

Tatsuo Miyamoto

List of Publications by Year in descending order

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Version: 2024-02-01

36
papers

1,691
citations

430843

18
h-index

361001

35
g-index

38
all docs

38
docs citations

38
times ranked

2879
citing authors

#	ARTICLE	IF	CITATIONS
1	YAP is essential for tissue tension to ensure vertebrate 3D body shape. <i>Nature</i> , 2015, 521, 217-221.	27.8	237
2	Repeating pattern of non-RVD variations in DNA-binding modules enhances TALEN activity. <i>Scientific Reports</i> , 2013, 3, 3379.	3.3	195
3	Efficient TALEN construction and evaluation methods for human cell and animal applications. <i>Genes To Cells</i> , 2013, 18, 315-326.	1.2	190
4	Tight junctions in Schwann cells of peripheral myelinated axons. <i>Journal of Cell Biology</i> , 2005, 169, 527-538.	5.2	176
5	Compartmentalization established by claudin-11-based tight junctions in stria vascularis is required for hearing through generation of endocochlear potential. <i>Journal of Cell Science</i> , 2004, 117, 5087-5096.	2.0	169
6	The Microtubule-Depolymerizing Activity of a Mitotic Kinesin Protein KIF2A Drives Primary Cilia Disassembly Coupled with Cell Proliferation. <i>Cell Reports</i> , 2015, 10, 664-673.	6.4	128
7	Space Radiation Biology for "Living in Space". <i>BioMed Research International</i> , 2020, 2020, 1-25.	1.9	75
8	Insufficiency of BUBR1, a mitotic spindle checkpoint regulator, causes impaired ciliogenesis in vertebrates. <i>Human Molecular Genetics</i> , 2011, 20, 2058-2070.	2.9	52
9	TALEN-mediated single-base-pair editing identification of an intergenic mutation upstream of <i>BUB1B</i> as causative of PCS (MVA) syndrome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 1461-1466.	7.1	52
10	Two unrelated patients with MRE11A mutations and Nijmegen breakage syndrome-like severe microcephaly. <i>DNA Repair</i> , 2011, 10, 314-321.	2.8	49
11	Zinc-finger nuclease-mediated targeted insertion of reporter genes for quantitative imaging of gene expression in sea urchin embryos. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 10915-10920.	7.1	40
12	Insufficiency of ciliary cholesterol in hereditary Zellweger syndrome. <i>EMBO Journal</i> , 2020, 39, e103499.	7.8	35
13	Genome-wide association meta-analysis identifies GP2 gene risk variants for pancreatic cancer. <i>Nature Communications</i> , 2020, 11, 3175.	12.8	34
14	HpSulf, a heparan sulfate 6-O-endosulfatase, is involved in the regulation of VEGF signaling during sea urchin development. <i>Mechanisms of Development</i> , 2010, 127, 235-245.	1.7	33
15	PLK1-mediated phosphorylation of WDR62/MCPH2 ensures proper mitotic spindle orientation. <i>Human Molecular Genetics</i> , 2017, 26, 4429-4440.	2.9	32
16	Cytoskeleton-related regulation of primary cilia shortening mediated by melanin-concentrating hormone receptor 1. <i>General and Comparative Endocrinology</i> , 2017, 253, 44-52.	1.8	24
17	Functional consequence of fibulin-4 missense mutations associated with vascular and skeletal abnormalities and cutis laxa. <i>Matrix Biology</i> , 2016, 56, 132-149.	3.6	19
18	Evaluation of ATM heterozygous mutations underlying individual differences in radiosensitivity using genome editing in human cultured cells. <i>Scientific Reports</i> , 2017, 7, 5996.	3.3	18

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19	Sulforaphane suppresses metastasis of triple-negative breast cancer cells by targeting the RAF/MEK/ERK pathway. <i>Npj Breast Cancer</i> , 2022, 8, 40.	5.2	17
20	Characterization of Functional Primary Cilia in Human Induced Pluripotent Stem Cell-Derived Neurons. <i>Neurochemical Research</i> , 2019, 44, 1736-1744.	3.3	16
21	Cilia play a role in breaking left-right symmetry of the sea urchin embryo. <i>Genes To Cells</i> , 2016, 21, 568-578.	1.2	12
22	Coordinated regulation of the dorsal-ventral and anterior-posterior patterning of <i>Xenopus</i> embryos by the <i>BTB</i> / <i>POZ</i> zinc finger protein <i>Zbtb14</i> . <i>Development Growth and Differentiation</i> , 2018, 60, 158-173.	1.5	11
23	Exploration of genetic basis underlying individual differences in radiosensitivity within human populations using genome editing technology. <i>Journal of Radiation Research</i> , 2018, 59, ii75-ii82.	1.6	11
24	Ciliary GPCR-based transcriptome as a key regulator of cilia length control. <i>FASEB BioAdvances</i> , 2021, 3, 744-767.	2.4	11
25	Premature aging syndrome showing random chromosome number instabilities with <i>CDC20</i> mutation. <i>Aging Cell</i> , 2020, 19, e13251.	6.7	10
26	Analysis of individual differences in radiosensitivity using genome editing. <i>Annals of the ICRP</i> , 2016, 45, 290-296.	3.8	8
27	Generation of transgenic medaka expressing claudin7-EGFP for imaging of tight junctions in living medaka embryos. <i>Cell and Tissue Research</i> , 2009, 335, 465-471.	2.9	6
28	Updated summary of genome editing technology in human cultured cells linked to human genetics studies. <i>Journal of Human Genetics</i> , 2018, 63, 133-143.	2.3	6
29	<i>Albatross</i> / <i>FBF1</i> contributes to both centriole duplication and centrosome separation. <i>Genes To Cells</i> , 2018, 23, 1023-1042.	1.2	6
30	Applications of Genome Editing Technology in Research on Chromosome Aneuploidy Disorders. <i>Cells</i> , 2020, 9, 239.	4.1	5
31	iPSC reprogramming-mediated aneuploidy correction in autosomal trisomy syndromes. <i>PLoS ONE</i> , 2022, 17, e0264965.	2.5	4
32	A novel CDK-independent function of p27Kip1 in preciliary vesicle trafficking during ciliogenesis. <i>Biochemical and Biophysical Research Communications</i> , 2020, 527, 716-722.	2.1	3
33	<i>NBS1</i> I171V variant underlies individual differences in chromosomal radiosensitivity within human populations. <i>Scientific Reports</i> , 2021, 11, 19661.	3.3	3
34	In Vivo Imaging of Tight Junctions Using Claudin-EGFP Transgenic Medaka. <i>Methods in Molecular Biology</i> , 2011, 762, 171-178.	0.9	2
35	Imaging of the Ciliary Cholesterol Underlying the Sonic Hedgehog Signal Transduction. <i>Methods in Molecular Biology</i> , 2022, 2374, 49-57.	0.9	1
36	Ciliopathy in PCS (MVA) syndrome. <i>Oncotarget</i> , 2015, 6, 24582-24583.	1.8	1