

Masato Ohtsuka

List of Publications by Year in descending order

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82
papers

2,608
citations

270111

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all docs

87
docs citations

87
times ranked

3375
citing authors

#	ARTICLE	IF	CITATIONS
1	Cyclin D1 Binding Protein 1 Responds to DNA Damage through the ATM-CHK2 Pathway. <i>Journal of Clinical Medicine</i> , 2022, 11, 851.	1.0	2
2	Dll1 Can Function as a Ligand of Notch1 and Notch2 in the Thymic Epithelium. <i>Frontiers in Immunology</i> , 2022, 13, 852427.	2.2	3
3	Use of anti-inhibin monoclonal antibody for increasing the litter size of mouse strains and its application to i-GONAD. <i>Biology of Reproduction</i> , 2022, , .	1.2	7
4	AMBRA1 controls antigen-driven activation and proliferation of naive T cells. <i>International Immunology</i> , 2021, 33, 107-118.	1.8	3
5	Designing and generating a mouse model: frequently asked questions. <i>Journal of Biomedical Research</i> , 2021, 35, 76.	0.7	6
6	Response to correspondence on "Reproducibility of CRISPR-Cas9 methods for generation of conditional mouse alleles: a multi-center evaluation". <i>Genome Biology</i> , 2021, 22, 99.	3.8	4
7	Interleukin-11-expressing fibroblasts have a unique gene signature correlated with poor prognosis of colorectal cancer. <i>Nature Communications</i> , 2021, 12, 2281.	5.8	60
8	Novel reporter mouse models useful for evaluating in vivo gene editing and for optimization of methods of delivering genome editing tools. <i>Molecular Therapy - Nucleic Acids</i> , 2021, 24, 325-336.	2.3	10
9	GONAD: A new method for germline genome editing in mice and rats. <i>Development Growth and Differentiation</i> , 2021, 63, 439-447.	0.6	11
10	Alopecia areata susceptibility variant in MHC region impacts expressions of genes contributing to hair keratinization and is involved in hair loss. <i>EBioMedicine</i> , 2020, 57, 102810.	2.7	19
11	Genetically modified mouse models to help fight COVID-19. <i>Nature Protocols</i> , 2020, 15, 3777-3787.	5.5	26
12	Monitoring the autophagy-endolysosomal system using monomeric Keima-fused MAP1LC3B. <i>PLoS ONE</i> , 2020, 15, e0234180.	1.1	2
13	<i>Thy1</i> promoter activity in the <i>Rosa26</i> locus in mice: lessons from Dre-ox<i> conditional expression system. <i>Experimental Animals</i> , 2020, 69, 287-294.	0.7	1
14	Effect of Diphtheria Toxin-Based Gene Therapy for Hepatocellular Carcinoma. <i>Cancers</i> , 2020, 12, 472.	1.7	13
15	Sequential i-GONAD: An Improved In Vivo Technique for CRISPR/Cas9-Based Genetic Manipulations in Mice. <i>Cells</i> , 2020, 9, 546.	1.8	13
16	Acrosin is essential for sperm penetration through the zona pellucida in hamsters. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 2513-2518.	3.3	64
17	Modification of i-GONAD Suitable for Production of Genome-Edited C57BL/6 Inbred Mouse Strain. <i>Cells</i> , 2020, 9, 957.	1.8	10
18	Creation of CRISPR-based germline-genome-engineered mice without ex vivo handling of zygotes by i-GONAD. <i>Nature Protocols</i> , 2019, 14, 2452-2482.	5.5	93

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19	Reproducibility of CRISPR-Cas9 methods for generation of conditional mouse alleles: a multi-center evaluation. <i>Genome Biology</i> , 2019, 20, 171.	3.8	69
20	i-GONAD : A method for generating genome-edited animals without ex vivo handling of embryos. <i>Development Growth and Differentiation</i> , 2019, 61, 306-315.	0.6	19
21	SAM68-Specific Splicing Is Required for Proper Selection of Alternative 5' UTR Isoforms in the Nervous System. <i>iScience</i> , 2019, 22, 318-335.	1.9	15
22	Isolation and Analysis of a Genome-Edited Single-Hepatocyte from a Cas9 Transgenic Mouse Line. <i>Methods in Molecular Biology</i> , 2019, 1874, 257-271.	0.4	0
23	Easi-CRISPR for creating knock-in and conditional knockout mouse models using long ssDNA donors. <i>Nature Protocols</i> , 2018, 13, 195-215.	5.5	209
24	Successful production of genome-edited rats by the rGONAD method. <i>BMC Biotechnology</i> , 2018, 18, 19.	1.7	40
25	i-GONAD: a robust method for in situ germline genome engineering using CRISPR nucleases. <i>Genome Biology</i> , 2018, 19, 25.	3.8	130
26	Cell-cell interactions between monocytes/macrophages and synoviocyte-like cells promote inflammatory cell infiltration mediated by augmentation of MCP-1 production in temporomandibular joint. <i>Bioscience Reports</i> , 2018, 38, .	1.1	15
27	Timing of CRISPR/Cas9-related mRNA microinjection after activation as an important factor affecting genome editing efficiency in porcine oocytes. <i>Theriogenology</i> , 2018, 108, 29-38.	0.9	31
28	Intraoviductal Instillation of a Solution as an Effective Route for Manipulating Preimplantation Mammalian Embryos in vivo. , 2018, , .		10
29	Intravenous Delivery of piggyBac Transposons as a Useful Tool for Liver-Specific Gene-Switching. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3452.	1.8	10
30	In vivo genome editing targeted towards the female reproductive system. <i>Archives of Pharmacal Research</i> , 2018, 41, 898-910.	2.7	7
31	i-GONAD (improved genome-editing via oviductal nucleic acids delivery), a convenient in vivo tool to produce genome-edited rats. <i>Scientific Reports</i> , 2018, 8, 12059.	1.6	34
32	Easi-CRISPR: a robust method for one-step generation of mice carrying conditional and insertion alleles using long ssDNA donors and CRISPR ribonucleoproteins. <i>Genome Biology</i> , 2017, 18, 92.	3.8	375
33	Simplified CRISPR tools for efficient genome editing and streamlined protocols for their delivery into mammalian cells and mouse zygotes. <i>Methods</i> , 2017, 121-122, 16-28.	1.9	121
34	Bone marrow-derived mesenchymal stem cells propagate immunosuppressive/anti-inflammatory macrophages in cell-to-cell contact-independent and -dependent manners under hypoxic culture. <i>Experimental Cell Research</i> , 2017, 358, 411-420.	1.2	61
35	Efficient Generation of Somatic Cell Nuclear Transfer-Competent Porcine Cells with Mutated Alleles at Multiple Target Loci by Using CRISPR/Cas9 Combined with Targeted Toxin-Based Selection System. <i>International Journal of Molecular Sciences</i> , 2017, 18, 2610.	1.8	7
36	The piggyBac-Based Gene Delivery System Can Confer Successful Production of Cloned Porcine Blastocysts with Multigene Constructs. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1424.	1.8	5

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37	Establishment of Nephhrin Reporter Mice and Use for Chemical Screening. <i>PLoS ONE</i> , 2016, 11, e0157497.	1.1	6
38	CRISPR: a versatile tool for both forward and reverse genetics research. <i>Human Genetics</i> , 2016, 135, 971-976.	1.8	41
39	GONAD: A Novel CRISPR/Cas9 Genome Editing Method that Does Not Require Ex Vivo Handling of Embryos. <i>Current Protocols in Human Genetics</i> , 2016, 88, 15.8.1-15.8.12.	3.5	27
40	Three-dimensional X-ray visualization of axonal tracts in mouse brain hemisphere. <i>Scientific Reports</i> , 2016, 6, 35061.	1.6	15
41	Effective Prevention of Liver Fibrosis by Liver-targeted Hydrodynamic Gene Delivery of Matrix Metalloproteinase-13 in a Rat Liver Fibrosis Model. <i>Molecular Therapy - Nucleic Acids</i> , 2016, 5, e276.	2.3	33
42	Effects of Fibrotic Tissue on Liver-targeted Hydrodynamic Gene Delivery. <i>Molecular Therapy - Nucleic Acids</i> , 2016, 5, e359.	2.3	10
43	Pronuclear Injection-Based Targeted Transgenesis. <i>Current Protocols in Human Genetics</i> , 2016, 91, 15.10.1-15.10.28.	3.5	10
44	CRISPR/Cas9 and the Paradigm Shift in Mouse Genome Manipulation Technologies. , 2016, , 65-77.		3
45	Nucleic acids delivery methods for genome editing in zygotes and embryos: the old, the new, and the old-new. <i>Biology Direct</i> , 2016, 11, 16.	1.9	28
46	Conditional knockout of <i>Foxc2</i> gene in kidney: efficient generation of conditional alleles of single-exon gene by double-selection system. <i>Mammalian Genome</i> , 2016, 27, 62-69.	1.0	8
47	CRISPR/Cas9-based generation of knockdown mice by intronic insertion of artificial microRNA using longer single-stranded DNA. <i>Scientific Reports</i> , 2015, 5, 12799.	1.6	119
48	Direct Injection of CRISPR/Cas9-Related mRNA into Cytoplasm of Parthenogenetically Activated Porcine Oocytes Causes Frequent Mosaicism for Indel Mutations. <i>International Journal of Molecular Sciences</i> , 2015, 16, 17838-17856.	1.8	55
49	Assessment of Artificial MiRNA Architectures for Higher Knockdown Efficiencies without the Undesired Effects in Mice. <i>PLoS ONE</i> , 2015, 10, e0135919.	1.1	6
50	Insertion of sequences at the original provirus integration site of mouse <i>ROSA26</i> locus using the CRISPR/Cas9 system. <i>FEBS Open Bio</i> , 2015, 5, 191-197.	1.0	30
51	Establishment of immortalized mesenchymal stem cells derived from the submandibular glands of tdTomato transgenic mice. <i>Experimental and Therapeutic Medicine</i> , 2015, 10, 1380-1386.	0.8	5
52	A combination of targeted toxin technology and the piggyBac-mediated gene transfer system enables efficient isolation of stable transfectants in nonhuman mammalian cells. <i>Biotechnology Journal</i> , 2015, 10, 143-153.	1.8	6
53	GONAD: Genome-editing via Oviductal Nucleic Acids Delivery system: a novel microinjection independent genome engineering method in mice. <i>Scientific Reports</i> , 2015, 5, 11406.	1.6	98
54	One-step generation of multiple transgenic mouse lines using an improved Pronuclear Injection-based Targeted Transgenesis (i-PITT). <i>BMC Genomics</i> , 2015, 16, 274.	1.2	19

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55	PiggyBac transposon-mediated gene delivery efficiently generates stable transfectants derived from cultured primary human deciduous tooth dental pulp cells (HDDPCs) and HDDPC-derived iPS cells. <i>International Journal of Oral Science</i> , 2015, 7, 144-154.	3.6	17
56	Mouse Genome Editing Using the CRISPR/Cas System. <i>Current Protocols in Human Genetics</i> , 2014, 83, 15.7.1-27.	3.5	90
57	The combinational use of CRISPR/Cas9-based gene editing and targeted toxin technology enables efficient biallelic knockout of the β -galactosyltransferase gene in porcine embryonic fibroblasts. <i>Xenotransplantation</i> , 2014, 21, 291-300.	1.6	47
58	Development of Pronuclear Injection-Based Targeted Transgenesis in Mice Through Cre-loxP Site-Specific Recombination. <i>Methods in Molecular Biology</i> , 2014, 1194, 3-19.	0.4	9
59	Improvement of pronuclear injection-based targeted transgenesis (PITT) by iCre mRNA-mediated site-specific recombination. <i>Transgenic Research</i> , 2013, 22, 873-875.	1.3	9
60	piggyBac-mediated generation of stable transfectants with surface human leukocyte antigen expression from a small number of cells. <i>Analytical Biochemistry</i> , 2013, 437, 29-31.	1.1	8
61	X-ray microtomographic visualization of <i>Escherichia coli</i> by metalloprotein overexpression. <i>Journal of Synchrotron Radiation</i> , 2013, 20, 581-586.	1.0	5
62	Targeted Toxin-Based Selectable Drug-Free Enrichment of Mammalian Cells with High Transgene Expression. <i>Biology</i> , 2013, 2, 341-355.	1.3	11
63	In vivo gene transfer in mouse preimplantation embryos after intraoviductal injection of plasmid DNA and subsequent in vivo electroporation. <i>Systems Biology in Reproductive Medicine</i> , 2012, 58, 278-287.	1.0	11
64	PITT: Pronuclear Injection-Based Targeted Transgenesis, a Reliable Transgene Expression Method in Mice. <i>Experimental Animals</i> , 2012, 61, 489-502.	0.7	21
65	Fluorescent transgenic mice suitable for multi-color aggregation chimera studies. <i>Cell and Tissue Research</i> , 2012, 350, 251-260.	1.5	15
66	Targeted transgenesis through pronuclear injection of improved vectors into in vitro fertilized eggs. <i>Transgenic Research</i> , 2012, 21, 225-226.	1.3	11
67	Simple cloning strategy using GFPuv gene as positive/negative indicator. <i>Analytical Biochemistry</i> , 2011, 416, 237-239.	1.1	7
68	Double Anal Fin (Da): A Medaka Mutant Exhibiting a Mirror-Image Pattern Duplication of the Dorsal-Ventral Axis. , 2011, , 201-215.		2
69	Enrichment of xenograft-competent genetically modified pig cells using a targeted toxin, isolectin BS-I-B4 conjugate. <i>Xenotransplantation</i> , 2010, 17, 81-89.	1.6	12
70	Pronuclear injection-based mouse targeted transgenesis for reproducible and highly efficient transgene expression. <i>Nucleic Acids Research</i> , 2010, 38, e198-e198.	6.5	53
71	Recombinant DNA Technologies for Construction of Precisely Designed Transgene Constructs. <i>Current Pharmaceutical Biotechnology</i> , 2009, 10, 244-251.	0.9	20
72	Development of CRTEIL and CETRIZ, Cre-loxP-Based Systems, Which Allow Change of Expression of Red to Green or Green to Red Fluorescence upon Transfection with a Cre-Expression Vector. <i>Journal of Biomedicine and Biotechnology</i> , 2009, 2009, 1-9.	3.0	1

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73	A transposon-based chromosomal engineering method to survey a large cis-regulatory landscape in mice. <i>Nature Genetics</i> , 2009, 41, 946-952.	9.4	58
74	Cre ⁺ loxP system as a versatile tool for conferring increased levels of tissue-specific gene expression from a weak promoter. <i>Molecular Reproduction and Development</i> , 2008, 75, 1085-1093.	1.0	14
75	Major histocompatibility complex (Mhc) class Ib gene duplications, organization and expression patterns in mouse strain C57BL/6. <i>BMC Genomics</i> , 2008, 9, 178.	1.2	65
76	One-step generation of recombinering constructs by asymmetric-end ligation and negative selection. <i>Analytical Biochemistry</i> , 2007, 360, 306-308.	1.1	7
77	Construction of Mouse 129/Ola BAC Library for Targeting Experiments Using E14 Embryonic Stem Cells. <i>Genes and Genetic Systems</i> , 2006, 81, 143-146.	0.2	11
78	CHOP: visualization of 'wobbling' and isolation of highly conserved regions from aligned DNA sequences. <i>Nucleic Acids Research</i> , 2004, 32, W55-W58.	6.5	1
79	Possible roles of zic1 and zic4, identified within the medaka Double anal fin (Da) locus, in dorsoventral patterning of the trunk-tail region (related to phenotypes of the Da mutant). <i>Mechanisms of Development</i> , 2004, 121, 873-882.	1.7	29
80	Comparative analysis of a 229-kb medaka genomic region, containing the zic1 and zic4 genes, with Fugu, human, and mouse. <i>Genomics</i> , 2004, 83, 1063-1071.	1.3	6
81	Rapid Screening of a Novel Arrayed Medaka (<i>Oryzias latipes</i>) Cosmid Library. <i>Marine Biotechnology</i> , 2002, 4, 173-178.	1.1	6
82	Construction of a Linkage Map of the Medaka (<i>Oryzias latipes</i>) and Mapping of the Da Mutant Locus Defective in Dorsoventral Patterning. <i>Genome Research</i> , 1999, 9, 1277-1287.	2.4	31