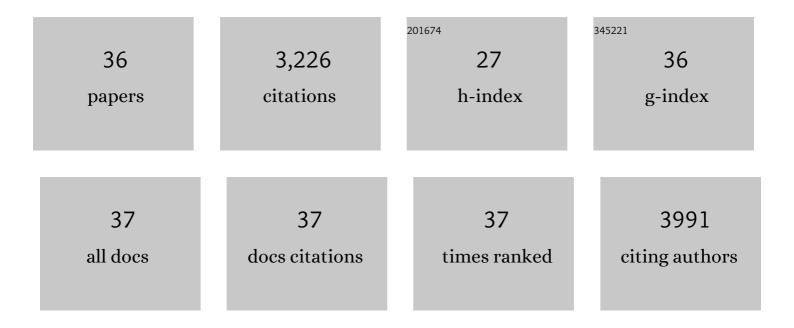
## Daisuke Miki

List of Publications by Year in descending order

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DAISLIKE MIKI

#	Article	IF	CITATIONS
1	Three highly conserved hydrophobic residues in the predicted α2â€helix of rice NLR protein Pit contribute to its localization and immune induction. Plant, Cell and Environment, 2022, , .	5.7	2
2	Roles of DEMETER in regulating DNA methylation in vegetative tissues and pathogen resistance. Journal of Integrative Plant Biology, 2021, 63, 691-706.	8.5	26
3	CRISPR/Cas9-Based Genome Editing Toolbox for Arabidopsis thaliana. Methods in Molecular Biology, 2021, 2200, 121-146.	0.9	14
4	Gene Targeting Facilitated by Engineered Sequence-Specific Nucleases: Potential Applications for Crop Improvement. Plant and Cell Physiology, 2021, 62, 752-765.	3.1	6
5	Genomeâ€wide distribution and functions of the AAE complex in epigenetic regulation in <i>Arabidopsis</i> . Journal of Integrative Plant Biology, 2021, 63, 707-722.	8.5	18
6	Gene targeting in <i>Arabidopsis</i> via an allâ€inâ€one strategy that uses a translational enhancer to aid Cas9 expression. Plant Biotechnology Journal, 2020, 18, 892-894.	8.3	23
7	DNA demethylases are required for myo-inositol-mediated mutualism between plants and beneficial rhizobacteria. Nature Plants, 2020, 6, 983-995.	9.3	48
8	Histone acetylation recruits the SWR1 complex to regulate active DNA demethylation in <i>Arabidopsis</i> . Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 16641-16650.	7.1	73
9	The genome of broomcorn millet. Nature Communications, 2019, 10, 436.	12.8	130
10	DNA demethylase ROS1 negatively regulates the imprinting of <i>DOGL4</i> and seed dormancy in <i>Arabidopsis thaliana</i> . Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E9962-E9970.	7.1	46
11	Four putative SWI2/SNF2 chromatin remodelers have dual roles in regulating DNA methylation in Arabidopsis. Cell Discovery, 2018, 4, 55.	6.7	22
12	CRISPR/Cas9-mediated gene targeting in Arabidopsis using sequential transformation. Nature Communications, 2018, 9, 1967.	12.8	178
13	The developmental regulator PKL is required to maintain correct DNA methylation patterns at RNA-directed DNA methylation loci. Genome Biology, 2017, 18, 103.	8.8	44
14	Efficient Generation of diRNAs Requires Components in the Posttranscriptional Gene Silencing Pathway. Scientific Reports, 2017, 7, 301.	3.3	34
15	A pair of transposon-derived proteins function in a histone acetyltransferase complex for active DNA demethylation. Cell Research, 2017, 27, 226-240.	12.0	80
16	SAC3B, a central component of the mRNA export complex TREX-2, is required for prevention of epigenetic gene silencing in <i>Arabidopsis</i> . Nucleic Acids Research, 2017, 45, 181-197.	14.5	21
17	The Methyl-CpG-Binding Protein MBD7 Facilitates Active DNA Demethylation to Limit DNA Hyper-Methylation and Transcriptional Gene Silencing. Molecular Cell, 2015, 57, 971-983.	9.7	112
18	Involvement of Multiple Gene-Silencing Pathways in a Paramutation-like Phenomenon in Arabidopsis. Cell Reports, 2015, 11, 1160-1167.	6.4	13

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19	<i>Arabidopsis</i> EDM2 promotes <i>IBM1</i> distal polyadenylation and regulates genome DNA methylation patterns. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 527-532.	7.1	102
20	Regulation of Active DNA Demethylation by an α-Crystallin Domain Protein in Arabidopsis. Molecular Cell, 2014, 55, 361-371.	9.7	44
21	Insights into the Localization and Function of the Membrane Trafficking Regulator GNOM ARF-GEF at the Golgi Apparatus in <i>Arabidopsis</i> Â. Plant Cell, 2014, 26, 3062-3076.	6.6	121
22	Overproduction of stomatal lineage cells in Arabidopsis mutants defective in active DNA demethylation. Nature Communications, 2014, 5, 4062.	12.8	90
23	A Pre-mRNA-Splicing Factor Is Required for RNA-Directed DNA Methylation in Arabidopsis. PLoS Genetics, 2013, 9, e1003779.	3.5	58
24	Sulfamethazine Suppresses Epigenetic Silencing in <i>Arabidopsis</i> by Impairing Folate Synthesis. Plant Cell, 2012, 24, 1230-1241.	6.6	77
25	A DNA 3′ Phosphatase Functions in Active DNA Demethylation in Arabidopsis. Molecular Cell, 2012, 45, 357-370.	9.7	81
26	A Histone Acetyltransferase Regulates Active DNA Demethylation in <i>Arabidopsis</i> . Science, 2012, 336, 1445-1448.	12.6	224
27	An RNA polymerase II- and AGO4-associated protein acts in RNA-directed DNA methylation. Nature, 2010, 465, 106-109.	27.8	228
28	Knock-down of OsDCL2 in Rice Negatively Affects Maintenance of the Endogenous dsRNA Virus, Oryza sativa Endornavirus. Plant and Cell Physiology, 2010, 51, 58-67.	3.1	35
29	Analysis of the Rac/Rop Small GTPase Family in Rice: Expression, Subcellular Localization and Role in Disease Resistance. Plant and Cell Physiology, 2010, 51, 585-595.	3.1	113
30	DNA Replication Factor C1 Mediates Genomic Stability and Transcriptional Gene Silencing in <i>Arabidopsis</i> Â Â. Plant Cell, 2010, 22, 2336-2352.	6.6	72
31	ROS3 is an RNA-binding protein required for DNA demethylation in Arabidopsis. Nature, 2008, 455, 1259-1262.	27.8	150
32	Small interfering RNA (siRNA) targeting of endogenous promoters induces DNA methylation, but not necessarily gene silencing, in rice. Plant Journal, 2008, 53, 65-77.	5.7	65
33	<i>De novo</i> DNA methylation induced by siRNA targeted to endogenous transcribed sequences is geneâ€specific and <i>OsMet1</i> â€independent in rice. Plant Journal, 2008, 56, 539-549.	5.7	29
34	RNAi-mediated Silencing of OsGEN-L (OsGEN-like), a New Member of the RAD2/XPG Nuclease Family, Causes Male Sterility by Defect of Microspore Development in Rice. Plant and Cell Physiology, 2005, 46, 699-715.	3.1	75
35	RNA Silencing of Single and Multiple Members in a Gene Family of Rice. Plant Physiology, 2005, 138, 1903-1913.	4.8	270
36	Simple RNAi Vectors for Stable and Transient Suppression of Gene Function in Rice. Plant and Cell Physiology, 2004, 45, 490-495.	3.1	502