## Jonathan J Park

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/424720/publications.pdf

Version: 2024-02-01

1040056 1281871 1,483 11 9 11 citations h-index g-index papers 12 12 12 4274 docs citations times ranked citing authors all docs

| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Comprehensive functional genomic resource and integrative model for the human brain. Science, 2018, 362, .   | 12.6 | 618       |
| 2  | Systematic Immunotherapy Target Discovery Using Genome-Scale InÂVivo CRISPR Screens in CD8ÂT Cells. Cell, 2019, 178, 1189-1204.e23.  | 28.9 | 189       |
| 3  | Nonstructural Protein 1 of SARS-CoV-2 Is a Potent Pathogenicity Factor Redirecting Host Protein Synthesis Machinery toward Viral RNA. Molecular Cell, 2020, 80, 1055-1066.e6.                        | 9.7  | 152       |
| 4  | In vivo CRISPR screening in CD8 T cells with AAV–Sleeping Beauty hybrid vectors identifies membrane targets for improving immunotherapy for glioblastoma. Nature Biotechnology, 2019, 37, 1302-1313. | 17.5 | 123       |
| 5  | exceRpt: A Comprehensive Analytic Platform for Extracellular RNA Profiling. Cell Systems, 2019, 8, 352-357.e3.   | 6.2  | 118       |
| 6  | One-step generation of modular CAR-T cells with AAV–Cpf1. Nature Methods, 2019, 16, 247-254.   | 19.0 | 101       |
| 7  | Multiplexed activation of endogenous genes by CRISPRa elicits potent antitumor immunity. Nature Immunology, 2019, 20, 1494-1505.   | 14.5 | 83        |
| 8  | A genome-scale gain-of-function CRISPR screen in CD8 TÂcells identifies proline metabolism as a means to enhance CAR-T therapy. Cell Metabolism, 2022, 34, 595-614.e14.                              | 16.2 | 70        |
| 9  | Convergent Identification and Interrogation of Tumor-Intrinsic Factors that Modulate Cancer Immunity InÂVivo. Cell Systems, 2019, 8, 136-151.e7.   | 6.2  | 14        |
| 10 | Variant-specific vaccination induces systems immune responses and potent inÂvivo protection against SARS-CoV-2. Cell Reports Medicine, 2022, 3, 100634.  | 6.5  | 10        |
| 11 | Metaviromic identification of discriminative genomic features in SARS-CoV-2 using machine learning. Patterns, 2022, 3, 100407.   | 5.9  | 4         |