

Mari Mito

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8409005/publications.pdf>

Version: 2024-02-01

13
papers

972
citations

932766

10
h-index

1125271

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16
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docs citations

16
times ranked

1439
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Structural, super-resolution microscopy analysis of paraspeckle nuclear body organization. <i>Journal of Cell Biology</i> , 2016, 214, 817-830. | 2.3 | 262 |
| 2 | The lncRNA <i>Neat1</i> is required for corpus luteum formation and the establishment of pregnancy in a subpopulation of mice. <i>Development (Cambridge)</i> , 2014, 141, 4618-4627. | 1.2 | 229 |
| 3 | Complete chemical structures of human mitochondrial tRNAs. <i>Nature Communications</i> , 2020, 11, 4269. | 5.8 | 144 |
| 4 | The Translation Inhibitor Rocaglamide Targets a Bimolecular Cavity between eIF4A and Polypurine RNA. <i>Molecular Cell</i> , 2019, 73, 738-748.e9. | 4.5 | 128 |
| 5 | Protocol for Disome Profiling to Survey Ribosome Collision in Humans and Zebrafish. <i>STAR Protocols</i> , 2020, 1, 100168. | 0.5 | 40 |
| 6 | The long noncoding RNA <i>NEAT1_1</i> is seemingly dispensable for normal tissue homeostasis and cancer cell growth. <i>Rna</i> , 2019, 25, 1681-1695. | 1.6 | 39 |
| 7 | Dual targeting of DDX3 and eIF4A by the translation inhibitor rocaglamide A. <i>Cell Chemical Biology</i> , 2021, 28, 475-486.e8. | 2.5 | 37 |
| 8 | Simultaneous multicolor detection of RNA and proteins using super-resolution microscopy. <i>Methods</i> , 2016, 98, 158-165. | 1.9 | 36 |
| 9 | Forced isoform switching of <i>Neat1_1</i> to <i>Neat1_2</i> leads to the loss of <i>Neat1_1</i> and the hyperformation of paraspeckles but does not affect the development and growth of mice. <i>Rna</i> , 2020, 26, 251-264. | 1.6 | 27 |
| 10 | Regulation of gene expression via retrotransposon insertions and the noncoding <i>4.5S RNA_H</i> . <i>Genes To Cells</i> , 2015, 20, 887-901. | 0.5 | 15 |
| 11 | Cell Type-Specific Survey of Epigenetic Modifications by Tandem Chromatin Immunoprecipitation Sequencing. <i>Scientific Reports</i> , 2018, 8, 1143. | 1.6 | 5 |
| 12 | UPA-seq: prediction of functional lncRNAs using differential sensitivity to UV crosslinking. <i>Rna</i> , 2018, 24, 1785-1802. | 1.6 | 4 |
| 13 | TChIP-Seq: Cell-Type-Specific Epigenome Profiling. <i>Journal of Visualized Experiments</i> , 2019, , . | 0.2 | 1 |