

# Charles P Fulco

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5550320/publications.pdf>

Version: 2024-02-01

10  
papers

2,202  
citations

1040056

9  
h-index

1281871

11  
g-index

17  
all docs

17  
docs citations

17  
times ranked

4016  
citing authors

#	ARTICLE	IF	CITATIONS
1	Activity-by-contact model of enhancer-promoter regulation from thousands of CRISPR perturbations. <i>Nature Genetics</i> , 2019, 51, 1664-1669.	21.4	631
2	Systematic mapping of functional enhancer-promoter connections with CRISPR interference. <i>Science</i> , 2016, 354, 769-773.	12.6	512
3	Inherited causes of clonal haematopoiesis in 97,691 whole genomes. <i>Nature</i> , 2020, 586, 763-768.	27.8	376
4	Genome-wide enhancer maps link risk variants to disease genes. <i>Nature</i> , 2021, 593, 238-243.	27.8	332
5	Compatibility rules of human enhancer and promoter sequences. <i>Nature</i> , 2022, 607, 176-184.	27.8	67
6	Combining SNP-to-gene linking strategies to identify disease genes and assess disease omnigenicity. <i>Nature Genetics</i> , 2022, 54, 827-836.	21.4	61
7	Prioritizing disease and trait causal variants at the TNFAIP3 locus using functional and genomic features. <i>Nature Communications</i> , 2020, 11, 1237.	12.8	38
8	CRISPR-SURF: discovering regulatory elements by deconvolution of CRISPR tiling screen data. <i>Nature Methods</i> , 2018, 15, 992-993.	19.0	33
9	HyPR-seq: Single-cell quantification of chosen RNAs via hybridization and sequencing of DNA probes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 33404-33413.	7.1	21
10	Systematic identification of genomic elements that regulate <i>FCGR2A</i> expression and harbor variants linked with autoimmune disease. <i>Human Molecular Genetics</i> , 2022, 31, 1946-1961.	2.9	7