Marcella Devoto

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/4517212/marcella-devoto-publications-by-year.pdf
Version: 2024-04-23

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.
The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

136 papers	9,510 citations	45 h-index	96 g-index
143 ext. papers	10,643 ext. citations	9.6 avg, IF	4.85 L-index

#	Paper	IF	Citations
136	Variants in STXBP3 are Associated with Very Early Onset Inflammatory Bowel Disease, Bilateral Sensorineural Hearing Loss and Immune Dysregulation. *Journal of Crohn's and Colitis*, **2021**, 15, 1908-1919	7.5	0
135	Immune Dysregulation in Human ITCH Deficiency Successfully Treated with Hematopoietic Cell Transplantation. *Journal of Allergy and Clinical Immunology: in Practice*, **2021**, 9, 2885-2893.e3	5.4	0
134	Colonoids From Patients With Pediatric Inflammatory Bowel Disease Exhibit Decreased Growth Associated With Inflammation Severity and Durable Upregulation of Antigen Presentation Genes. *Inflammatory Bowel Diseases*, **2021**, 27, 256-267	4.5	1
133	The Unique Disease Course of Children with Very Early onset-Inflammatory Bowel Disease. *Inflammatory Bowel Diseases*, **2020**, 26, 909-918	4.5	14
132	Impaired Redox and Protein Homeostasis as Risk Factors and Therapeutic Targets in Toxin-Induced Biliary Atresia. *Gastroenterology*, **2020**, 159, 1068-1084.e2	13.3	0
131	Exome Sequencing in Individuals with Isolated Biliary Atresia. *Scientific Reports*, **2020**, 10, 2709	4.9	7
130	Rare copy number variants in over 100,000 European ancestry subjects reveal multiple disease associations. *Nature Communications*, **2020**, 11, 255	17.4	17
129	Complex genetic signatures in immune cells underlie autoimmunity and inform therapy. *Nature Genetics*, **2020**, 52, 1036-1045	36.3	16
128	Neural crest-derived tumor neuroblastoma and melanoma share 1p13.2 as susceptibility locus that shows a long-range interaction with the SLC16A1 gene. *Carcinogenesis*, **2020**, 41, 284-295	4.6	11
127	Exploring Shared Susceptibility between Two Neural Crest Cells Originating Conditions: Neuroblastoma and Congenital Heart Disease. *Genes*, **2019**, 10,	4.2	8
126	Expanding the spectrum of genes responsible for hereditary motor neuropathies. *Journal of Neurology, Neurosurgery and Psychiatry*, **2019**, 90, 1171-1179	5.5	19
125	An Exploratory Association Analysis of the Insulin Gene Region With Diabetes Mellitus in Two Dog Breeds. *Journal of Heredity*, **2019**, 110, 793-800	2.4	2
124	Subphenotype meta-analysis of testicular cancer genome-wide association study data suggests a role for RBFOX family genes in cryptorchidism susceptibility. *Human Reproduction*, **2018**, 33, 967-977	5.7	7
123	Fine mapping of 2q35 high-risk neuroblastoma locus reveals independent functional risk variants and suggests full-length BARD1 as tumor-suppressor. *International Journal of Cancer*, **2018**, 143, 2828-2837	7.5	34
122	A genome-wide association study identifies a susceptibility locus for biliary atresia on 2p16.1 within the gene EFEMP1. *PLoS Genetics*, **2018**, 14, e1007532	6	23
121	Commentary on Mutations in Interleukin-10 Receptor and Clinical Phenotypes in Patients with Very Early-onset Inflammatory Bowel Disease: A Chinese VEO-IBD Collaboration Group Survey. *Inflammatory Bowel Diseases*, **2017**, 23, 591-592	4.5	2
120	Novel ZBTB24 Mutation Associated with Immunodeficiency, Centromere Instability, and Facial Anomalies Type-2 Syndrome Identified in a Patient with Very Early Onset Inflammatory Bowel Disease. *Inflammatory Bowel Diseases*, **2017**, 23, 2252-2255	4.5	15

119	Common variants upstream of MLF1 at 3q25 and within CPZ at 4p16 associated with neuroblastoma. *PLoS Genetics*, **2017**, 13, e1006787	6	40
118	IQ and hemizygosity for the Val Met functional polymorphism of COMT in 22q11DS. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2016**, 171, 1112-1115	3.5	5
117	Variants of the ACTG2 gene correlate with degree of severity and presence of megacystis in chronic intestinal pseudo-obstruction. *European Journal of Human Genetics*, **2016**, 24, 1211-5	5.3	27
116	P-197 Identification of a Homozygous Mutation in the ZBTB24 Gene in a Patient with Very Early Onset Inflammatory Bowel Disease. *Inflammatory Bowel Diseases*, **2016**, 22, S68	4.5	
115	O-003 Understanding the Relevance of Whole Exome Sequencing Identified Variants in Patients with Very Early-Onset-IBD. *Inflammatory Bowel Diseases*, **2016**, 22, S1-S2	4.5	
114	Rare copy number variants and congenital heart defects in the 22q11.2 deletion syndrome. *Human Genetics*, **2016**, 135, 273-85	6.3	31
113	Polygenic inheritance of cryptorchidism susceptibility in the LE/orl rat. *Molecular Human Reproduction*, **2016**, 22, 18-34	4.4	7
112	Compound heterozygous mutations in NEK8 in siblings with end-stage renal disease with hepatic and cardiac anomalies. *American Journal of Medical Genetics, Part A*, **2016**, 170, 750-3	2.5	17
111	Genomic copy number variation association study in Caucasian patients with nonsyndromic cryptorchidism. *BMC Urology*, **2016**, 16, 62	2.2	4
110	Is a Candidate Modifier of Liver Disease Severity in Alagille Syndrome. *Cellular and Molecular Gastroenterology and Hepatology*, **2016**, 2, 663-675.e2	7.9	21
109	Phenotype specific association of the TGFBR3 locus with nonsyndromic cryptorchidism. *Journal of Urology*, **2015**, 193, 1637-45	2.5	16
108	Integration of Omics Data in Genetic Epidemiology. *Human Heredity*, **2015**, 79, 109-10	1.1	4
107	Pathway analysis supports association of nonsyndromic cryptorchidism with genetic loci linked to cytoskeleton-dependent functions. *Human Reproduction*, **2015**, 30, 2439-51	5.7	19
106	Exome sequencing analysis reveals variants in primary immunodeficiency genes in patients with very early onset inflammatory bowel disease. *Gastroenterology*, **2015**, 149, 1415-24	13.3	68
105	Copy-Number Variation of the Glucose Transporter Gene SLC2A3 and Congenital Heart Defects in the 22q11.2 Deletion Syndrome. *American Journal of Human Genetics*, **2015**, 96, 753-64	11	54
104	Association of a Novel ACTA1 Mutation With a Dominant Progressive Scapuloperoneal Myopathy in an Extended Family. *JAMA Neurology*, **2015**, 72, 689-98	17.2	27
103	Heterozygous deletion of FOXA2 segregates with disease in a family with heterotaxy, panhypopituitarism, and biliary atresia. *Human Mutation*, **2015**, 36, 631-7	4.7	31
102	Exome sequencing reveals compound heterozygous mutations in ATP8B1 in a JAG1/NOTCH2 mutation-negative patient with clinically diagnosed Alagille syndrome. *American Journal of Medical Genetics, Part A*, **2015**, 167A, 891-3	2.5	9

101	A de novo whole gene deletion of XIAP detected by exome sequencing analysis in very early onset inflammatory bowel disease: a case report. *BMC Gastroenterology*, **2015**, 15, 160	3	31
100	Rare variants in TP53 and susceptibility to neuroblastoma. *Journal of the National Cancer Institute*, **2014**, 106, dju047	9.7	83
99	Common genetic variants in NEFL influence gene expression and neuroblastoma risk. *Cancer Research*, **2014**, 74, 6913-24	10.1	69
98	Recessive and dominant mutations in COL12A1 cause a novel EDS/myopathy overlap syndrome in humans and mice. *Human Molecular Genetics*, **2014**, 23, 2339-52	5.6	78
97	Replication of a GWAS signal in a Caucasian population implicates ADD3 in susceptibility to biliary atresia. *Human Genetics*, **2014**, 133, 235-43	6.3	45
96	A candidate gene approach to identify modifiers of the palatal phenotype in 22q11.2 deletion syndrome patients. *International Journal of Pediatric Otorhinolaryngology*, **2013**, 77, 123-7	1.7	11
95	Novel COCH mutation in a family with autosomal dominant late onset sensorineural hearing impairment and tinnitus. *American Journal of Otolaryngology - Head and Neck Medicine and Surgery*, **2013**, 34, 230-5	2.8	20
94	Evidence from human and zebrafish that GPC1 is a biliary atresia susceptibility gene. *Gastroenterology*, **2013**, 144, 1107-1115.e3	13.3	87
93	Developmental dysplasia of the hip: linkage mapping and whole exome sequencing identify a shared variant in CX3CR1 in all affected members of a large multigeneration family. *Journal of Bone and Mineral Research*, **2013**, 28, 2540-9	6.3	36
92	Replication of GWAS-identified neuroblastoma risk loci strengthens the role of BARD1 and affirms the cumulative effect of genetic variations on disease susceptibility. *Carcinogenesis*, **2013**, 34, 605-11	4.6	82
91	Computerized neurocognitive profile in young people with 22q11.2 deletion syndrome compared to youths with schizophrenia and at-risk for psychosis. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2012**, 159B, 87-93	3.5	31
90	Overt cleft palate phenotype and TBX1 genotype correlations in velo-cardio-facial/DiGeorge/22q11.2 deletion syndrome patients. *American Journal of Medical Genetics, Part A*, **2012**, 158A, 2781-7	2.5	19
89	Common variation at 6q16 within HACE1 and LIN28B influences susceptibility to neuroblastoma. *Nature Genetics*, **2012**, 44, 1126-30	36.3	198
88	Genetic mapping of quantitative trait loci for disease-related phenotypes. *Methods in Molecular Biology*, **2012**, 871, 281-311	1.4	2
87	Replication of neuroblastoma SNP association at the BARD1 locus in African-Americans. *Cancer Epidemiology Biomarkers and Prevention*, **2012**, 21, 658-63	4	48
86	Common variation at BARD1 results in the expression of an oncogenic isoform that influences neuroblastoma susceptibility and oncogenicity. *Cancer Research*, **2012**, 72, 2068-78	10.1	75
85	Genetic modifiers of β-thalassemia and clinical severity as assessed by age at first transfusion. *Haematologica*, **2012**, 97, 989-93	6.6	51
84	Genome-wide copy number variation study associates metabotropic glutamate receptor gene networks with attention deficit hyperactivity disorder. *Nature Genetics*, **2011**, 44, 78-84	36.3	279

83	Phenotype restricted genome-wide association study using a gene-centric approach identifies three low-risk neuroblastoma susceptibility Loci. *PLoS Genetics*, **2011**, 7, e1002026	6	123
82	Integrative genomics identifies LMO1 as a neuroblastoma oncogene. *Nature*, **2011**, 469, 216-20	50.4	231
81	Genotype and cardiovascular phenotype correlations with TBX1 in 1,022 velo-cardio-facial/DiGeorge/22q11.2 deletion syndrome patients. *Human Mutation*, **2011**, 32, 1278-89	4.7	48
80	Genome-wide linkage analysis to identify genetic modifiers of ALK mutation penetrance in familial neuroblastoma. *Human Heredity*, **2011**, 71, 135-9	1.1	23
79	Next-generation linkage analysis. *Human Heredity*, **2011**, 72, 227	1.1	4
78	Variants of DENND1B associated with asthma in children. *New England Journal of Medicine*, **2010**, 362, 36-44	59.2	261
77	The Otto Aufranc Award: Identification of a 4 Mb region on chromosome 17q21 linked to developmental dysplasia of the hip in one 18-member, multigeneration family. *Clinical Orthopaedics and Related Research*, **2010**, 468, 337-44	2.2	39
76	Analysis of GWAS top hits in ADHD suggests association to two polymorphisms located in genes expressed in the cerebellum. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2010**, 153B, 1127-33	3.5	17
75	Comparative analysis of different approaches for dealing with candidate regions in the context of a genome-wide association study. *BMC Proceedings*, **2009**, 3 Suppl 7, S93	2.3	4
74	Nocturnal enuresis: a suggestive endophenotype marker for a subgroup of inattentive attention-deficit/hyperactivity disorder. *Journal of Pediatrics*, **2009**, 155, 239-44.e5	3.6	41
73	Copy number variation at 1q21.1 associated with neuroblastoma. *Nature*, **2009**, 459, 987-91	50.4	285
72	Common variations in BARD1 influence susceptibility to high-risk neuroblastoma. *Nature Genetics*, **2009**, 41, 718-23	36.3	226
71	Candidate gene analysis in an on-going genome-wide association study of attention-deficit hyperactivity disorder: suggestive association signals in ADRA1A. *Psychiatric Genetics*, **2009**, 19, 134-41	2.9	27
70	Identification of ALK as a major familial neuroblastoma predisposition gene. *Nature*, **2008**, 455, 930-5	50.4	960
69	Association of a polymorphic variant of the adiponectin gene with insulin resistance in african americans. *Clinical and Translational Science*, **2008**, 1, 194-9	4.9	6
68	Estimates of genetic and environmental contribution to 43 quantitative traits support sharing of a homogeneous environment in an isolated population from South Tyrol, Italy. *Human Heredity*, **2008**, 65, 175-82	1.1	28
67	Advances in family-based association analysis. Introduction. *Human Heredity*, **2008**, 66, 65-6	1.1	3
66	A novel susceptibility locus for type 1 diabetes on Chr12q13 identified by a genome-wide association study. *Diabetes*, **2008**, 57, 1143-6	0.9	118

65	Chromosome 6p22 locus associated with clinically aggressive neuroblastoma. *New England Journal of Medicine*, **2008**, 358, 2585-93	59.2	224
64	Variants of ST8SIA1 are associated with risk of developing multiple sclerosis. *PLoS ONE*, **2008**, 3, e2653	3.7	10
63	Genetic loci linked to type 1 diabetes and multiple sclerosis families in Sardinia. *BMC Medical Genetics*, **2008**, 9, 3	2.1	18
62	Association of variants of the interleukin-23 receptor gene with susceptibility to pediatric Crohn's disease. *Clinical Gastroenterology and Hepatology*, **2007**, 5, 972-6	6.9	53
61	A novel locus for syndromic chronic idiopathic intestinal pseudo-obstruction maps to chromosome 8q23-q24. *European Journal of Human Genetics*, **2007**, 15, 889-97	5.3	24
60	A genome-wide association study identifies KIAA0350 as a type 1 diabetes gene. *Nature*, **2007**, 448, 591-4	30.4	424
59	ADHD genetics: 2007 update. *Current Psychiatry Reports*, **2007**, 9, 434-9	9.1	31
58	Association of the T300A non-synonymous variant of the ATG16L1 gene with susceptibility to paediatric Crohn's disease. *Gut*, **2007**, 56, 1171-3	19.2	53
57	Genetic predisposition to familial neuroblastoma: identification of two novel genomic regions at 2p and 12p. *Human Heredity*, **2007**, 63, 205-11	1.1	31
56	IRAK-M is involved in the pathogenesis of early-onset persistent asthma. *American Journal of Human Genetics*, **2007**, 80, 1103-14	11	125
55	Meta-analysis of genome-wide scans provides evidence for sex- and site-specific regulation of bone mass. *Journal of Bone and Mineral Research*, **2007**, 22, 173-183	6.3	128
54	Genetic linkage study of high-grade myopia in a Hutterite population from South Dakota. *Molecular Vision*, **2007**, 13, 229-36	2.3	57
53	Contribution of IBD5 locus to clinical features of IBD patients. *American Journal of Gastroenterology* , **2006**, 101, 318-25	0.7	24
52	Ophthalmologic findings in Cornelia de Lange syndrome: a genotype-phenotype correlation study. *JAMA Ophthalmology*, **2006**, 124, 552-7		21
51	Statistical tools for linkage analysis and genetic association studies. *Expert Review of Molecular Diagnostics*, **2005**, 5, 781-96	3.8	11
50	Univariate and bivariate variance component linkage analysis of a whole-genome scan for loci contributing to bone mineral density. *European Journal of Human Genetics*, **2005**, 13, 781-8	5.3	37
49	A common haplotype at the 5'end of the RET proto-oncogene, overrepresented in Hirschsprung patients, is associated with reduced gene expression. *Human Mutation*, **2005**, 25, 189-95	4.7	44
48	Precocious sister chromatid separation (PSCS) in Cornelia de Lange syndrome. *American Journal of Medical Genetics, Part A*, **2005**, 138, 27-31	2.5	72

47	Identification of a novel locus on 2q for autosomal dominant high-grade myopia. *Investigative Ophthalmology and Visual Science*, **2005**, 46, 2300-7		100
46	Heterogeneity in the magnitude of the insulin gene effect on HLA risk in type 1 diabetes. *Diabetes*, **2004**, 53, 3286-91	0.9	24
45	A genotype-phenotype correlation with gender-effect for hearing impairment caused by TECTA mutations. *Cellular Physiology and Biochemistry*, **2004**, 14, 369-76	3.9	46
44	Cornelia de Lange syndrome is caused by mutations in NIPBL, the human homolog of Drosophila melanogaster Nipped-B. *Nature Genetics*, **2004**, 36, 631-5	36.3	560
43	NIPBL mutational analysis in 120 individuals with Cornelia de Lange syndrome and evaluation of genotype-phenotype correlations. *American Journal of Human Genetics*, **2004**, 75, 610-23	11	245
42	CARD15 genotyping in inflammatory bowel disease patients by multiplex pyrosequencing. *Clinical Chemistry*, **2003**, 49, 1675-9	5.5	29
41	New locus for autosomal dominant high myopia maps to the long arm of chromosome 17. *Investigative Ophthalmology and Visual Science*, **2003**, 44, 1830-6		149
40	Linkage analysis in families with recurrent neuroblastoma. *Annals of the New York Academy of Sciences*, **2002**, 963, 74-84	6.5	16
39	Weak linkage at 4p16 to predisposition for human neuroblastoma. *Oncogene*, **2002**, 21, 8356-60	9.2	34
38	Second family with hearing impairment linked to 19q13 and refined DFNA4 localisation. *European Journal of Human Genetics*, **2002**, 10, 95-9	5.3	11
37	Linkage analysis conditional on HLA status in a large North American pedigree supports the presence of a multiple sclerosis susceptibility locus on chromosome 12p12. *Human Molecular Genetics*, **2002**, 11, 295-300	5.6	27
36	A rare haplotype of the RET proto-oncogene is a risk-modifying allele in hirschsprung disease. *American Journal of Human Genetics*, **2002**, 71, 969-74	11	39
35	Comparison of sib pair-based approaches for identifying quantitative trait loci underlying asthma in the Busselton families. *Genetic Epidemiology*, **2001**, 21 Suppl 1, S198-203	2.6	3
34	Exclusion of linkage to the CDL1 gene region on chromosome 3q26.3 in some familial cases of Cornelia de Lange syndrome. *American Journal of Medical Genetics Part A*, **2001**, 101, 120-129		26
33	Novel X-linked mental retardation syndrome with short stature maps to Xq24. *American Journal of Medical Genetics Part A*, **2001**, 103, 1-8		12
32	Charcot-Marie-Tooth type 4B is caused by mutations in the gene encoding myotubularin-related protein-2. *Nature Genetics*, **2000**, 25, 17-9	36.3	419
31	A single-nucleotide polymorphic variant of the RET proto-oncogene is underrepresented in sporadic Hirschsprung disease. *European Journal of Human Genetics*, **2000**, 8, 721-4	5.3	42
30	Combined segregation and linkage analysis of inflammatory bowel disease in the IBD1 region using severity to characterise Crohn's disease and ulcerative colitis. On behalf of the GISC. *European Journal of Human Genetics*, **2000**, 8, 846-52	5.3	16

29	A refined physical and transcriptional map of the SPG9 locus on 10q23.3-q24.2. *European Journal of Human Genetics*, **2000**, 8, 777-82	5.3	15
28	Genetic refinement and physical mapping of the CMT4B gene on chromosome 11q22. *Genomics*, **2000**, 63, 271-8	4.3	17
27	Linkage study of early-onset obesity to leptin receptor gene in Italian children. *Nutrition Research*, **2000**, 20, 1059-1063	4	2
26	Genetic analysis in Italian families with inflammatory bowel disease supports linkage to the IBD1 locus--a GISC study. *European Journal of Human Genetics*, **1999**, 7, 567-73	5.3	73
25	Genetic mapping to 10q23.3-q24.2, in a large Italian pedigree, of a new syndrome showing bilateral cataracts, gastroesophageal reflux, and spastic paraparesis with amyotrophy. *American Journal of Human Genetics*, **1999**, 64, 586-93	11	97
24	Five families with arginine 519-cysteine mutation in COL2A1: evidence for three distinct founders. *Human Mutation*, **1998**, 12, 172-6	4.7	23
23	A common mutation in Sardinian autoimmune polyendocrinopathy-candidiasis-ectodermal dystrophy patients. *Human Genetics*, **1998**, 103, 428-34	6.3	192
22	Polymorphisms of alpha-adducin and salt sensitivity in patients with essential hypertension. *Lancet, The*, **1997**, 349, 1353-7	40	473
21	A linkage study of schizophrenia to markers within Xp11 near the MAOB gene. *Psychiatry Research*, **1997**, 70, 131-43	9.9	55
20	Exclusion of linkage between RET and neuronal intestinal dysplasia type B. *American Journal of Medical Genetics Part A*, **1996**, 62, 195-8		15
19	A common molecular basis for three inherited kidney stone diseases. *Nature*, **1996**, 379, 445-9	50.4	614
18	Gender-dependent disease severity in autosomal polycystic kidney disease of rats. *Kidney International*, **1995**, 48, 496-500	9.9	34
17	Analysis of linkage disequilibrium between different cystic fibrosis mutations and three intragenic microsatellites in the Italian population. *Human Mutation*, **1995**, 5, 23-7	4.7	9
16	Search for linkage to schizophrenia on the X and Y chromosomes. *American Journal of Medical Genetics Part A*, **1994**, 54, 113-21		60
15	Close linkage with the RET protooncogene and boundaries of deletion mutations in autosomal dominant Hirschsprung disease. *Human Molecular Genetics*, **1993**, 2, 1803-8	5.6	77
14	Mapping a gene for familial situs abnormalities to human chromosome Xq24-q27.1. *Nature Genetics* , **1993**, 5, 403-7	36.3	122
13	A nonsense mutation (R1158X) and a splicing mutation (3849 + 4A----G) in exon 19 of the cystic fibrosis transmembrane conductance regulator gene. *Genomics*, **1992**, 12, 417-8	4.3	21
12	Four new mutations of the CFTR gene (541delC, R347H, R352Q, E585X) detected by DGGE analysis in Italian CF patients, associated with different clinical phenotypes. *Human Mutation*, **1992**, 1, 314-9	4.7	39

11	Frequency of the delta F508 mutation in a sample of 175 Italian cystic fibrosis patients. *Human Genetics*, **1990**, 85, 400-2	6.3	9
10	Preliminary results on the frequency of the delta F508 mutation in cystic fibrosis patients from the USSR. *Human Genetics*, **1990**, 85, 423-4	6.3	
9	Genetic differences in cystic fibrosis patients with and without pancreatic insufficiency. An Italian collaborative study. *Human Genetics*, **1990**, 84, 435-8	6.3	10
8	Haplotype distribution and molecular defects at the phenylalanine hydroxylase locus in Italy. *Human Genetics*, **1990**, 86, 69-72	6.3	15
7	Prenatal diagnosis and linkage disequilibrium with cystic fibrosis for markers surrounding D7S8. *Human Genetics*, **1990**, 85, 275-8	6.3	2
6	Why is the cystic fibrosis gene so frequent?. *Human Genetics*, **1989**, 84, 1-5	6.3	84
5	N-myc oncogene amplification and catecholamine metabolism in patients with neuroblastoma. *Lancet, The*, **1987**, 2, 795	40	1
4	Frequency of consanguineous marriages among parents and grandparents of Down patients. *Human Genetics*, **1985**, 70, 256-8	6.3	12
3	Attention-deficit hyperactivity disorder168-182		
2	Impaired redox and protein homeostasis as risk factors and therapeutic targets in toxin-induced biliary atresia		1
1	Sequencing of over 100,000 individuals identifies multiple genes and rare variants associated with Crohns disease susceptibility		2