

# Alejandro Chavez

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/299786/publications.pdf>

Version: 2024-02-01

25  
papers

4,265  
citations

361413

20  
h-index

552781

26  
g-index

30  
all docs

30  
docs citations

30  
times ranked

6366  
citing authors

#	ARTICLE	IF	CITATIONS
1	Highly efficient Cas9-mediated transcriptional programming. <i>Nature Methods</i> , 2015, 12, 326-328.	19.0	1,245
2	Comparison of Cas9 activators in multiple species. <i>Nature Methods</i> , 2016, 13, 563-567.	19.0	438
3	Programmable transcriptional repression in mycobacteria using an orthogonal CRISPR interference platform. <i>Nature Microbiology</i> , 2017, 2, 16274.	13.3	368
4	An enhanced CRISPR repressor for targeted mammalian gene regulation. <i>Nature Methods</i> , 2018, 15, 611-616.	19.0	361
5	Safeguarding CRISPR-Cas9 gene drives in yeast. <i>Nature Biotechnology</i> , 2015, 33, 1250-1255.	17.5	291
6	Cas9 gRNA engineering for genome editing, activation and repression. <i>Nature Methods</i> , 2015, 12, 1051-1054.	19.0	272
7	An Integrated Genome-wide CRISPRa Approach to Functionalize lncRNAs in Drug Resistance. <i>Cell</i> , 2018, 173, 649-664.e20.	28.9	238
8	sgRNA Scorer 2.0: A Species-Independent Model To Predict CRISPR/Cas9 Activity. <i>ACS Synthetic Biology</i> , 2017, 6, 902-904.	3.8	189
9	Daisy-chain gene drives for the alteration of local populations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 8275-8282.	7.1	154
10	A CRISPR-Cas9-based gene drive platform for genetic interaction analysis in <i>Candida albicans</i> . <i>Nature Microbiology</i> , 2018, 3, 73-82.	13.3	135
11	Enhanced Bacterial Immunity and Mammalian Genome Editing via RNA-Polymerase-Mediated Dislodging of Cas9 from Double-Strand DNA Breaks. <i>Molecular Cell</i> , 2018, 71, 42-55.e8.	9.7	112
12	CRISPR-based genomic tools for the manipulation of genetically intractable microorganisms. <i>Nature Reviews Microbiology</i> , 2018, 16, 333-339.	28.6	88
13	Lead compounds for the development of SARS-CoV-2 3CL protease inhibitors. <i>Nature Communications</i> , 2021, 12, 2016.	12.8	65
14	A CRISPR Interference Platform for Efficient Genetic Repression in <i>Candida albicans</i> . <i>MSphere</i> , 2019, 4, .	2.9	49
15	Development of optimized drug-like small molecule inhibitors of the SARS-CoV-2 3CL protease for treatment of COVID-19. <i>Nature Communications</i> , 2022, 13, 1891.	12.8	45
16	Reduced apoptosis in Chinese hamster ovary cells via optimized CRISPR interference. <i>Biotechnology and Bioengineering</i> , 2019, 116, 1813-1819.	3.3	39
17	An antibody class with a common CDRH3 motif broadly neutralizes sarbecoviruses. <i>Science Translational Medicine</i> , 2022, 14, eabn6859.	12.4	31
18	Precise Cas9 targeting enables genomic mutation prevention. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 3669-3673.	7.1	28

#	ARTICLE	IF	CITATIONS
19	Synthetic immunomodulation with a CRISPR super-repressor in vivo. <i>Nature Cell Biology</i> , 2020, 22, 1143-1154.	10.3	27
20	Inhibitors of Coronavirus 3CL Proteases Protect Cells from Protease-Mediated Cytotoxicity. <i>Journal of Virology</i> , 2021, 95, e0237420.	3.4	27
21	Design, execution, and analysis of CRISPR-Cas9-based deletions and genetic interaction networks in the fungal pathogen <i>Candida albicans</i> . <i>Nature Protocols</i> , 2019, 14, 955-975.	12.0	25
22	Enabling multiplexed testing of pooled donor cells through whole-genome sequencing. <i>Genome Medicine</i> , 2018, 10, 31.	8.2	10
23	CRISPR Guide RNA Cloning for Mammalian Systems. <i>Journal of Visualized Experiments</i> , 2018, , .	0.3	6
24	Quantifying protein abundance on single cells using split-pool sequencing on DNA-barcoded antibodies for diagnostic applications. <i>Scientific Reports</i> , 2022, 12, 884.	3.3	3
25	Clonal Hematopoiesis with Somatic Mutations Is a Common, Age-Related Condition Associated with Adverse Outcomes. <i>Blood</i> , 2014, 124, 840-840.	1.4	1