases. 2019,	
342	Rab GTPases as Physiological Substrates of LRRK2 Kinase. 2019 , 28, 134-145	14
341	Genetic and Genomic Markers for Prognostication. 2019 , 323-331	

340	Genetic Studies of Inflammatory Bowel Disease-Focusing on Asian Patients. 2019, 8,		25
339	Common functional alterations identified in blood transcriptome of autoimmune cholestatic liver and inflammatory bowel diseases. 2019 , 9, 7190		10
338	Effective Use of the Laboratory in the Management of Patients with Inflammatory Bowel Diseases. 2019 , 48, 237-258		O
337	PRDM1 levels are associated with clinical diseases in chronic HBV infection and survival of patients with HBV-related hepatocellular carcinoma. <i>International Immunopharmacology</i> , 2019 , 73, 156-162	8	7
336	Understanding the importance of autophagy in human diseases using Drosophila. 2019, 46, 157-169		12
335	An update on the interactions between Alzheimer's disease, autophagy and inflammation. 2019 , 705, 157-166		38
334	Genetic Factors and the Intestinal Microbiome Guide Development of Microbe-Based Therapies for Inflammatory Bowel Diseases. 2019 , 156, 2174-2189		85
333	Dysregulation of Intestinal Epithelial Cell RIPK Pathways Promotes Chronic Inflammation in the IBD Gut. 2019 , 10, 1094		27
332	IgG and Fc[Receptors in Intestinal Immunity and Inflammation. 2019, 10, 805		37
331	Prediction of steroid demand in the treatment of patients with ulcerative colitis by immunohistochemical analysis of the mucosal microenvironment and immune checkpoint: role of macrophages and regulatory markers in disease severity. 2019 , 69, 260-271		6
330	Multiple Epistasis Interactions Within MHC Are Associated With Ulcerative Colitis. 2019, 10, 257		6
329	Intestinal Organoids as a Novel Complementary Model to Dissect Inflammatory Bowel Disease. 2019 , 2019, 8010645		6
328	Ulcerative Colitis: Shifting Sands. 2019 , 19, 227-234		9
327	Contribution of Zinc and Zinc Transporters in the Pathogenesis of Inflammatory Bowel Diseases. 2019 , 2019, 8396878		20
326	The Synergistic Role of Diet and Exercise in the Prevention, Pathogenesis, and Management of Ulcerative Colitis: An Underlying Metabolic Mechanism. 2019 , 12, 1178638819834526		4
325	Autoimmune genetic risk variants as germline biomarkers of response to melanoma immune-checkpoint inhibition. 2019 , 68, 897-905		21
324	Molecular mechanisms underlying prostaglandin E2-exacerbated inflammation and immune diseases. 2019 , 31, 597-606		73
323	Impact of red meat, processed meat and fibre intake on risk of late-onset chronic inflammatory diseases: prospective cohort study on lifestyle factors using the Danish 'Diet, Cancer and Health' cohort (PROCID-DCH):. 2019 , 9, e024555		5

322 Very Early Onset Inflammatory Bowel Disease (VEOIBD). **2019**, 383-404

321	Colonic epithelial cell diversity in health and inflammatory bowel disease. 2019 , 567, 49-55	213
320	Inflammatory bowel disease - one entity with many molecular faces. 2019 , 14, 228-232	1
319	Flip the coin: IL-7 and IL-7R in health and disease. 2019 , 20, 1584-1593	83
318	Association of rosacea with inflammatory bowel disease: A MOOSE-compliant meta-analysis. 2019 , 98, e16448	6
317	Vascular endothelial growth factor receptor 1 tyrosine kinase signaling facilitates healing of DSS-induced colitis by accumulation of Tregs in ulcer area. 2019 , 111, 131-141	8
316	Moderate-to-severe asthma in individuals of European ancestry: a genome-wide association study. 2019 , 7, 20-34	109
315	Protective Mechanisms of Butyrate on Inflammatory Bowel Disease. 2018 , 24, 4154-4166	50
314	Identification of novel susceptibility genes associated with seven autoimmune disorders using whole genome molecular interaction networks. 2019 , 97, 48-58	7
313	Integrated serum proteins and fatty acids analysis for putative biomarker discovery in inflammatory bowel disease. 2019 , 195, 138-149	15
312	Peripheral disease contributes significantly to the level of disease activity in axial spondyloarthritis. 2019 , 5, e000802	20
311	Application of Proteomics to Inflammatory Bowel Disease Research: Current Status and Future Perspectives. 2019 , 2019, 1426954	9
310	Genetic variants of SMAD2/3/4/7 are associated with susceptibility to ulcerative colitis in a Japanese genetic background. 2019 , 207, 64-72	9
309	Big data in IBD: a look into the future. 2019 , 16, 312-321	58
308	The role of obesity in inflammatory bowel disease. 2019 , 1865, 63-72	24
307	Emergent Behavior of IBD-Associated Escherichia coli During Disease. 2019 , 25, 33-44	6
306	Comorbid immune-mediated diseases in inflammatory bowel disease: a nation-wide population-based study. 2019 , 49, 165-172	10
305	Use of oral contraceptives and risk of ulcerative colitis - A systematic review and meta-analysis. 2019 , 139, 367-374	6

304 Current Medical Therapies for Ulcerative Colitis. **2019**, 1-15

303	Copy number variation-based gene set analysis reveals cytokine signalling pathways associated with psychiatric comorbidity in patients with inflammatory bowel disease. 2020 , 112, 683-693	4
302	Emerging views of mitophagy in immunity and autoimmune diseases. 2020 , 16, 3-17	111
301	Live cell optical assay for precise characterization of receptors coupling to G#2. 2020 , 126 Suppl 6, 88-95	1
300	Genetic Predisposition, Humans. 2020 , 383-418	2
299	RIPK1 Mediates TNF-Induced Intestinal Crypt Apoptosis During Chronic NF- B Activation. 2020 , 9, 295-312	14
298	Multiple outcome meta-analysis of gene-expression data in inflammatory bowel disease. 2020 , 112, 1761-176	67 ₅
297	Altered gene regulation as a candidate mechanism by which ciliopathy gene SDCCAG8 contributes to schizophrenia and cognitive function. 2020 , 29, 407-417	1
296	Molecular systems in inflammatory bowel disease. 2020 , 367-388	1
295	Epithelial Barrier Function. 2020 , 300-313	1
294	Epigenetic methylation in Eosinophilic Esophagitis: Molecular ageing and novel biomarkers for treatment response. 2020 , 50, 1372-1380	3
293	Rosacea. 2020 ,	O
292	The crucial role of non-coding RNAs in the pathophysiology of inflammatory bowel disease. 2020 , 129, 110507	11
291	Modulation of TCR Signaling by Tyrosine Phosphatases: From Autoimmunity to Immunotherapy. 2020 , 8, 608747	9
290	Deep sequencing reveals a DAP1 regulatory haplotype that potentiates autoimmunity in systemic lupus erythematosus. 2020 , 21, 281	3
289	Genome-wide association study of prevalent and persistent cervical high-risk human papillomavirus (HPV) infection. 2020 , 21, 231	4
288	Beyond Low-Earth Orbit: Characterizing Immune and microRNA Differentials following Simulated Deep Spaceflight Conditions in Mice. 2020 , 23, 101747	8
287	Biochemical transformation of bacterial lipopolysaccharides by acyloxyacyl hydrolase reduces host injury and promotes recovery. 2020 , 295, 17842-17851	6

286	ESRRA (estrogen related receptor alpha) is a critical regulator of intestinal homeostasis through activation of autophagic flux via gut microbiota. 2021 , 17, 2856-2875	8
285	Trans-Ancestral Fine-Mapping and Epigenetic Annotation as Tools to Delineate Functionally Relevant Risk Alleles at and in Systemic Lupus Erythematosus. <i>International Journal of Molecular</i> 6.3 Sciences, 2020 , 21,	1
284	The association of genetic variants IL2RA rs2104286, IFI30 rs11554159 and IKZF3 rs12946510 with multiple sclerosis onset and severity in patients from Serbia. 2020 , 347, 577346	1
283	Anxiety, depression, chronic inflammation and aortic stiffness in Crohn's disease: the braingutvascular axis. 2020 , 38, 2008-2017	4
282	PARK7 Diminishes Oxidative Stress-Induced Mucosal Damage in Celiac Disease. 2020 , 2020, 4787202	3
281	Gut-Brain Axis: Potential Factors Involved in the Pathogenesis of Parkinson's Disease. 2020 , 11, 849	5
280	A Novel Genetic Variant Mapping on the Genomic Region Targeted by Mongersen Is Associated with Crohn's Disease. 2020 , 8,	2
279	Identification of novel loci controlling inflammatory bowel disease susceptibility utilizing the genetic diversity of wild-derived mice. 2020 , 21, 311-325	3
278	Detecting ulcerative colitis from colon samples using efficient feature selection and machine learning. 2020 , 10, 13744	13
277	-Cymene and Rosmarinic Acid Ameliorate TNBS-Induced Intestinal Inflammation Upkeeping ZO-1 and MUC-2: Role of Antioxidant System and Immunomodulation. <i>International Journal of Molecular 6.3 Sciences</i> , 2020 , 21,	14
276	Inflammatory Bowel Disease: The Emergence of New Trends in Lifestyle and Nanomedicine as the Modern Tool for Pharmacotherapy. 2020 , 10,	8
275	IMAGINE Network's ind nd ut nteractions ohort (MAGIC) Study: a protocol for a prospective observational multicentre cohort study in inflammatory bowel disease and irritable bowel syndrome. 2020 , 10, e041733	2
274	Association between IL12B polymorphisms and inflammatory bowel disease in Caucasian population: A meta-analysis. 2020 , 136, 155296	1
273	Treatment of Allergic Asthma with Fenretinide Formulation (LAU-7b) Downregulates ORMDL Sphingolipid Biosynthesis Regulator 3 () Expression and Normalizes Ceramide Imbalance. 2020 , 373, 476-487	3
272	Donor-derived hypouricemia in irrelevant recipients caused by kidney transplantation. 2020 , 8, 330	1
271	Genetic Variants of the MGAT5 Gene Are Functionally Implicated in the Modulation of T Cells Glycosylation and Plasma IgG Glycome Composition in Ulcerative Colitis. 2020 , 11, e00166	6
271 270		4

268	Genome-Wide Association Study of Cryptosporidiosis in Infants Implicates. 2020, 11,		12
267	Genetics of systemic sclerosis 2020 , 5, 192-201		1
266	Interplay between Cytokine Circuitry and Transcriptional Regulation Shaping Helper T Cell Pathogenicity and Plasticity in Inflammatory Bowel Disease. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	6
265	The Role of Tissue-Specific Ubiquitin Ligases, RNF183, RNF186, RNF182 and RNF152, in Disease and Biological Function. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	10
264	Candidate polymorphisms and susceptibility to inflammatory bowel disease: A systematic review and meta-analysis. 2020 , 753, 144814		0
263	Pitfalls in Single Clone CRISPR-Cas9 Mutagenesis to Fine-map Regulatory Intervals. 2020 , 11,		2
262	JASS: command line and web interface for the joint analysis of GWAS results. 2020 , 2, lqaa003		4
261	Metagenomics for taxonomy profiling: tools and approaches. 2020 , 11, 356-374		62
260	Role of IRF8 in immune cells functions, protection against infections, and susceptibility to inflammatory diseases. 2020 , 139, 707-721		22
259	Clinical Pharmacology of Janus Kinase Inhibitors in Inflammatory Bowel Disease. 2020 , 14, S725-S736		9
258	Genetic drug target validation using Mendelian randomisation. 2020, 11, 3255		34
257	Genome-Wide Association Study Data Reveal Genetic Susceptibility to Chronic Inflammatory Intestinal Diseases and Pancreatic Ductal Adenocarcinoma Risk. 2020 , 80, 4004-4013		1
256	miR-125b and miR-223 Contribute to Inflammation by Targeting the Key Molecules of NF B Pathway. 2019 , 6, 313		14
255	Discovery of inflammatory bowel disease-associated miRNAs using a novel bipartite clustering approach. 2020 , 13, 10		7
254	Deficiency in the anti-apoptotic protein DJ-1 promotes intestinal epithelial cell apoptosis and aggravates inflammatory bowel disease via p53. 2020 , 295, 4237-4251		11
253	Diseases of the digestive system. 2020 , 443-491		O
252	Genetic Analysis of Ulcerative Colitis in Japanese Individuals Using Population-specific SNP Array. 2020 , 26, 1177-1187		1
251	Germline Features Associated with Immune Infiltration in Solid Tumors. 2020 , 30, 2900-2908.e4		14

250 Genetics and Risk of Autoimmune Liver Diseases. **2020**, 47-64

249	Targeting Janus Kinases and Signal Transducer and Activator of Transcription 3 to Treat Inflammation, Fibrosis, and Cancer: Rationale, Progress, and Caution. 2020 , 72, 486-526	62
248	Mapping the 17q12-21.1 Locus for Variants Associated with Early-Onset Asthma in African Americans. 2021 , 203, 424-436	5
247	Genetic analyses identify GSDMB associated with asthma severity, exacerbations, and antiviral pathways. 2021 , 147, 894-909	15
246	N-glycans as functional effectors of genetic and epigenetic disease risk. 2021 , 79, 100891	13
245	Inflammatory Bowel Diseases (IBD) and the Microbiome-Searching the Crime Scene for Clues. 2021 , 160, 524-537	67
244	Association of a Gene Haplotype with Behcet's Disease in a Chinese Han Population. 2021, 29, 219-227	1
243	Epigenomics of intestinal disease. 2021 , 213-230	
242	Identification of susceptibility loci for Takayasu arteritis through a large multi-ancestral genome-wide association study. 2021 , 108, 84-99	8
241	Ulcerative Colitis in Children and Adolescents. 2021 , 474-492.e8	Ο
240	The Interrelationships between Intestinal Permeability and Phlegm Syndrome and Therapeutic Potential of Some Medicinal Herbs. 2021 , 11,	1
239	Plectin ensures intestinal epithelial integrity and protects colon against colitis. 2021 , 14, 691-702	7
238	Expanding the drug discovery space with predicted metabolite-target interactions. 2021 , 4, 288	1
237	Distinct transcriptomic signature of mRNA and microRNA in ulcerative colitis and irritable bowel syndrome.	1
236	Targeting RIP Kinases in Chronic Inflammatory Disease. 2021 , 11,	7
235	Using Hierarchical Similarity To Examine The Genetics of Beh⊞t⊠ Disease.	
234	Novel and Emerging Therapies for Inflammatory Bowel Disease. 2021 , 12, 651415	17
233	Alleviation of colonic inflammation by Lypd8 in a mouse model of inflammatory bowel disease. 2021 , 33, 359-372	2

232	Intestinal immunoregulation: lessons from human mendelian diseases. 2021 , 14, 1017-1037	2
231	Interleukin-28A induces epithelial barrier dysfunction in CD patient-derived intestinal organoids. 2021 , 320, G689-G699	4
230	G-protein coupled receptor 35 (GPR35) regulates the colonic epithelial cell response to enterotoxigenic Bacteroides fragilis. 2021 , 4, 585	7
229	Genome-wide association study identifies TNFSF15 associated with childhood asthma. 2021 ,	2
228	Gene Polymorphisms of NOD2, IL23R, PTPN2 and ATG16L1 in Patients with Crohn's Disease: On the Way to Personalized Medicine?. 2021 , 12,	1
227	A netrin domain-containing protein secreted by the human hookworm Necator americanus protects against CD4 T cell transfer colitis. 2021 , 232, 88-102	4
226	Small Effects: The Indispensable Foundation for a Cumulative Psychological Science. 2021 , 17456916209844	18342
225	Associations of GWAS-Supported Non-MHC Genes with Autoimmune Thyroiditis in Patients with Type 1 Diabetes. 2021 , 14, 3017-3026	
224	A susceptibility locus in the IL12B but not LILRA3 region is associated with vascular damage in Takayasu arteritis. 2021 , 11, 13667	2
223	LRRK2 is required for CD38-mediated NAADP-Ca signaling and the downstream activation of TFEB (transcription factor EB) in immune cells. 2021 , 1-19	2
222	Role of the IL33 and IL1RL1 pathway in the pathogenesis of Immunoglobulin A vasculitis. 2021, 11, 16163	1
221	Inflammasomes and Colorectal Cancer. 2021 , 10,	2
220	Hepatitis B Virus DNA Integration, Chronic Infections and Hepatocellular Carcinoma. 2021, 9,	2
219	The Candidate IBD Risk Gene Is Dispensable for Intestinal Epithelial Homeostasis. 2021, 10,	1
218	Genetic and Environmental Determinants of T Helper 17 Pathogenicity in Spondyloarthropathies. 2021 , 12, 703242	
217	Machine Learning Modeling from Omics Data as Prospective Tool for Improvement of Inflammatory Bowel Disease Diagnosis and Clinical Classifications. 2021 , 12,	O
216	Quantifying mediation between omics layers and complex traits.	0
215	Genetics of Asthma: Insights From Genome Wide Association Studies. 2022 , 308-325	

214	Genetic Epidemiology of Inflammatory Bowel Disease, Early Twin and Family Studies. 2013, 23-43	1
213	Genetics of Ulcerative Colitis. 2013 , 119-134	1
212	Antimicrobial Peptides and Inflammatory Bowel Disease. 2013, 255-273	1
211	Genetics of lupus. 2015 , 1045-1051	2
210	Inflammatory Bowel Disease. 2011 , 1294-1304.e1	3
209	GETECCU 2020 guidelines for the treatment of ulcerative colitis. Developed using the GRADE approach. 2020 , 43 Suppl 1, 1-57	3
208	It Takes Three Receptors to Raise a B Cell. 2020 , 41, 629-642	1
207	An xQTL map integrates the genetic architecture of the human brain's transcriptome and epigenome. 2017 , 20, 1418-1426	201
206	Host-microbiota interactions in inflammatory bowel disease. 2020 , 20, 411-426	133
205	Activation of pH-Sensing Receptor OGR1 (GPR68) Induces ER Stress Via the IRE1別K Pathway in an Intestinal Epithelial Cell Model. 2020 , 10, 1438	21
204	Increased risk of ischemic heart disease and diabetes in înflammatory bowel disease. 2021 , 59, 117-124	3
203	Genome-wide association studies suggest limited immune gene enrichment in schizophrenia compared to five autoimmune diseases.	2
202	Genome-wide association study identifies 17 new loci influencing concentrations of circulating cytokines and growth factors.	2
201	LD Hub: a centralized database and web interface to perform LD score regression that maximizes the potential of summary level GWAS data for SNP heritability and genetic correlation analysis.	2
200	Genome-wide association study implicates immune activation of multiple integrin genes in inflammatory bowel disease.	4
199	Exploring the genetic architecture of inflammatory bowel disease by whole genome sequencing identifies association at ADCY7.	3
198	Genetic correlations among psychiatric and immune-related phenotypes based on genome-wide association data.	2
197	GARFIELD - GWAS Analysis of Regulatory or Functional Information Enrichment with LD correction.	16

196	Shared activity patterns arising at genetic susceptibility loci reveal underlying genomic and cellular architecture of human disease.	2
195	Type 1 diabetes genome-wide association analysis with imputation identifies five new risk regions.	8
194	Genome-wide association study reveals genetic link between diarrhea-associated Entamoeba histolytica infection and inflammatory bowel disease.	2
193	A large-scale genome-wide enrichment analysis identifies new trait-associated genes, pathways and tissues across 31 human phenotypes*.	5
192	Genetic associations at regulatory phenotypes improve fine-mapping of causal variants for twelve immune-mediated diseases.	8
191	Family-based genome-wide association study of leprosy in Vietnam.	1
190	Population Structure Discovery in Meta-Analyzed Microbial Communities and Inflammatory Bowel Disease.	3
189	Degradation of the extracellular matrix is part of the pathology of ulcerative colitis□	2
188	Genetic and Epigenetic Fine Mapping of Complex Trait Associated Loci in the Human Liver.	2
187	TFF3 is a ligand for LINGO2 that de-represses EGFR to control disease outcome during colitis and gastrointestinal nematode infection.	1
186	Genome-wide association study of cryptosporidiosis in infants implicatesPRKCA.	3
185	IL-10 induces a STAT3-dependent autoregulatory loop in T2 cells that promotes Blimp-1 restriction of cell expansion via antagonism of STAT5 target genes. 2016 , 1,	19
184	IL-7 receptor influences anti-TNF responsiveness and T cell gut homing in inflammatory bowel disease. 2019 , 129, 1910-1925	38
183	Regulation of dendritic cell activation by microRNA let-7c and BLIMP1. 2013 , 123, 823-33	61
182	Genetics and Therapeutics in Pediatric Ulcerative Colitis: the Past, Present and Future. 2016 , 5,	3
181	Genetic associations of inflammatory bowel disease in a South Asian population. 2018 , 6, 908-915	5
180	Investigating the Causal Relationship of C-Reactive Protein with 32 Complex Somatic and Psychiatric Outcomes: A Large-Scale Cross-Consortium Mendelian Randomization Study. 2016 , 13, e1001976	100
179	Investigation of multiple susceptibility loci for inflammatory bowel disease in an Italian cohort of patients. 2011 , 6, e22688	45

178	Genetic evidence supporting the association of protease and protease inhibitor genes with inflammatory bowel disease: a systematic review. 2011 , 6, e24106	24
177	The role of osteopontin (OPN/SPP1) haplotypes in the susceptibility to Crohn's disease. 2011 , 6, e29309	31
176	A functional variant in ERAP1 predisposes to multiple sclerosis. 2012 , 7, e29931	41
175	PTPN2 gene variants are associated with susceptibility to both Crohn's disease and ulcerative colitis supporting a common genetic disease background. 2012 , 7, e33682	46
174	Genome-wide association study of multiple sclerosis confirms a novel locus at 5p13.1. 2012 , 7, e36140	41
173	Association of single nucleotide polymorphisms of IL23R and IL17 with ulcerative colitis risk in a Chinese Han population. 2012 , 7, e44380	22
172	IRGM variants and susceptibility to inflammatory bowel disease in the German population. 2013 , 8, e54338	51
171	Genetic Association of Peptidoglycan Recognition Protein Variants with Inflammatory Bowel Disease. 2013 , 8, e67393	19
170	Intestinal DMBT1 expression is modulated by Crohn's disease-associated IL23R variants and by a DMBT1 variant which influences binding of the transcription factors CREB1 and ATF-2. 2013 , 8, e77773	18
169	A modular organization of the human intestinal mucosal microbiota and its association with inflammatory bowel disease. 2013 , 8, e80702	124
169 168		124
	Job strain and the risk of inflammatory bowel diseases: individual-participant meta-analysis of	
168	inflammatory bowel disease. 2013, 8, e80702 Job strain and the risk of inflammatory bowel diseases: individual-participant meta-analysis of 95,000 men and women. 2014, 9, e88711 Transcriptome profiling of human ulcerative colitis mucosa reveals altered expression of pathways	14
168 167	inflammatory bowel disease. 2013, 8, e80702 Job strain and the risk of inflammatory bowel diseases: individual-participant meta-analysis of 95,000 men and women. 2014, 9, e88711 Transcriptome profiling of human ulcerative colitis mucosa reveals altered expression of pathways enriched in genetic susceptibility loci. 2014, 9, e96153	14
168 167 166	Job strain and the risk of inflammatory bowel diseases: individual-participant meta-analysis of 95,000 men and women. 2014, 9, e88711 Transcriptome profiling of human ulcerative colitis mucosa reveals altered expression of pathways enriched in genetic susceptibility loci. 2014, 9, e96153 Association of NOD2 and IL23R with inflammatory bowel disease in Puerto Rico. 2014, 9, e108204 A candidate gene approach identifies an IL33 genetic variant as a novel genetic risk factor for GCA.	14 8 9
168167166165	Job strain and the risk of inflammatory bowel diseases: individual-participant meta-analysis of 95,000 men and women. 2014, 9, e88711 Transcriptome profiling of human ulcerative colitis mucosa reveals altered expression of pathways enriched in genetic susceptibility loci. 2014, 9, e96153 Association of NOD2 and IL23R with inflammatory bowel disease in Puerto Rico. 2014, 9, e108204 A candidate gene approach identifies an IL33 genetic variant as a novel genetic risk factor for GCA. 2014, 9, e113476 Effective population size, extended linkage disequilibrium and signatures of selection in the rare	14 8 9
168167166165164	Job strain and the risk of inflammatory bowel diseases: individual-participant meta-analysis of 95,000 men and women. 2014, 9, e88711 Transcriptome profiling of human ulcerative colitis mucosa reveals altered expression of pathways enriched in genetic susceptibility loci. 2014, 9, e96153 Association of NOD2 and IL23R with inflammatory bowel disease in Puerto Rico. 2014, 9, e108204 A candidate gene approach identifies an IL33 genetic variant as a novel genetic risk factor for GCA. 2014, 9, e113476 Effective population size, extended linkage disequilibrium and signatures of selection in the rare dog breed lundehund. 2015, 10, e0122680 Abundant genetic overlap between blood lipids and immune-mediated diseases indicates shared	14 8 9 14

160	The CARD9 Polymorphisms rs4077515, rs10870077 and rs10781499 Are Uncoupled from Susceptibility to and Severity of Pulmonary Tuberculosis. 2016 , 11, e0163662		5
159	ShinyGPA: An interactive visualization toolkit for investigating pleiotropic architecture using GWAS datasets. 2018 , 13, e0190949		2
158	IL-23R mutation is associated with ulcerative colitis: A systemic review and meta-analysis. 2017 , 8, 484	9-4863	14
157	Associations between , and and Beh∄t's disease but not VKH syndrome in Han Chinese. 2017 , 8, 10503	7-1050	46 ₄
156	Erbin exerts a protective effect against inflammatory bowel disease by suppressing autophagic cell death. 2018 , 9, 12035-12049		11
155	Cancer-predicting transcriptomic and epigenetic signatures revealed for ulcerative colitis in patient-derived epithelial organoids. 2018 , 9, 28717-28730		14
154	Micro-RNAs -106a and -362-3p in Peripheral Blood of Inflammatory Bowel Disease Patients. 2018 , 12, 78-86		15
153	The role of diet in the prevention and treatment of Inflammatory Bowel Diseases. 2018 , 89, 60-75		48
152	Current issues of pediatric inflammatory bowel disease in Korea. 2014 , 57, 465-71		8
151	Recurrence and rejection in liver transplantation for primary sclerosing cholangitis. <i>World Journal of Gastroenterology</i> , 2012 , 18, 1-15	5.6	98
150	Analysis of single nucleotide polymorphisms in the region of CLDN2-MORC4 in relation to inflammatory bowel disease. <i>World Journal of Gastroenterology</i> , 2013 , 19, 4935-43	5.6	14
149	Synergistic effect of interleukin-10-receptor variants in a case of early-onset ulcerative colitis. World Journal of Gastroenterology, 2013 , 19, 8659-70	5.6	23
148	Susceptibility to ulcerative colitis in Hungarian patients determined by gene-gene interactions. World Journal of Gastroenterology, 2014 , 20, 219-27	5.6	10
147	Interleukin and interleukin receptor gene polymorphisms in inflammatory bowel diseases susceptibility. World Journal of Gastroenterology, 2014 , 20, 3208-22	5.6	15
146	Inflammatory bowel disease in pediatric and adolescent patients: a biomolecular and histopathological review. <i>World Journal of Gastroenterology</i> , 2014 , 20, 10262-78	5.6	27
145	Immunogenetic biomarkers in inflammatory bowel diseases: role of the IBD3 region. <i>World Journal of Gastroenterology</i> , 2014 , 20, 15037-48	5.6	11
144	Associations between CD24 gene polymorphisms and inflammatory bowel disease: A meta-analysis. <i>World Journal of Gastroenterology</i> , 2015 , 21, 6052-9	5.6	2
143	Protein tyrosine phosphatase non-receptor type 2 and inflammatory bowel disease. World Journal of Gastroenterology, 2016, 22, 1034-44	5.6	22

(2013-2020)

142	Differential regulation of JAK/STAT-signaling in patients with ulcerative colitis and Crohn's disease. <i>World Journal of Gastroenterology</i> , 2020 , 26, 4055-4075	5.6	12
141	Inflammatory bowel disease: Therapeutic limitations and prospective of the stem cell therapy. 2020 , 12, 1050-1066		6
140	Is Whole Exome Sequencing Clinically Practical in the Management of Pediatric Crohn's Disease?. 2015 , 9, 767-75		8
139	Epithelial-microbial diplomacy: escalating border tensions drive inflammation in inflammatory bowel disease. 2019 , 17, 177-191		8
138	Recent updates on the basic mechanisms and pathogenesis of inflammatory bowel diseases in experimental animal models. 2020 , 18, 151-167		42
137	Cholesteryl ester transfer protein (CETP) as a drug target for cardiovascular disease. 2021 , 12, 5640		7
136	Mitochondria and Inflammatory Bowel Diseases: Toward a Stratified Therapeutic Intervention. 2021 ,		3
135	Genetic control of the dynamic transcriptional response to immune stimuli and glucocorticoids at single cell resolution 2022 ,		O
134	An extremes of phenotype approach confirms significant genetic heterogeneity in patients with ulcerative colitis.		
133	The Genetics of Inflammatory Bowel Disease. 2012 , 3-16		
132	Molecular Genetics of Coeliac Disease.		
131	Genetics of Inflammatory Bowel Diseases. 2013 , 3-12		
130	Complex Disease Genes and Their Discovery. 2013 , 87-97		
129	Inflammatory Bowel Disease at the Intersection of Autophagy and Immunity: Insights from Human Genetics. 2013 , 241-264		1
128	Encyclopedia of Inflammatory Diseases. 2013 , 1-20		
127	The IL23-Th17 Axis in Intestinal Inflammation. 2013 , 219-240		
126	Genetic Overlap Between Inflammatory Bowel Disease and Other Diseases. 2013, 135-150		
125	The Genetics of Crohn⊠ Disease. 2013 , 99-118		

124	The Epithelial Barrier. 2013, 265-280	
123	Insights from Recent Advances in Animal Models of Inflammatory Bowel Disease. 2013 , 45-83	1
122	Genetics of inflammatory bowel disease: the state of play. 2013 , 28, 759-60	
121	Methylation QTLs are associated with coordinated changes in transcription factor binding, histone modifications, and gene expression levels.	1
120	Autophagy Restricts Interleukin-1 (5) ignaling via Regulation of P62 Stability. 2015, 223-229	
119	Inflammatory Bowel Disease. 2016 , 183-201	
118	Compendium of Inflammatory Diseases. 2016 , 643-659	
117	Mucosal Surfaces: Immunological Protection. 1-12	
116	An ancient fecundability-associated polymorphism switches a repressor into an enhancer of endometrial TAP2 Expression.	
115	High-resolution DNA accessibility profiles increase the discovery and interpretability of genetic associations.	
114	Trauma, Regulated Cell Death, and Inflammation. 2017 , 253-281	
113	Genetics of Inflammatory Bowel Diseases. 2017 , 3-14	1
112	Narrating Genes: How Patients with Chronic Inflammatory Bowel Diseases Interpret an Emerging Disease Aetiology and How We Can Make Sense Out of It by Developing a Historically and Sociologically Informed Framework. 2017 , 325-336	
111	Histopathological image QTL discovery of immune infiltration variants.	1
110	Immune-related genetic enrichment in frontotemporal dementia.	
109	Genetic Influences on the Periodontal Microbial-Host Crosstalk. 2018 , 87-95	
108	An ancient fecundability-associated polymorphism creates a new GATA2 binding site in a distal enhancer of HLA-F.	
107	Evidence for a potential role of miR-1908-5p and miR-3614-5p in autoimmune disease risk using genome-wide analyses.	O

106	Systems genetic discovery of host-microbiome interactions reveals mechanisms of microbial involvement in disease.	1
105	Mikrobiomische Selbstwirksamkeit. 2019 , 43-66	
104	IBD Genomic Risk Loci and Overlap with Other Inflammatory Diseases. 2019 , 91-115	
103	Presentation and Natural Course of Ulcerative Colitis. 2019 , 17-28	
102	Drugs for Soft Tissue Autoimmune Disorders. 2019 , 751-775	
101	Inflammatory Bowel Disease and Epigenetics. 2019 , 183-201	O
100	The IL-23/Th17 Axis in Intestinal Inflammation. 2019 , 281-303	
99	Inflammatory Bowel Disease at the Intersection of Autophagy and Immunity: Insights from Human Genetics. 2019 , 305-328	1
98	Complex Disease Genes and Their Discovery. 2019 , 79-89	
97	Sequencing and Mapping IBD Genes to Individual Causative Variants and Their Clinical Relevance. 2019 , 117-139	
96	Germline features associated with immune infiltration in solid tumors.	
95	JASS: Command Line and Web interface for the joint analysis of GWAS results.	1
94	The role of TNF-alpha gene (-238G/A and -308G/A) polymorphisms in the etiology and pathogenesis of inflammatory bowel diseases in various ethnic groups. 2019 , 47, 548-558	O
93	Genetics of Rosacea. 2020 , 23-34	
92	Peptide YY: a novel Paneth cell antimicrobial peptide that maintains fungal commensalism.	1
91	Genome-Wide Association Study of Prevalent and Persistent Cervical High-Risk Human Papillomavirus (HPV) Infection.	
90	Genome-Wide Association Study of Prevalent and Persistent Cervical High-Risk Human Papillomavirus(HPV) Infection.	
89	Regulation of meprin metalloproteases in mucosal homeostasis. 2022 , 1869, 119158	1

88 Inflammatory bowel diseases and genetic.

87	Bflamatuvar barsak hastal&lar⊕e genetik.		
86	Plectin Ensures Intestinal Epithelial Integrity and Protects Colon Against Colitis.		
85	Current and future role of serogenomics in ulcerative colitis. 2011 , 7, 720-7		
84	Potential treatment of inflammatory bowel disease: a review of helminths therapy. 2014 , 7, 9-16		28
83	The history of genetics in inflammatory bowel disease. 2014 , 27, 294-303		83
82	Study on the Multi-marker Components Quantitative HPLC Fingerprint of the Compound Chinese Medicine Wuwei Changyanning Granule. 2014 , 13, 1191-201		2
81	TNF#nhibitor induced lupus-like syndrome (TAILS) in a patient with IBD. 2014 , 40, 285-8		1
80	Causal Estimation with Functional Confounders. 2020 , 33, 5115-5125		
79	Association Analysis of , and Polymorphisms in Chinese Patients With Parkinson's Disease and Multiple System Atrophy. 2021 , 12, 765833		O
78	Identification of a gut microbiota member that ameliorates DSS-induced colitis in intestinal barrier enhanced Dusp6-deficient mice. 2021 , 37, 110016		3
77	Hygiene Hypothesis as the Etiology of Kawasaki Disease: Dysregulation of Early B Cell Development. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	O
76	Research-Based Product Innovation to Address Critical Unmet Needs of Patients with Inflammatory Bowel Diseases. 2021 , 27, S1-S16		1
75	Pharmacogenetics of Biological Agents Used in Inflammatory Bowel Disease: A Systematic Review 2021 , 9,		1
74	Nedd4-2-dependent Ubiquitination Potentiates the Inhibition of Human NHE3 by Cholera Toxin and Enteropathogenic Escherichia coli. 2021 ,		1
73	Finding the sweet spot: glycosylation mediated regulation of intestinal inflammation. 2021,		2
72	Radioiodination of balsalazide, bioevaluation and characterization as a highly selective radiotracer for imaging of ulcerative colitis in mice 2022 ,		6
71	Developing Biliary Atresia-like Model by Treating Human Liver Organoids with Polyinosinic:Polycytidylic Acid (Poly (I:C)). 2022 , 44, 644-653		O

(2019-2021)

70	Type 2 Inflammation in Eosinophilic Esophagitis: From Pathophysiology to Therapeutic Targets 2021 , 12, 815842		4
69	Probiotics, prebiotics and synbiotics: Safe options for next-generation therapeutics 2022 , 106, 505		7
68	pH-Sensing G Protein-Coupled Receptor OGR1 (GPR68) Expression and Activation Increases in Intestinal Inflammation and Fibrosis <i>International Journal of Molecular Sciences</i> , 2022 , 23,	6.3	1
67	Conserved and Unique Functions of Blimp1 in Immune Cells 2021 , 12, 805260		3
66	Genetics of ulcerative colitis: putting into perspective the incremental gains from Indian studies. 2018 , 97, 1493-1507		2
65	Micronutrient Improvement of Epithelial Barrier Function in Various Disease States: A Case for Adjuvant Therapy <i>International Journal of Molecular Sciences</i> , 2022 , 23,	6.3	2
64	Role of Energy Metabolism and Mitochondrial Function in Inflammatory Bowel Disease 2022,		1
63	Biomedical consequences of elevated cholesterol-containing lipoproteins and apolipoproteins.		
62	Variants in Interferon Lambda are Associated with Very Early Onset Inflammatory Bowel Disease.		
61	SMetABF: A rapid algorithm for Bayesian GWAS meta-analysis with a large number of studies included 2022 , 18, e1009948		O
60	Genetic associations at regulatory phenotypes improve fine-mapping of causal variants for 12 immune-mediated diseases <i>Nature Genetics</i> , 2022 , 54, 251-262	36.3	1
59	miR-374a-5p regulates inflammatory genes and monocyte function in patients with inflammatory bowel disease 2022 , 219,		O
58	Organisation of gene programs revealed by unsupervised analysis of diverse gene-trait associations.		
57	Prostaglandins and calprotectin are genetically and functionally linked to the Inflammatory Bowel Diseases.		
56	Genetic variants involved in innate immunity modulate the risk of inflammatory bowel diseases in an understudied Malaysian population. 2021 ,		1
55	An Update of Research Animal Models of Inflammatory Bowel Disease 2021 , 2021, 7479540		2
54	Data_Sheet_1.ZIP. 2019 ,		
53	Presentation_1.pdf. 2019,		

52	Table_1.XLSX. 2018 ,	
51	Table_2.XLSX. 2018 ,	
50	Table_3.XLSX. 2018 ,	
49	Image_1.JPEG. 2018 ,	
48	Image_2.JPEG. 2018 ,	
47	Image_3.JPEG. 2018 ,	
46	Image_4.JPEG. 2018 ,	
45	Image_5.JPEG. 2018 ,	
44	Image_1.JPEG. 2018 ,	
43	Table_1.xlsx. 2018 ,	
42	Table_2.xlsx. 2018 ,	
41	Table_3.XLSX. 2018 ,	
40	Glucocorticoid receptor modulates dendritic cell function in ulcerative colitis. 2020, 35, 1379-1389	2
39	OUP accepted manuscript.	O
38	The role of the BTLA-HVEM complex in the pathogenesis of autoimmune diseases 2022 , 376, 104532	0
37	Significance of IL-7 and IL-7R in RA and autoimmunity. 2022 , 21, 103120	1
36	Intestine-Specific NHE3 Deletion in Adulthood Causes Microbial Dysbiosis. 2022 , 12,	1
35	Investigating immunoregulatory effects of myeloid cell autophagy in acute and chronic inflammation.	

34	PARK7/DJ-1 as a Therapeutic Target in Gut-Brain Axis Diseases. <i>International Journal of Molecular Sciences</i> , 2022 , 23, 6626	6.3	2
33	Inverse and Concordant Mucosal Pathway Gene Expressions in Inflamed and Non-Inflamed Ulcerative Colitis Patients: Potential Relevance to Aetiology and Pathogenesis. <i>International Journal of Molecular Sciences</i> , 2022 , 23, 6944	6.3	O
32	Epigenetics in IBD: a conceptual framework for disease pathogenesis. <i>Frontline Gastroenterology</i> , 2022 , 13, e22-e27	2.6	1
31	Co-existing ulcerative colitis and takayasu arteritis: A case-based review. <i>Indian Journal of Rheumatology</i> , 2022 ,	0.5	
30	Characteristics of inflammatory bowel diseases in patients with concurrent immune-mediated inflammatory diseases. <i>World Journal of Gastroenterology</i> , 2022 , 28, 2843-2853	5.6	
29	CD30L is involved in the regulation of the inflammatory response through inducing homing and differentiation of monocytes via CCL2/CCR2 axis and NF-B pathway in mice with colitis. International Immunopharmacology, 2022, 110, 108934	5.8	O
28	A large-scale genome-wide cross-trait analysis reveals shared genetic architecture between Alzheimer disease and gastrointestinal tract disorders. 2022 , 5,		2
27	Inflammatory Bowel Diseases Before and After 1990. 2022 ,		
26	Current and projected incidence trends of pediatric-onset inflammatory bowel disease in Germany based on the Saxon Pediatric IBD Registry 2000\(\textbf{Q} 014 \textbf{B} 15-year evaluation of trends. \textbf{2022}, 17, e0274117	,	О
25	Etiology of Ulcerative Colitis.		O
24	Prostaglandins and calprotectin are genetically and functionally linked to the Inflammatory Bowel Diseases. 2022 , 18, e1010189		О
23	An extremes of phenotype approach confirms significant genetic heterogeneity in patients with ulcerative colitis.		O
22	Population structure discovery in meta-analyzed microbial communities and inflammatory bowel disease using MMUPHin. 2022 , 23,		1
21	Trans-ancestry, Bayesian Meta-analysis Discovers 20 Novel Risk Loci for Inflammatory Bowel Disease in an African American, East Asian, and European Cohort.		1
20	Overlapping genetic susceptibility of seven autoimmune diseases:SPU tests based on genome-wide association summary statistics. 2023 , 851, 147036		О
19	Heat Shock Transcription Factor 2 Is Significantly Involved in Neurodegenerative Diseases, Inflammatory Bowel Disease, Cancer, Male Infertility, and Fetal Alcohol Spectrum Disorder: The Novel Mechanisms of Several Severe Diseases. 2022 , 23, 13763		1
18	Mitochondrial Control in Inflammatory Gastrointestinal Diseases. 2022 , 23, 14890		О
17	Fecal Luminal Factors from Patients with Gastrointestinal Diseases Alter Gene Expression Profiles in Caco-2 Cells and Colonoids. 2022 , 23, 15505		O

16	Machine Learning Reveals Genetic Modifiers of the Immune Microenvironment of Cancer.	O
15	Biomedical consequences of elevated cholesterol-containing lipoproteins and apolipoproteins on cardiovascular and non-cardiovascular outcomes. 2023 , 3,	1
14	The landscape of immune dysregulation in Crohn disease revealed through single-cell transcriptomic profiling in the ileum and colon. 2023 ,	0
13	Protein kinase modulation for anti-aging intervention. 2023 , 285-304	0
12	The genetics of non-monogenic IBD.	0
11	Gastrointestinal disorders and intestinal bacteria: Advances in research and applications in therapy. 9,	O
10	Genetics of Inflammatory Bowel Diseases. 2023 , 3-14	0
9	Translating non-coding genetic associations into a better understanding of immune-mediated disease. 2023 , 16,	O
8	A family with ulcerative colitis maps to 7p21.1 and comprises a region with regulatory activity for the aryl hydrocarbon receptor gene.	0
7	Immunoregulation by antigen-presenting cells in human intestinal lamina propria. 14,	1
6	Treatment Effects of Natural Products on Inflammatory Bowel Disease In Vivo and Their Mechanisms: Based on Animal Experiments. 2023 , 15, 1031	O
5	Comparative Analysis of the Effect of Different Concentrations of Dextran Sodium Sulfate on the Severity and Extent of Inflammation in Experimental Ulcerative Colitis. 2023 , 13, 3233	O
4	Phenome-wide Mendelian randomization study evaluating the association of circulating vitamin D with complex diseases. 10,	0
3	The Study of the Association of Polymorphisms in LSP1, GPNMB, PDPN, TAGLN, TSPO, and TUBB6 Genes with the Risk and Outcome of Ischemic Stroke in the Russian Population. 2023 , 24, 6831	O
2	Identification of hub genes and immune infiltration in ulcerative colitis using bioinformatics. 2023 , 13,	0
1	Clinical characteristics and genotype analysis of a Chinese patient with juvenile arthritis due to novel LACC1 frameshift mutation and literature review.	O