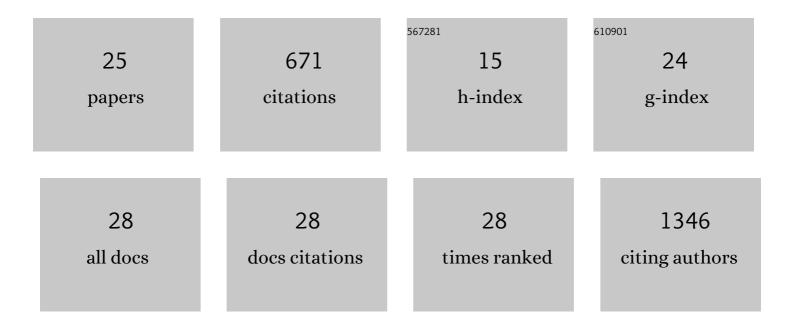
## Alex J Cornish

List of Publications by Year in descending order

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ALEY CODNICH

| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Relationship between genetically determined telomere length and glioma risk. Neuro-Oncology, 2022, 24, 171-181.   | 1.2  | 21        |
| 2  | Signatures of TOP1 transcription-associated mutagenesis in cancer and germline. Nature, 2022, 602, 623-631.   | 27.8 | 38        |
| 3  | Elongin C ( <i>ELOC</i> / <i>TCEB1</i> )-associated von Hippel–Lindau disease. Human Molecular Genetics,<br>2022, 31, 2728-2737.  | 2.9  | 11        |
| 4  | 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         |
| 5  | Searching for causal relationships of glioma: a phenome-wide Mendelian randomisation study. British<br>Journal of Cancer, 2021, 124, 447-454.   | 6.4  | 9         |
| 6  | Lack of an association between gallstone disease and bilirubin levels with risk of colorectal cancer: a<br>Mendelian randomisation analysis. British Journal of Cancer, 2021, 124, 1169-1174. | 6.4  | 6         |
| 7  | Cancer drivers and clonal dynamics in acute lymphoblastic leukaemia subtypes. Blood Cancer Journal, 2021, 11, 177.  | 6.2  | 9         |
| 8  | Lack of association between modifiable exposures and glioma risk: A Mendelian randomisation analysis. Neuro-Oncology, 2020, 22, 207-215.  | 1.2  | 19        |
| 9  | Modifiable pathways for colorectal cancer: a mendelian randomisation analysis. The Lancet<br>Gastroenterology and Hepatology, 2020, 5, 55-62.   | 8.1  | 79        |
| 10 | An enhanced genetic model of relapsed IGH-translocated multiple myeloma evolutionary dynamics.<br>Blood Cancer Journal, 2020, 10, 101.  | 6.2  | 11        |
| 11 | Impact of mitochondrial DNA mutations in multiple myeloma. Blood Cancer Journal, 2020, 10, 46.  | 6.2  | 8         |
| 12 | Search for multiple myeloma risk factors using Mendelian randomization. Blood Advances, 2020, 4,<br>2172-2179.  | 5.2  | 27        |
| 13 | Genomic landscape of platinum resistant and sensitive testicular cancers. Nature Communications, 2020, 11, 2189.  | 12.8 | 43        |
| 14 | Reference bias in the Illumina Isaac aligner. Bioinformatics, 2020, 36, 4671-4672.  | 4.1  | 5         |
| 15 | Mutational processes contributing to the development of multiple myeloma. Blood Cancer Journal, 2019, 9, 60.  | 6.2  | 41        |
| 16 | Phenome-wide association analysis of LDL-cholesterol lowering genetic variants in PCSK9. BMC<br>Cardiovascular Disorders, 2019, 19, 240.  | 1.7  | 22        |
| 17 | Mendelian randomization provides support for obesity as a risk factor for meningioma. Scientific<br>Reports, 2019, 9, 309.  | 3.3  | 21        |
| 18 | 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        |

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| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 19 | Identification of recurrent noncoding mutations in B-cell lymphoma using capture Hi-C. Blood Advances, 2019, 3, 21-32.                                      | 5.2  | 20        |
| 20 | Mendelian randomisation study of the relationship between vitamin D and risk of glioma. Scientific<br>Reports, 2018, 8, 2339.                               | 3.3  | 23        |
| 21 | Impact of atopy on risk of glioma: a Mendelian randomisation study. BMC Medicine, 2018, 16, 42.   | 5.5  | 38        |
| 22 | Influence of obesity-related risk factors in the aetiology of glioma. British Journal of Cancer, 2018, 118, 1020-1027.                                      | 6.4  | 32        |
| 23 | Whole-genome sequencing of multiple myeloma reveals oncogenic pathways are targeted somatically through multiple mechanisms. Leukemia, 2018, 32, 2459-2470. | 7.2  | 68        |
| 24 | Promoter capture Hi-C-based identification of recurrent noncoding mutations in colorectal cancer.<br>Nature Genetics, 2018, 50, 1375-1380.                  | 21.4 | 49        |
| 25 | Genome-wide association analysis identifies a meningioma risk locus at 11p15.5. Neuro-Oncology, 2018, 20, 1485-1493.  | 1.2  | 23        |