

# Joelle Mbatchou

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3470551/publications.pdf>

Version: 2024-02-01

6  
papers

1,204  
citations

1478505

6  
h-index

1872680

6  
g-index

9  
all docs

9  
docs citations

9  
times ranked

1715  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide analysis provides genetic evidence that ACE2 influences COVID-19 risk and yields risk scores associated with severe disease. <i>Nature Genetics</i> , 2022, 54, 382-392.	21.4	97
2	Computationally efficient whole-genome regression for quantitative and binary traits. <i>Nature Genetics</i> , 2021, 53, 1097-1103.	21.4	457
3	Pan-ancestry exome-wide association analyses of COVID-19 outcomes in 586,157 individuals. <i>American Journal of Human Genetics</i> , 2021, 108, 1350-1355.	6.2	72
4	Sequencing of 640,000 exomes identifies <i>GPR75</i> variants associated with protection from obesity. <i>Science</i> , 2021, 373, .	12.6	130
5	Exome sequencing and analysis of 454,787 UK Biobank participants. <i>Nature</i> , 2021, 599, 628-634.	27.8	377
6	Retrospective Association Analysis of Binary Traits: Overcoming Some Limitations of the Additive Polygenic Model. <i>Human Heredity</i> , 2015, 80, 187-195.	0.8	13