Christopher J Cardinale

List of Publications by Year
in descending order

22	1,223	687363 13	794594 19
papers	citations	h-index	g-index

23	23	23	2755
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Termination Factor Rho and Its Cofactors NusA and NusG Silence Foreign DNA in E. coli. Science, 2008, 320, 935-938.	12.6	266
2	Meta-analysis of shared genetic architecture across ten pediatric autoimmune diseases. Nature Medicine, 2015, 21, 1018-1027.	30.7	212
3	Large Sample Size, Wide Variant Spectrum, and Advanced Machine-Learning Technique Boost Risk Prediction for Inflammatory Bowel Disease. American Journal of Human Genetics, 2013, 92, 1008-1012.	6.2	162
4	An Allosteric Path to Transcription Termination. Molecular Cell, 2007, 28, 991-1001.	9.7	114
5	A NovelBHLHE41Variant is Associated with Short Sleep and Resistance to Sleep Deprivation in Humans. Sleep, 2014, 37, 1327-1336.	1.1	104
6	Whole-genome DNA/RNA sequencing identifies truncating mutations in RBCK1 in a novel Mendelian disease with neuromuscular and cardiac involvement. Genome Medicine, 2013, 5, 67.	8.2	87
7	Genetic sharing and heritability of paediatric age of onset autoimmune diseases. Nature Communications, 2015, 6, 8442.	12.8	58
8	Rare copy number variants in over 100,000 European ancestry subjects reveal multiple disease associations. Nature Communications, 2020, 11, 255.	12.8	48
9	The B Cell SH2/PH Domain-Containing Adaptor Bam32/DAPP1 Is Required for T Cell-Independent II Antigen Responses. Current Biology, 2003, 13, 1858-1866.	3.9	42
10	Extended haplotype association study in Crohnâ€™s disease identifies a novel, Ashkenazi Jewish-specific missense mutation in the NF-Î²B pathway gene, HEATR3. Genes and Immunity, 2013, 14, 310-316.	4.1	31
11	Impact of exome sequencing in inflammatory bowel disease. World Journal of Gastroenterology, 2013, 19, 6721.	3.3	20
12	CNV Analysis Associates AKNAD1 with Type-2 Diabetes in Jordan Subpopulations. Scientific Reports, 2015, 5, 13391.	3.3	18
13	Targeted resequencing identifies defective variants of decoy receptor 3 in pediatric-onset inflammatory bowel disease. Genes and Immunity, 2013, 14, 447-452.	4.1	16
14	Association of a rare NOTCH4 coding variant with systemic sclerosis: a family-based whole exome sequencing study. BMC Musculoskeletal Disorders, 2016, 17, 462.	1.9	12
15	Pathway-based Genome-wide Association Studies Reveal the Association Between Growth Factor Activity and Inflammatory Bowel Disease. Inflammatory Bowel Diseases, 2016, 22, 1540-1551.	1.9	8
16	Transcriptome Profiling of Human Ulcerative Colitis Mucosa Reveals Altered Expression of Pathways Enriched in Genetic Susceptibility Loci. PLoS ONE, 2014, 9, e96153.	2.5	8
17	Multiple Epistasis Interactions Within MHC Are Associated With Ulcerative Colitis. Frontiers in Genetics, 2019, 10, 257.	2.3	7
18	Regulation of Janus Kinase 2 by an Inflammatory Bowel Disease Causal Non-coding Single Nucleotide Polymorphism. Journal of Crohn's and Colitis, 2020, 14, 646-653.	1.3	5

#	Article	IF	Citations
19	A novel <scp>MBTPS2</scp> variant associated with <scp>BRESHECK</scp> syndrome impairs <scp>sterolâ€regulated</scp> transcription and the endoplasmic reticulum stress response. American Journal of Medical Genetics, Part A, 2022, 188, 463-472.	1.2	4
20	Genetics of Inflammatory Bowel Diseases. , 2017, , 3-14.		1
21	P027 DIFFERENTIAL BINDING OF RBPJ AND CUX1 TO IBD CAUSAL SNP RS1887428 MODIFIES EXPRESSION OF JAK2. Gastroenterology, 2018, 154, S14-S15.	1.3	0
22	Genetic Underpinnings of Asthma and Related Traits. , 2020, , 341-360.		0