Yuichiro Mishima

List of Publications by Year in descending order

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19	3,331	687220	839398
papers	citations	h-index	g-index
19 all docs	19 docs citations	19 times ranked	4316 citing authors
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#	Article	IF	CITATIONS
1	Zebrafish MiR-430 Promotes Deadenylation and Clearance of Maternal mRNAs. Science, 2006, 312, 75-79.	6.0	1,405
2	A Novel miRNA Processing Pathway Independent of Dicer Requires Argonaute2 Catalytic Activity. Science, 2010, 328, 1694-1698.	6.0	718
3	Differential Regulation of Germline mRNAs in Soma and Germ Cells by Zebrafish miR-430. Current Biology, 2006, 16, 2135-2142.	1.8	280
4	Codon Usage and 3′ UTR Length Determine Maternal mRNA Stability in Zebrafish. Molecular Cell, 2016, 61, 874-885.	4.5	229
5	Zebrafish miR-1 and miR-133 shape muscle gene expression and regulate sarcomeric actin organization. Genes and Development, 2009, 23, 619-632.	2.7	149
6	MicroRNAs Trigger Dissociation of elF4Al and elF4All from Target mRNAs in Humans. Molecular Cell, 2014, 56, 79-89.	4.5	117
7	Translational inhibition by deadenylation-independent mechanisms is central to microRNA-mediated silencing in zebrafish. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 1104-1109.	3.3	86
8	DAZL Relieves miRNA-Mediated Repression of Germline mRNAs by Controlling Poly(A) Tail Length in Zebrafish. PLoS ONE, 2009, 4, e7513.	1.1	85
9	Elements and machinery of nonâ€coding <scp>RNA</scp> s: toward their taxonomy. EMBO Reports, 2014, 15, 489-507.	2.0	84
10	miR-1-2 Gets to the Heart of the Matter. Cell, 2007, 129, 247-249.	13.5	42
10	miR-1-2 Gets to the Heart of the Matter. Cell, 2007, 129, 247-249. Widespread roles of microRNAs during zebrafish development and beyond. Development Growth and Differentiation, 2012, 54, 55-65.	0.6	42
	Widespread roles of microRNAs during zebrafish development and beyond. Development Growth and		
11	Widespread roles of microRNAs during zebrafish development and beyond. Development Growth and Differentiation, 2012, 54, 55-65. Protocol for Disome Profiling to Survey Ribosome Collision in Humans and Zebrafish. STAR	0.6	41
11 12	Widespread roles of microRNAs during zebrafish development and beyond. Development Growth and Differentiation, 2012, 54, 55-65. Protocol for Disome Profiling to Survey Ribosome Collision in Humans and Zebrafish. STAR Protocols, 2020, 1, 100168. Ribosome slowdown triggers codonâ€mediated mRNA decay independently of ribosome quality control.	0.6	41
11 12 13	Widespread roles of microRNAs during zebrafish development and beyond. Development Growth and Differentiation, 2012, 54, 55-65. Protocol for Disome Profiling to Survey Ribosome Collision in Humans and Zebrafish. STAR Protocols, 2020, 1, 100168. Ribosome slowdown triggers codonâ€mediated mRNA decay independently of ribosome quality control. EMBO Journal, 2022, 41, e109256. Pervasive yet nonuniform contributions of Dcp2 and Cnot7 to maternal ⟨scp⟩mRNA⟨/scp⟩ clearance in	0.6 0.5 3.5	41 40 25
11 12 13 14	Widespread roles of microRNAs during zebrafish development and beyond. Development Growth and Differentiation, 2012, 54, 55-65. Protocol for Disome Profiling to Survey Ribosome Collision in Humans and Zebrafish. STAR Protocols, 2020, 1, 100168. Ribosome slowdown triggers codonâ€mediated mRNA decay independently of ribosome quality control. EMBO Journal, 2022, 41, e109256. Pervasive yet nonuniform contributions of Dcp2 and Cnot7 to maternal <scp>mRNA</scp> clearance in zebrafish. Genes To Cells, 2017, 22, 670-678. Deadenylation by the <scp>CCR</scp> 4â€ <scp>NOT</scp> complex contributes to the turnover of <i>hairy</i> i>hairyi>â€related <scp>mRNA</scp> s in the zebrafish segmentation clock. FEBS Letters, 2018, 592,	0.6 0.5 3.5	41 40 25 10
11 12 13 14	Widespread roles of microRNAs during zebrafish development and beyond. Development Growth and Differentiation, 2012, 54, 55-65. Protocol for Disome Profiling to Survey Ribosome Collision in Humans and Zebrafish. STAR Protocols, 2020, 1, 100168. Ribosome slowdown triggers codonâ€mediated mRNA decay independently of ribosome quality control. EMBO Journal, 2022, 41, e109256. Pervasive yet nonuniform contributions of Dcp2 and Cnot7 to maternal ⟨scp⟩mRNA⟨/scp⟩ clearance in zebrafish. Genes To Cells, 2017, 22, 670-678. Deadenylation by the ⟨scp⟩CCR⟨/scp⟩4â€⟨scp⟩NOT⟨/scp⟩ complex contributes to the turnover of ⟨i⟩hairy⟨ i⟩á€related ⟨scp⟩mRNA⟨/scp⟩s in the zebrafish segmentation clock. FEBS Letters, 2018, 592, 3388-3398. Roles of mRNA Fate Modulators Dhh1 and Pat1 in TNRC6-dependent Gene Silencing Recapitulated in	0.6 0.5 3.5 0.5	41 40 25 10

#	Article	IF	CITATIONS
19	Tethered Function Assay to Study RNA-Regulatory Proteins in Zebrafish. Methods in Molecular Biology, 2021, 2218, 347-354.	0.4	0