Alex J Cornish

List of Publications by Year
in descending order

25 papers	671 citations	567281 15 h-index	610901 24 g-index
28 all docs	28 docs citations	28 times ranked	1346 citing authors

#	ARTICLE	IF	CITATIONS
1	Modifiable pathways for colorectal cancer: a mendelian randomisation analysis. The Lancet Gastroenterology and Hepatology, 2020, 5, 55-62.	8.1	79
2	Whole-genome sequencing of multiple myeloma reveals oncogenic pathways are targeted somatically through multiple mechanisms. Leukemia, 2018, 32, 2459-2470.	7.2	68
3	Promoter capture Hi-C-based identification of recurrent noncoding mutations in colorectal cancer. Nature Genetics, 2018, 50, 1375-1380.	21.4	49
4	Genomic landscape of platinum resistant and sensitive testicular cancers. Nature Communications, 2020, 11, 2189.	12.8	43
5	Mutational processes contributing to the development of multiple myeloma. Blood Cancer Journal, 2019, 9, 60.	6.2	41
6	Mendelian randomisation: A powerful and inexpensive method for identifying and excluding non-genetic risk factors for colorectal cancer. Molecular Aspects of Medicine, 2019, 69, 41-47.	6.4	39
7	Impact of atopy on risk of glioma: a Mendelian randomisation study. BMC Medicine, 2018, 16, 42.	5.5	38
8	Signatures of TOP1 transcription-associated mutagenesis in cancer and germline. Nature, 2022, 602, 623-631.	27.8	38
9	Influence of obesity-related risk factors in the aetiology of glioma. British Journal of Cancer, 2018, 118, 1020-1027.	6.4	32
10	Search for multiple myeloma risk factors using Mendelian randomization. Blood Advances, 2020, 4, 2172-2179.	5.2	27
11	Mendelian randomisation study of the relationship between vitamin D and risk of glioma. Scientific Reports, 2018, 8, 2339.	3.3	23
12	Genome-wide association analysis identifies a meningioma risk locus at 11p15.5. Neuro-Oncology, 2018, 20, 1485-1493.	1.2	23
13	Phenome-wide association analysis of LDL-cholesterol lowering genetic variants in PCSK9. BMC Cardiovascular Disorders, 2019, 19, 240.	1.7	22
14	Mendelian randomization provides support for obesity as a risk factor for meningioma. Scientific Reports, 2019, 9, 309.	3.3	21
15	Relationship between genetically determined telomere length and glioma risk. Neuro-Oncology, 2022, 24, 171-181.	1.2	21
16	Identification of recurrent noncoding mutations in B-cell lymphoma using capture Hi-C. Blood Advances, 2019, 3, 21-32.	5.2	20
17	Lack of association between modifiable exposures and glioma risk: A Mendelian randomisation analysis. Neuro-Oncology, 2020, 22, 207-215.	1.2	19
18	An enhanced genetic model of relapsed IGH-translocated multiple myeloma evolutionary dynamics. Blood Cancer Journal, 2020, 10, 101.	6.2	11

#	Article	IF	Citations
19	Elongin C (<i>ELOC</i>/<i>TCEB1</i>)-associated von Hippel–Lindau disease. Human Molecular Genetics, 2022, 31, 2728-2737.	2.9	11
20	Searching for causal relationships of glioma: a phenome-wide Mendelian randomisation study. British Journal of Cancer, 2021, 124, 447-454.	6.4	9
21	Cancer drivers and clonal dynamics in acute lymphoblastic leukaemia subtypes. Blood Cancer Journal, 2021, 11, 177.	6.2	9
22	Frequency of pathogenic germline variants in cancer susceptibility genes in 1336 renal cell carcinoma cases. Human Molecular Genetics, 2022, 31, 3001-3011.	2.9	9
23	Impact of mitochondrial DNA mutations in multiple myeloma. Blood Cancer Journal, 2020, 10, 46.	6.2	8
24	Lack of an association between gallstone disease and bilirubin levels with risk of colorectal cancer: a Mendelian randomisation analysis. British Journal of Cancer, 2021, 124, 1169-1174.	6.4	6
25	Reference bias in the Illumina Isaac aligner. Bioinformatics, 2020, 36, 4671-4672.	4.1	5