

# Mikiko C Siomi

## List of Publications by Year in descending order

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67  
papers

13,383  
citations

100601

38  
h-index

116156

66  
g-index

69  
all docs

69  
docs citations

69  
times ranked

12739  
citing authors

#	ARTICLE	IF	CITATIONS
1	T-hairpin structure found in the RNA element involved in piRNA biogenesis. <i>Rna</i> , 2022, 28, 541-550.	1.6	4
2	Maelstrom functions in the production of Siwi-piRISC capable of regulating transposons in <i>Bombyx</i> germ cells. <i>iScience</i> , 2022, 25, 103914.	1.9	5
3	Siwi cooperates with Par-1 kinase to resolve the autoinhibitory effect of Papi for Siwi-piRISC biogenesis. <i>Nature Communications</i> , 2022, 13, 1518.	5.8	1
4	DEAD-box polypeptide 43 facilitates piRNA amplification by actively liberating RNA from Ago3-piRISC. <i>EMBO Reports</i> , 2021, 22, e51313.	2.0	14
5	Hamster PIWI proteins bind to piRNAs with stage-specific size variations during oocyte maturation. <i>Nucleic Acids Research</i> , 2021, 49, 2700-2720.	6.5	26
6	piRNA- and siRNA-mediated transcriptional repression in <i>Drosophila</i> , mice, and yeast: new insights and biodiversity. <i>EMBO Reports</i> , 2021, 22, e53062.	2.0	31
7	Japan: prize diversity, not conformity, to boost research. <i>Nature</i> , 2021, 599, 201-201.	13.7	1
8	Armitage determines Piwi-piRISC processing from precursor formation and quality control to interorganellar translocation. <i>EMBO Reports</i> , 2020, 21, e48769.	2.0	19
9	Piwi suppresses transcription of Brahma-dependent transposons via Maelstrom in ovarian somatic cells. <i>Science Advances</i> , 2020, 6, .	4.7	18
10	The Mi-2 nucleosome remodeler and the Rpd3 histone deacetylase are involved in piRNA-guided heterochromatin formation. <i>Nature Communications</i> , 2020, 11, 2818.	5.8	30
11	The piRNA pathway in <i>Drosophila</i> ; ovarian germ and somatic cells. <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 2020, 96, 32-42.	1.6	50
12	Crystal structure of <i>Drosophila</i> Piwi. <i>Nature Communications</i> , 2020, 11, 858.	5.8	42
13	Siwi levels reversibly regulate secondary piRISC biogenesis by affecting Ago3 body morphology in <i>Bombyx mori</i> . <i>EMBO Journal</i> , 2020, 39, e105130.	3.5	13
14	Assembly and Function of Gonad-Specific Non-Membranous Organelles in <i>Drosophila</i> piRNA Biogenesis. <i>Non-coding RNA</i> , 2019, 5, 52.	1.3	5
15	Distinct and Collaborative Functions of Yb and Armitage in Transposon-Targeting piRNA Biogenesis. <i>Cell Reports</i> , 2019, 27, 1822-1835.e8.	2.9	37
16	Requirements for multivalent Yb body assembly in transposon silencing in <i>Drosophila</i> . <i>EMBO Reports</i> , 2019, 20, e47708.	2.0	25
17	Essential roles of Wndei and nuclear monoubiquitination of Eggless/SETDB1 in transposon silencing. <i>EMBO Reports</i> , 2019, 20, e48296.	2.0	34
18	Nuclear RNA export factor variant initiates piRNA-guided co-transcriptional silencing. <i>EMBO Journal</i> , 2019, 38, e102870.	3.5	57

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19	Hierarchical roles of mitochondrial Papi and Zucchini in Bombyx germline piRNA biogenesis. <i>Nature</i> , 2018, 555, 260-264.	13.7	44
20	PIWI-Interacting RNA in <i>Drosophila</i> : Biogenesis, Transposon Regulation, and Beyond. <i>Chemical Reviews</i> , 2018, 118, 4404-4421.	23.0	82
21	The PIWI-Interacting RNA Molecular Pathway: Insights From Cultured Silkworm Germline Cells. <i>BioEssays</i> , 2018, 40, 1700068.	1.2	21
22	Two distinct transcriptional controls triggered by nuclear Piwi-piRISCs in the <i>Drosophila</i> piRNA pathway. <i>Current Opinion in Structural Biology</i> , 2018, 53, 69-76.	2.6	20
23	Piwi Nuclear Localization and Its Regulatory Mechanism in <i>Drosophila</i> Ovarian Somatic Cells. <i>Cell Reports</i> , 2018, 23, 3647-3657.	2.9	45
24	Use of the CRISPR-Cas9 system for genome editing in cultured <i>Drosophila</i> ovarian somatic cells. <i>Methods</i> , 2017, 126, 186-192.	1.9	8
25	Loss of <i>l(3)mbt</i> leads to acquisition of the ping-pong cycle in <i>Drosophila</i> ovarian somatic cells. <i>Genes and Development</i> , 2016, 30, 1617-1622.	2.7	30
26	Inheritance of a Nuclear PIWI from Pluripotent Stem Cells by Somatic Descendants Ensures Differentiation by Silencing Transposons in Planarian. <i>Developmental Cell</i> , 2016, 37, 226-237.	3.1	71
27	Crystal Structure of Silkworm PIWI-Clade Argonaute Siwi Bound to piRNA. <i>Cell</i> , 2016, 167, 484-497.e9.	13.5	116
28	Piwi Modulates Chromatin Accessibility by Regulating Multiple Factors Including Histone H1 to Repress Transposons. <i>Molecular Cell</i> , 2016, 63, 408-419.	4.5	110
29	piRNA biogenesis in the germline: From transcription of piRNA genomic sources to piRNA maturation. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2016, 1859, 82-92.	0.9	87
30	Somatic Primary piRNA Biogenesis Driven by cis-Acting RNA Elements and trans-Acting Yb. <i>Cell Reports</i> , 2015, 12, 429-440.	2.9	63
31	Tudor-domain containing proteins act to make the piRNA pathways more robust in <i>Drosophila</i> . <i>Fly</i> , 2015, 9, 86-90.	0.9	13
32	Respective Functions of Two Distinct Siwi Complexes Assembled during PIWI-Interacting RNA Biogenesis in Bombyx Germ Cells. <i>Cell Reports</i> , 2015, 10, 193-203.	2.9	94
33	PIWI-Interacting RNA: Its Biogenesis and Functions. <i>Annual Review of Biochemistry</i> , 2015, 84, 405-433.	5.0	579
34	Krimper Enforces an Antisense Bias on piRNA Pools by Binding AGO3 in the <i>Drosophila</i> Germline. <i>Molecular Cell</i> , 2015, 59, 553-563.	4.5	61
35	Functional and structural insights into the piRNA factor Maelstrom. <i>FEBS Letters</i> , 2015, 589, 1688-1693.	1.3	25
36	Crystal Structure and Activity of the Endoribonuclease Domain of the piRNA Pathway Factor Maelstrom. <i>Cell Reports</i> , 2015, 11, 366-375.	2.9	36

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37	Phased piRNAs tackle transposons. <i>Science</i> , 2015, 348, 756-757.	6.0	12
38	Immuno-Electron Microscopy and Electron Microscopic In Situ Hybridization for Visualizing piRNA Biogenesis Bodies in <i>Drosophila</i> Ovaries. <i>Methods in Molecular Biology</i> , 2015, 1328, 163-178.	0.4	21
39	piRNA clusters and open chromatin structure. <i>Mobile DNA</i> , 2014, 5, 22.	1.3	86
40	Yb Integrates piRNA Intermediates and Processing Factors into Perinuclear Bodies to Enhance piRISC Assembly. <i>Cell Reports</i> , 2014, 8, 103-113.	2.9	62
41	Small RNA profiling and characterization of piRNA clusters in the adult testes of the common marmoset, a model primate. <i>Rna</i> , 2014, 20, 1223-1237.	1.6	80
42	Roles of R2D2, a Cytoplasmic D2 Body Component, in the Endogenous siRNA Pathway in <i>Drosophila</i> . <i>Molecular Cell</i> , 2013, 49, 680-691.	4.5	62
43	DmGTSF1 is necessary for Piwi-mediated piRISC-mediated transcriptional transposon silencing in the <i>Drosophila</i> ovary. <i>Genes and Development</i> , 2013, 27, 1656-1661.	2.7	122
44	Biology of PIWI-interacting RNAs: new insights into biogenesis and function inside and outside of germlines. <i>Genes and Development</i> , 2012, 26, 2361-2373.	2.7	305
45	Structure and function of Zucchini endoribonuclease in piRNA biogenesis. <i>Nature</i> , 2012, 491, 284-287.	13.7	298
46	Gender-Specific Hierarchy in Nuage Localization of PIWI-Interacting RNA Factors in <i>Drosophila</i> . <i>Frontiers in Genetics</i> , 2011, 2, 55.	1.1	33
47	PIWI-interacting small RNAs: the vanguard of genome defence. <i>Nature Reviews Molecular Cell Biology</i> , 2011, 12, 246-258.	16.1	1,114
48	Maelstrom coordinates microtubule organization during <i>Drosophila</i> oogenesis through interaction with components of the MTOC. <i>Genes and Development</i> , 2011, 25, 2361-2373.	2.7	65
49	How does the Royal Family of Tudor rule the PIWI-interacting RNA pathway?. <i>Genes and Development</i> , 2010, 24, 636-646.	2.7	172
50	Biogenesis pathways of piRNAs loaded onto AGO3 in the <i>Drosophila</i> testis. <i>Rna</i> , 2010, 16, 2503-2515.	1.6	109
51	Roles for the Yb body components Armitage and Yb in primary piRNA biogenesis in <i>Drosophila</i> . <i>Genes and Development</i> , 2010, 24, 2493-2498.	2.7	261
52	Small RNA-Mediated Quiescence of Transposable Elements in Animals. <i>Developmental Cell</i> , 2010, 19, 687-697.	3.1	156
53	Characterization of the miRNA-RISC loading complex and miRNA-RISC formed in the <i>Drosophila</i> miRNA pathway. <i>Rna</i> , 2009, 15, 1282-1291.	1.6	96
54	RNA silencing in germlines: exquisite collaboration of Argonaute proteins with small RNAs for germline survival. <i>Current Opinion in Cell Biology</i> , 2009, 21, 426-434.	2.6	35

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55	Functional involvement of Tudor and dPRMT5 in the piRNA processing pathway in <i>Drosophila</i> germlines. <i>EMBO Journal</i> , 2009, 28, 3820-3831.	3.5	174
56	On the road to reading the RNA-interference code. <i>Nature</i> , 2009, 457, 396-404.	13.7	583
57	A regulatory circuit for piwi by the large Maf gene traffic jam in <i>Drosophila</i> . <i>Nature</i> , 2009, 461, 1296-1299.	13.7	387
58	Biogenesis of small RNAs in animals. <i>Nature Reviews Molecular Cell Biology</i> , 2009, 10, 126-139.	16.1	2,885
59	<i>Drosophila</i> endogenous small RNAs bind to Argonaute2 in somatic cells. <i>Nature</i> , 2008, 453, 793-797.	13.7	417
60	Pimet, the <i>Drosophila</i> homolog of HEN1, mediates 2'-O-methylation of Piwi-interacting RNAs at their 3' ends. <i>Genes and Development</i> , 2007, 21, 1603-1608.	2.7	400
61	Gene silencing mechanisms mediated by Aubergine piRNA complexes in <i>Drosophila</i> male gonad. <i>Rna</i> , 2007, 13, 1911-1922.	1.6	245
62	A Slicer-Mediated Mechanism for Repeat-Associated siRNA 5' End Formation in <i>Drosophila</i> . <i>Science</i> , 2007, 315, 1587-1590.	6.0	1,065
63	Specific association of Piwi with rasiRNAs derived from retrotransposon and heterochromatic regions in the <i>Drosophila</i> genome. <i>Genes and Development</i> , 2006, 20, 2214-2222.	2.7	566
64	Slicer function of <i>Drosophila</i> Argonautes and its involvement in RISC formation. <i>Genes and Development</i> , 2005, 19, 2837-2848.	2.7	343
65	Processing of Pre-microRNAs by the Dicer-1 Loquacious Complex in <i>Drosophila</i> Cells. <i>PLoS Biology</i> , 2005, 3, e235.	2.6	352
66	A <i>Drosophila</i> fragile X protein interacts with components of RNAi and ribosomal proteins. <i>Genes and Development</i> , 2002, 16, 2497-2508.	2.7	513
67	Essential role for KH domains in RNA binding: Impaired RNA binding by a mutation in the KH domain of FMR1 that causes fragile X syndrome. <i>Cell</i> , 1994, 77, 33-39.	13.5	437