# 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 | 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 |
| 2 | Genetic Underpinnings of Asthma and Related Traits. , 2020, , 341-360. | | 0 |
| 3 | Rare copy number variants in over 100,000 European ancestry subjects reveal multiple disease associations. Nature Communications, 2020, 11, 255. | 12.8 | 48 |
| 4 | 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 |
| 5 | Multiple Epistasis Interactions Within MHC Are Associated With Ulcerative Colitis. Frontiers in Genetics, 2019, 10, 257. | 2.3 | 7 |
| 6 | P027 DIFFERENTIAL BINDING OF RBPJ AND CUX1 TO IBD CAUSAL SNP RS1887428 MODIFIES EXPRESSION OF JAK2. Gastroenterology, 2018, 154, S14-S15. | 1.3 | 0 |
| 7 | Genetics of Inflammatory Bowel Diseases. , 2017, , 3-14. | | 1 |
| 8 | 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 |
| 9 | 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 |
| 10 | CNV Analysis Associates AKNAD1 with Type-2 Diabetes in Jordan Subpopulations. Scientific Reports, 2015, 5, 13391. | 3.3 | 18 |
| 11 | Genetic sharing and heritability of paediatric age of onset autoimmune diseases. Nature Communications, 2015, 6, 8442. | 12.8 | 58 |
| 12 | Meta-analysis of shared genetic architecture across ten pediatric autoimmune diseases. Nature Medicine, 2015, 21, 1018-1027. | 30.7 | 212 |
| 13 | A NovelBHLHE41Variant is Associated with Short Sleep and Resistance to Sleep Deprivation in Humans. Sleep, 2014, 37, 1327-1336. | 1.1 | 104 |
| 14 | 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 |
| 15 | 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 |
| 16 | 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 |
| 17 | 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 |
| 18 | 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 |

| # | Article | IF | Citations |
|---|---|---|---|
| 19 | Impact of exome sequencing in inflammatory bowel disease. World Journal of Gastroenterology, 2013, 19, 6721. | 3.3 | 20 |
| 20 | Termination Factor Rho and Its Cofactors NusA and NusG Silence Foreign DNA in E. coli. Science, 2008, 320, 935-938. | 12.6 | 266 |
| 21 | An Allosteric Path to Transcription Termination. Molecular Cell, 2007, 28, 991-1001. | 9.7 | 114 |
| 22 | 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 |