

Emmanouil Metzakopian

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8566871/publications.pdf>

Version: 2024-02-01

14
papers

860
citations

840776

11
h-index

996975

15
g-index

18
all docs

18
docs citations

18
times ranked

1821
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Foxa1 and Foxa2 function both upstream of and cooperatively with Lmx1a and Lmx1b in a feedforward loop promoting mesodiencephalic dopaminergic neuron development. <i>Developmental Biology</i> , 2009, 333, 386-396. | 2.0 | 139 |
| 2 | UTX-mediated enhancer and chromatin remodeling suppresses myeloid leukemogenesis through noncatalytic inverse regulation of ETS and GATA programs. <i>Nature Genetics</i> , 2018, 50, 883-894. | 21.4 | 117 |
| 3 | Felodipine induces autophagy in mouse brains with pharmacokinetics amenable to repurposing. <i>Nature Communications</i> , 2019, 10, 1817. | 12.8 | 88 |
| 4 | Single-Cell Transcriptomics of Parkinson's Disease Human In Vitro Models Reveals Dopamine Neuron-Specific Stress Responses. <i>Cell Reports</i> , 2020, 33, 108263. | 6.4 | 79 |
| 5 | A conditional piggyBac transposition system for genetic screening in mice identifies oncogenic networks in pancreatic cancer. <i>Nature Genetics</i> , 2015, 47, 47-56. | 21.4 | 77 |
| 6 | Foxa1 and Foxa2 Are Required for the Maintenance of Dopaminergic Properties in Ventral Midbrain Neurons at Late Embryonic Stages. <i>Journal of Neuroscience</i> , 2013, 33, 8022-8034. | 3.6 | 73 |
| 7 | ER-mitochondria contact sites in neurodegeneration: genetic screening approaches to investigate novel disease mechanisms. <i>Cell Death and Differentiation</i> , 2021, 28, 1804-1821. | 11.2 | 70 |
| 8 | A single-copy Sleeping Beauty transposon mutagenesis screen identifies new PTEN-cooperating tumor suppressor genes. <i>Nature Genetics</i> , 2017, 49, 730-741. | 21.4 | 53 |
| 9 | Foxa1 and Foxa2 positively and negatively regulate Shh signalling to specify ventral midbrain progenitor identity. <i>Mechanisms of Development</i> , 2011, 128, 90-103. | 1.7 | 50 |
| 10 | Genome-Scale CRISPRa Screen Identifies Novel Factors for Cellular Reprogramming. <i>Stem Cell Reports</i> , 2019, 12, 757-771. | 4.8 | 45 |
| 11 | Enhancing the genome editing toolbox: genome wide CRISPR arrayed libraries. <i>Scientific Reports</i> , 2017, 7, 2244. | 3.3 | 35 |
| 12 | The transcription factor BCL11A defines distinct subsets of midbrain dopaminergic neurons. <i>Cell Reports</i> , 2021, 36, 109697. | 6.4 | 14 |
| 13 | Development and Application of High-Throughput Single Cell Lipid Profiling: A Study of SNCA-A53T Human Dopamine Neurons. <i>iScience</i> , 2020, 23, 101703. | 4.1 | 13 |
| 14 | Using Reactome to build an autophagy mechanism knowledgebase. <i>Autophagy</i> , 2021, 17, 1543-1554. | 9.1 | 5 |