

Jonathan S Gootenberg

List of Publications by Year in descending order

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Version: 2024-02-01

44
papers

30,024
citations

126708

33
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205818

48
g-index

62
all docs

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docs citations

62
times ranked

23701
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Structure and engineering of the type III-E CRISPR-Cas7-11 effector complex. <i>Cell</i> , 2022, 185, 2324-2337.e16. | 13.5 | 51 |
| 2 | CRISPR diagnostics. <i>Science</i> , 2021, 372, 914-915. | 6.0 | 52 |
| 3 | A global metagenomic map of urban microbiomes and antimicrobial resistance. <i>Cell</i> , 2021, 184, 3376-3393.e17. | 13.5 | 164 |
| 4 | CRISPR-based diagnostics. <i>Nature Biomedical Engineering</i> , 2021, 5, 643-656. | 11.6 | 492 |
| 5 | Programmable RNA targeting with the single-protein CRISPR effector Cas7-11. <i>Nature</i> , 2021, 597, 720-725. | 13.7 | 155 |
| 6 | A Survey of Genome Editing Activity for 16 Cas12a Orthologs. <i>Keio Journal of Medicine</i> , 2020, 69, 59-65. | 0.5 | 41 |
| 7 | Clinical validation of a Cas13-based assay for the detection of SARS-CoV-2 RNA. <i>Nature Biomedical Engineering</i> , 2020, 4, 1140-1149. | 11.6 | 442 |
| 8 | Detection of SARS-CoV-2 with SHERLOCK One-Pot Testing. <i>New England Journal of Medicine</i> , 2020, 383, 1492-1494. | 13.9 | 506 |
| 9 | Rapid and accurate species identification for ecological studies and monitoring using CRISPR-based SHERLOCK. <i>Molecular Ecology Resources</i> , 2020, 20, 961-970. | 2.2 | 35 |
| 10 | Rapid SARS-CoV-2 testing in primary material based on a novel multiplex RT-LAMP assay. <i>PLoS ONE</i> , 2020, 15, e0238612. | 1.1 | 58 |
| 11 | CRISPR Tools for Systematic Studies of RNA Regulation. <i>Cold Spring Harbor Perspectives in Biology</i> , 2019, 11, a035386. | 2.3 | 22 |
| 12 | A cytosine deaminase for programmable single-base RNA editing. <i>Science</i> , 2019, 365, 382-386. | 6.0 | 322 |
| 13 | Nucleic Acid Detection of Plant Genes Using CRISPR-Cas13. <i>CRISPR Journal</i> , 2019, 2, 165-171. | 1.4 | 92 |
| 14 | Programmable Inhibition and Detection of RNA Viruses Using Cas13. <i>Molecular Cell</i> , 2019, 76, 826-837.e11. | 4.5 | 286 |
| 15 | SHERLOCK: nucleic acid detection with CRISPR nucleases. <i>Nature Protocols</i> , 2019, 14, 2986-3012. | 5.5 | 851 |
| 16 | Chipping in on Diagnostics. <i>CRISPR Journal</i> , 2019, 2, 69-71. | 1.4 | 4 |
| 17 | Structural basis for the promiscuous PAM recognition by <i>Corynebacterium diphtheriae</i> Cas9. <i>Nature Communications</i> , 2019, 10, 1968. | 5.8 | 33 |
| 18 | High-Resolution Structure of Cas13b and Biochemical Characterization of RNA Targeting and Cleavage. <i>Cell Reports</i> , 2019, 26, 3741-3751.e5. | 2.9 | 102 |

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|----|--|------|-----------|
| 19 | S15.2â€¦Crispr diagnostics: expanding the nucleic acid detection toolbox by harnessing microbial diversity. , 2019, , . | | 0 |
| 20 | Multiplexed and portable nucleic acid detection platform with Cas13, Cas12a, and Csm6. <i>Science</i> , 2018, 360, 439-444. | 6.0 | 1,649 |
| 21 | Field-deployable viral diagnostics using CRISPR-Cas13. <i>Science</i> , 2018, 360, 444-448. | 6.0 | 982 |
| 22 | Engineered CRISPR-Cas9 nuclease with expanded targeting space. <i>Science</i> , 2018, 361, 1259-1262. | 6.0 | 783 |
| 23 | Pairwise library screen systematically interrogates <i>Staphylococcus aureus</i> Cas9 specificity in human cells. <i>Nature Communications</i> , 2018, 9, 2962. | 5.8 | 32 |
| 24 | Diversity and evolution of class 2 CRISPRâ€“Cas systems. <i>Nature Reviews Microbiology</i> , 2017, 15, 169-182. | 13.6 | 792 |
| 25 | Nucleic acid detection with CRISPR-Cas13a/C2c2. <i>Science</i> , 2017, 356, 438-442. | 6.0 | 2,275 |
| 26 | Multiplex gene editing by CRISPRâ€“Cpf1 using a single crRNA array. <i>Nature Biotechnology</i> , 2017, 35, 31-34. | 9.4 | 736 |
| 27 | Genome-scale CRISPR-Cas9 knockout and transcriptional activation screening. <i>Nature Protocols</i> , 2017, 12, 828-863. | 5.5 | 858 |
| 28 | Barcode extension for analysis and reconstruction of structures. <i>Nature Communications</i> , 2017, 8, 14698. | 5.8 | 17 |
| 29 | Crystal Structure of the Minimal Cas9 from <i>Campylobacter jejuni</i> Reveals the Molecular Diversity in the CRISPR-Cas9 Systems. <i>Molecular Cell</i> , 2017, 65, 1109-1121.e3. | 4.5 | 145 |
| 30 | Cas13b Is a Type VI-B CRISPR-Associated RNA-Guided RNase Differentially Regulated by Accessory Proteins Csx27 and Csx28. <i>Molecular Cell</i> , 2017, 65, 618-630.e7. | 4.5 | 445 |
| 31 | RNA editing with CRISPR-Cas13. <i>Science</i> , 2017, 358, 1019-1027. | 6.0 | 1,301 |
| 32 | RNA targeting with CRISPRâ€“Cas13. <i>Nature</i> , 2017, 550, 280-284. | 13.7 | 1,442 |
| 33 | Genome-scale activation screen identifies a lncRNA locus regulating a gene neighbourhood. <i>Nature</i> , 2017, 548, 343-346. | 13.7 | 336 |
| 34 | C2c2 is a single-component programmable RNA-guided RNA-targeting CRISPR effector. <i>Science</i> , 2016, 353, aaf5573. | 6.0 | 1,647 |
| 35 | Structure and Engineering of <i>Francisella novicida</i> Cas9. <i>Cell</i> , 2016, 164, 950-961. | 13.5 | 296 |
| 36 | Engineered bromodomains to explore the acetylproteome. <i>Proteomics</i> , 2015, 15, 1470-1475. | 1.3 | 10 |

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|----|--|------|-----------|
| 37 | In vivo genome editing using <i>Staphylococcus aureus</i> Cas9. <i>Nature</i> , 2015, 520, 186-191. | 13.7 | 2,237 |
| 38 | Orthogonal gene knockout and activation with a catalytically active Cas9 nuclease. <i>Nature Biotechnology</i> , 2015, 33, 1159-1161. | 9.4 | 231 |
| 39 | Cpf1 Is a Single RNA-Guided Endonuclease of a Class 2 CRISPR-Cas System. <i>Cell</i> , 2015, 163, 759-771. | 13.5 | 3,558 |
| 40 | Discovery and Functional Characterization of Diverse Class 2 CRISPR-Cas Systems. <i>Molecular Cell</i> , 2015, 60, 385-397. | 4.5 | 971 |
| 41 | Genome-scale transcriptional activation by an engineered CRISPR-Cas9 complex. <i>Nature</i> , 2015, 517, 583-588. | 13.7 | 2,272 |
| 42 | Double Nicking by RNA-Guided CRISPR Cas9 for Enhanced Genome Editing Specificity. <i>Cell</i> , 2013, 154, 1380-1389. | 13.5 | 2,862 |
| 43 | Double Nicking by RNA-Guided CRISPR Cas9 for Enhanced Genome Editing Specificity. <i>Cell</i> , 2013, 155, 479-480. | 13.5 | 45 |
| 44 | Abstract 1978: The poly-SUMO protein specific E3 ubiquitin ligase RNF4 is induced in multiple myeloma and reduces bortezomib-induced cell killing. , 2010, , . | | 0 |