Felix Jonas

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1060092/publications.pdf

Version: 2024-02-01

| | | 1478505 | 1372567 | |
|----------|----------------|--------------|----------------|--|
| 10 | 173 | 6 | 10 | |
| papers | citations | h-index | g-index | |
| | | | | |
| | | | | |
| 15 | 15 | 15 | 267 | |
| all docs | docs citations | times ranked | citing authors | |
| | | | | |

| # | Article | IF | CITATIONS |
|----|--|--------------|-----------|
| 1 | Rethinking cell growth models. FEMS Yeast Research, 2016, 16, fow081. | 2.3 | 42 |
| 2 | Iron affects Ire1 clustering propensity and the amplitude of endoplasmic reticulum stress signaling. Journal of Cell Science, 2017, 130, 3222-3233. | 2.0 | 35 |
| 3 | Evolution of binding preferences among whole-genome duplicated transcription factors. ELife, 2022, 11, . | 6.0 | 34 |
| 4 | Measurement of histone replacement dynamics with genetically encoded exchange timers in yeast. Nature Biotechnology, 2021, 39, 1434-1443. | 17.5 | 15 |
| 5 | A regulatory phosphorylation site on Mec1 controls chromatin occupancy of RNA polymerases during replication stress. EMBO Journal, 2021, 40, e108439. | 7.8 | 14 |
| 6 | A Visual Framework for Classifying Determinants of Cell Size. Cell Reports, 2018, 25, 3519-3529.e2. | 6.4 | 11 |
| 7 | Rtt109 slows replication speed by histone N-terminal acetylation. Genome Research, 2021, 31, 426-435. | 5.5 | 7 |
| 8 | Accumulation of <i>cis</i> - and <i>trans</i> - regulatory variations is associated with phenotypic divergence of a complex trait between yeast species. G3: Genes, Genomes, Genetics, 2021, 11, . | 1.8 | 6 |
| 9 | Generation and timing of graded responses to morphogen gradients. Development (Cambridge), 2021, 148, . | 2.5 | 5 |
| 10 | Rtt109 promotes nucleosome replacement ahead of the replication fork. Genome Research, 2022, 32, 1089-1098. | 5 . 5 | 1 |