## Tetsushi Sakuma

List of Publications by Year in descending order

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TETSUSHI SAKUMA

#	Article	IF	CITATIONS
1	Precise Correction of the Dystrophin Gene in Duchenne Muscular Dystrophy Patient Induced Pluripotent Stem Cells by TALEN and CRISPR-Cas9. Stem Cell Reports, 2015, 4, 143-154.	2.3	459
2	Robust InÂVitro Induction of Human Germ Cell Fate from Pluripotent Stem Cells. Cell Stem Cell, 2015, 17, 178-194.	5.2	428
3	Microhomology-mediated end-joining-dependent integration of donor DNA in cells and animals using TALENs and CRISPR/Cas9. Nature Communications, 2014, 5, 5560.	5.8	414
4	Multiplex genome engineering in human cells using all-in-one CRISPR/Cas9 vector system. Scientific Reports, 2014, 4, 5400.	1.6	318
5	MMEJ-assisted gene knock-in using TALENs and CRISPR-Cas9 with the PITCh systems. Nature Protocols, 2016, 11, 118-133.	5.5	315
6	Cloning-free CRISPR/Cas system facilitates functional cassette knock-in in mice. Genome Biology, 2015, 16, 87.	3.8	250
7	Precise in-frame integration of exogenous DNA mediated by CRISPR/Cas9 system in zebrafish. Scientific Reports, 2015, 5, 8841.	1.6	207
8	Sterol Side Chain Reductase 2 Is a Key Enzyme in the Biosynthesis of Cholesterol, the Common Precursor of Toxic Steroidal Glycoalkaloids in Potato Â. Plant Cell, 2014, 26, 3763-3774.	3.1	206
9	Repeating pattern of non-RVD variations in DNA-binding modules enhances TALEN activity. Scientific Reports, 2013, 3, 3379.	1.6	195
10	Efficient <scp>TALEN</scp> construction and evaluation methods for human cell and animal applications. Genes To Cells, 2013, 18, 315-326.	0.5	190
11	Human Induced Pluripotent Stem Cell–Derived Podocytes Mature into Vascularized Glomeruli upon Experimental Transplantation. Journal of the American Society of Nephrology: JASN, 2016, 27, 1778-1791.	3.0	186
12	Simple knockout by electroporation of engineered endonucleases into intact rat embryos. Scientific Reports, 2014, 4, 6382.	1.6	179
13	Generation of a Nonhuman Primate Model of Severe Combined Immunodeficiency Using Highly Efficient Genome Editing. Cell Stem Cell, 2016, 19, 127-138.	5.2	139
14	EDEM2 initiates mammalian glycoprotein ERAD by catalyzing the first mannose trimming step. Journal of Cell Biology, 2014, 206, 347-356.	2.3	131
15	The Microtubule-Depolymerizing Activity of a Mitotic Kinesin Protein KIF2A Drives Primary Cilia Disassembly Coupled with Cell Proliferation. Cell Reports, 2015, 10, 664-673.	2.9	128
16	Non-transgenic genome modifications in a hemimetabolous insect using zinc-finger and TAL effector nucleases. Nature Communications, 2012, 3, 1017.	5.8	115
17	Organoids from Nephrotic Disease-Derived iPSCs Identify Impaired NEPHRIN Localization and Slit Diaphragm Formation in Kidney Podocytes. Stem Cell Reports, 2018, 11, 727-740.	2.3	113
18	Efficient Targeted Mutagenesis in Medaka Using Custom-Designed Transcription Activator-Like Effector Nucleases. Genetics, 2013, 193, 739-749.	1.2	102

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19	Stochastic promoter activation affects Nanog expression variability in mouse embryonic stem cells. Scientific Reports, 2014, 4, 7125.	1.6	97
20	Scleraxis is a transcriptional activator that regulates the expression of Tenomodulin, a marker of mature tenocytes and ligamentocytes. Scientific Reports, 2018, 8, 3155.	1.6	95
21	C-Type Lectin Receptor DCAR Recognizes Mycobacterial Phosphatidyl-Inositol Mannosides to Promote a Th1 Response during Infection. Immunity, 2016, 45, 1245-1257.	6.6	80
22	Cas9, Cpf1 and C2c1/2/3―What's next?. Bioengineered, 2017, 8, 265-273.	1.4	80
23	Highly efficient biallelic genome editing of human ES/iPS cells using a CRISPR/Cas9 or TALEN system. Nucleic Acids Research, 2017, 45, 5198-5207.	6.5	80
24	Unliganded Thyroid Hormone Receptor α Regulates Developmental Timing via Gene Repression in Xenopus tropicalis. Endocrinology, 2015, 156, 735-744.	1.4	78
25	Systematic Cellular Disease Models Reveal Synergistic Interaction of Trisomy 21 and GATA1 Mutations in Hematopoietic Abnormalities. Cell Reports, 2016, 15, 1228-1241.	2.9	78
26	Depdc5 knockout rat: A novel model of mTORopathy. Neurobiology of Disease, 2016, 89, 180-189.	2.1	78
27	Efficient gene targeting by TAL effector nucleases coinjected with exonucleases in zygotes. Scientific Reports, 2013, 3, 1253.	1.6	76
28	Homologous Recombination-Independent Large Gene Cassette Knock-in in CHO Cells Using TALEN and MMEJ-Directed Donor Plasmids. International Journal of Molecular Sciences, 2015, 16, 23849-23866.	1.8	76
29	Targeted mutagenesis in the sea urchin embryo using zincâ€finger nucleases. Genes To Cells, 2010, 15, 875-885.	0.5	75
30	High efficiency TALENs enable F0 functional analysis by targeted gene disruption in <i>Xenopus laevis</i> embryos. Biology Open, 2013, 2, 448-452.	0.6	74
31	Establishment of InÂVitro FUS-Associated Familial Amyotrophic Lateral Sclerosis Model Using Human Induced Pluripotent Stem Cells. Stem Cell Reports, 2016, 6, 496-510.	2.3	74
32	Production of Sry knockout mouse using TALEN via oocyte injection. Scientific Reports, 2013, 3, 3136.	1.6	72
33	Tissue-specific and ubiquitous gene knockouts by TALEN electroporation provide new approaches to investigating gene function in <i>Ciona</i> . Development (Cambridge), 2014, 141, 481-487.	1.2	70
34	Gene cassette knock-in in mammalian cells and zygotes by enhanced MMEJ. BMC Genomics, 2016, 17, 979.	1.2	66
35	Genome Editing in Mouse Spermatogonial Stem Cell Lines Using TALEN and Double-Nicking CRISPR/Cas9. Stem Cell Reports, 2015, 5, 75-82.	2.3	65
36	Highly efficient targeted mutagenesis in one-cell mouse embryos mediated by the TALEN and CRISPR/Cas systems. Scientific Reports, 2015, 4, 5705.	1.6	64

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37	Nucleaseâ€mediated genome editing: At the frontâ€line of functional genomics technology. Development Growth and Differentiation, 2014, 56, 2-13.	0.6	60
38	Highly multiplexed CRISPRâ€Cas9â€nuclease and Cas9â€nickase vectors for inactivation of hepatitis B virus. Genes To Cells, 2016, 21, 1253-1262.	0.5	55
39	Efficient modification of the myostatin gene in porcine somatic cells and generation of knockout piglets. Molecular Reproduction and Development, 2016, 83, 61-70.	1.0	53
40	TALEN-mediated single-base-pair editing identification of an intergenic mutation upstream of <i>BUB1B</i> as causative of PCS (MVA) syndrome. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 1461-1466.	3.3	52
41	Microhomology-assisted scarless genome editing in human iPSCs. Nature Communications, 2018, 9, 939.	5.8	52
42	Functional Investigation of a Non-coding Variant Associated with Adolescent Idiopathic Scoliosis in Zebrafish: Elevated Expression of the Ladybird Homeobox Gene Causes Body Axis Deformation. PLoS Genetics, 2016, 12, e1005802.	1.5	51
43	A High Excision Potential of TALENs for Integrated DNA of HIV-Based Lentiviral Vector. PLoS ONE, 2015, 10, e0120047.	1.1	48
44	Relative contribution of four nucleases, CtIP, Dna2, Exo1 and Mre11, to the initial step of DNA doubleâ€strand break repair by homologous recombination in both the chicken DT40 and human TK6 cell lines. Genes To Cells, 2015, 20, 1059-1076.	0.5	46
45	Production of knockout mice by DNA microinjection of various CRISPR/Cas9 vectors into freeze-thawed fertilized oocytes. BMC Biotechnology, 2015, 15, 33.	1.7	45
46	Smarcal1 promotes double-strand-break repair by nonhomologous end-joining. Nucleic Acids Research, 2015, 43, 6359-6372.	6.5	42
47	ARHGAP10, which encodes Rho GTPase-activating protein 10, is a novel gene for schizophrenia risk. Translational Psychiatry, 2020, 10, 247.	2.4	42
48	Generation of mutant mice via the CRISPR/Cas9 system using Fokl-dCas9. Scientific Reports, 2015, 5, 11221.	1.6	41
49	Zinc-finger nuclease-mediated targeted insertion of reporter genes for quantitative imaging of gene expression in sea urchin embryos. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 10915-10920.	3.3	40
50	Efficient targeted mutagenesis of the chordate <i><scp>C</scp>iona intestinalis</i> genome with zincâ€finger nucleases. Development Growth and Differentiation, 2012, 54, 535-545.	0.6	39
51	Transcription activatorâ€like effector nucleases efficiently disrupt the target gene in Iberian ribbed newts ( <i><scp>P</scp>leurodeles waltl</i> ), an experimental model animal for regeneration. Development Growth and Differentiation, 2014, 56, 115-121.	0.6	38
52	Quantitative assay for TALEN activity at endogenous genomic loci. Biology Open, 2013, 2, 363-367.	0.6	36
53	Tailorâ€made TALEN system for highly efficient targeted gene replacement in the rice blast fungus. Biotechnology and Bioengineering, 2015, 112, 1335-1342.	1.7	36
54	ALC1/CHD1L, a chromatin-remodeling enzyme, is required for efficient base excision repair. PLoS ONE, 2017, 12, e0188320.	1.1	34

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55	PAX2 is dispensable for in vitro nephron formation from human induced pluripotent stem cells. Scientific Reports, 2017, 7, 4554.	1.6	32
56	Targeted knock-in of an scFv-Fc antibody gene into the hprt locus of Chinese hamster ovary cells using CRISPR/Cas9 and CRIS-PITCh systems. Journal of Bioscience and Bioengineering, 2018, 125, 599-605.	1.1	32
57	Efficient genome engineering using Platinum TALEN in potato. Plant Biotechnology, 2019, 36, 167-173.	0.5	32
58	EDEM2 stably disulfide-bonded to TXNDC11 catalyzes the first mannose trimming step in mammalian glycoprotein ERAD. ELife, 2020, 9, .	2.8	31
59	Targeted mutagenesis of multiple and paralogous genes in <i>Xenopus laevis</i> using two pairs of transcription activatorâ€like effector nucleases. Development Growth and Differentiation, 2014, 56, 108-114.	0.6	30
60	Establishment of knockout adult sea urchins by using a CRISPR as9 system. Development Growth and Differentiation, 2019, 61, 378-388.	0.6	30
61	<i>In vivo</i> tracking of histone H3 lysine 9 acetylation in <i>Xenopus laevis</i> during tail regeneration. Genes To Cells, 2016, 21, 358-369.	0.5	29
62	Murine neonatal ketogenesis preserves mitochondrial energetics by preventing protein hyperacetylation. Nature Metabolism, 2021, 3, 196-210.	5.1	29
63	Germ cell mutations of the ascidian <i>Ciona intestinalis</i> with TALE nucleases. Genesis, 2014, 52, 431-439.	0.8	28
64	Screening Methods to Identify TALEN-Mediated Knockout Mice. Experimental Animals, 2014, 63, 79-84.	0.7	28
65	Down syndrome-associated haematopoiesis abnormalities created by chromosome transfer and genome editing technologies. Scientific Reports, 2014, 4, 6136.	1.6	28
66	Hox10-regulated endodermal cell migration is essential for development of the ascidian intestine. Developmental Biology, 2015, 403, 43-56.	0.9	28
67	Ultra-superovulation for the CRISPR-Cas9-mediated production of gene-knockout, single-amino-acid-substituted, and floxed mice. Biology Open, 2016, 5, 1142-1148.	0.6	28
68	Establishment of expanded and streamlined pipeline of PITCh knock-in – a web-based design tool for MMEJ-mediated gene knock-in, PITCh designer, and the variations of PITCh, PITCh-TG and PITCh-KIKO. Bioengineered, 2017, 8, 302-308.	1.4	28
69	Tailor-made gene silencing of Staphylococcus aureus clinical isolates by CRISPR interference. PLoS ONE, 2018, 13, e0185987.	1.1	28
70	Magic wands of CRISPR—lots of choices for gene knock-in. Cell Biology and Toxicology, 2017, 33, 501-505.	2.4	27
71	DJ-1 is indispensable for the S-nitrosylation of Parkin, which maintains function of mitochondria. Scientific Reports, 2020, 10, 4377.	1.6	27
72	The 3′UTR of nanos2 directs enrichment in the germ cell lineage of the sea urchin. Developmental Biology, 2013, 377, 275-283.	0.9	26

5

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73	Detailed analysis of targeted gene mutations caused by the Platinum-Fungal TALENs in Aspergillus oryzae RIB40 strain and a ligD disruptant. Journal of Bioscience and Bioengineering, 2017, 123, 287-293.	1.1	26
74	Humanized UGT2 and CYP3A transchromosomic rats for improved prediction of human drug metabolism. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 3072-3081.	3.3	26
75	Electroporation-mediated genome editing in vitrified/warmed mouse zygotes created by IVF via ultra-superovulation. Experimental Animals, 2018, 67, 535-543.	0.7	25
76	Modification of single-nucleotide polymorphism in a fully humanized CYP3A mouse by genome editing technology. Scientific Reports, 2017, 7, 15189.	1.6	24
77	Biased genome editing using the local accumulation of DSB repair molecules system. Nature Communications, 2018, 9, 3270.	5.8	23
78	Activin Is Superior to BMP7 for Efficient Maintenance of Human iPSC-Derived Nephron Progenitors. Stem Cell Reports, 2019, 13, 322-337.	2.3	23
79	Differential transactivation of the upstream aggrecan enhancer regulated by PAX1/9 depends on SOX9-driven transactivation. Scientific Reports, 2019, 9, 4605.	1.6	23
80	Loss of HCN1 subunits causes absence epilepsy in rats. Brain Research, 2019, 1706, 209-217.	1.1	23
81	Efficient and multiplexable genome editing using Platinum TALENs in oleaginous microalga, <i>Nannochloropsis oceanica</i> NIESâ€2145. Genes To Cells, 2020, 25, 695-702.	0.5	23
82	Targeted mutagenesis in sea urchin embryos using TALENs. Development Growth and Differentiation, 2014, 56, 92-97.	0.6	21
83	Transcriptional regulation of a horizontally transferred gene from bacterium to chordate. Proceedings of the Royal Society B: Biological Sciences, 2016, 283, 20161712.	1.2	20
84	Germ cell regeneration-mediated, enhanced mutagenesis in the ascidian Ciona intestinalis reveals flexible germ cell formation from different somatic cells. Developmental Biology, 2017, 423, 111-125.	0.9	20
85	Acceleration of cancer science with genome editing and related technologies. Cancer Science, 2018, 109, 3679-3685.	1.7	20
86	<scp>TALEN</scp> â€induced gene knock out in <i><scp>D</scp>rosophila</i> . Development Growth and Differentiation, 2014, 56, 86-91.	0.6	19
87	Desmocollin-2 alone forms functional desmosomal plaques, with the plaque formation requiring the juxtamembrane region and plakophilins. Journal of Biochemistry, 2015, 158, 339-353.	0.9	19
88	Functional consequence of fibulin-4 missense mutations associated with vascular and skeletal abnormalities and cutis laxa. Matrix Biology, 2016, 56, 132-149.	1.5	19
89	GABA-Induced GnRH Release Triggers Chordate Metamorphosis. Current Biology, 2020, 30, 1555-1561.e4.	1.8	19
90	The third type III module of human fibronectin mediates cell adhesion and migration. Journal of Biochemistry, 2010, 147, 327-335.	0.9	18

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91	KLF1 mutation E325K induces cell cycle arrest in erythroid cells differentiated from congenital dyserythropoietic anemia patient-specific induced pluripotent stem cells. Experimental Hematology, 2019, 73, 25-37.e8.	0.2	17
92	Targeted gene disruption by use of transcription activator-like effector nuclease (TALEN) in the water flea Daphnia pulex. BMC Biotechnology, 2014, 14, 95.	1.7	16
93	Involvement of aspartoacylase in tremor expression in rats. Experimental Animals, 2016, 65, 293-301.	0.7	16
94	The Expression of TALEN before Fertilization Provides a Rapid Knock-Out Phenotype in Xenopus laevis Founder Embryos. PLoS ONE, 2015, 10, e0142946.	1.1	15
95	Hox-mediated endodermal identity patterns the pharyngeal muscle formation in the chordate pharynx. Development (Cambridge), 2017, 144, 1629-1634.	1.2	15
96	Generation of D1-1 TALEN isogenic control cell line from Dravet syndrome patient iPSCs using TALEN-mediated editing of the SCN1A gene. Stem Cell Research, 2018, 28, 100-104.	0.3	15
97	PDIP38/PolDIP2 controls the DNA damage tolerance pathways by increasing the relative usage of translesion DNA synthesis over template switching. PLoS ONE, 2019, 14, e0213383.	1.1	15
98	Culture time of vitrified/warmed zygotes before microinjection affects the production efficiency of CRISPR-Cas9-mediated knock-in mice. Biology Open, 2017, 6, 706-713.	0.6	14
99	PLEKHN1 promotes apoptosis by enhancing Bax-Bak hetro-oligomerization through interaction with Bid in human colon cancer. Cell Death Discovery, 2018, 4, 11.	2.0	14
100	Targeted mutagenesis of the ryanodine receptor by Platinum TALENs causes slow swimming behaviour in Pacific bluefin tuna (Thunnus orientalis). Scientific Reports, 2019, 9, 13871.	1.6	14
101	Engineering Customized TALENs Using the Platinum Gate TALEN Kit. Methods in Molecular Biology, 2016, 1338, 61-70.	0.4	13
102	Identification of a cell-penetrating peptide applicable to a protein-based transcription activator-like effector expression system for cell engineering. Biomaterials, 2018, 173, 11-21.	5.7	13
103	Three multi-allelic gene pairs are responsible for self-sterility in the ascidian Ciona intestinalis. Scientific Reports, 2020, 10, 2514.	1.6	13
104	Versatile strategy for isolating transcription activatorâ€like effector nucleaseâ€mediated knockout mutants in <i><scp>C</scp>aenorhabditis elegans</i> . Development Growth and Differentiation, 2014, 56, 78-85.	0.6	12
105	Application of Oocyte Cryopreservation Technology in TALEN-Mediated Mouse Genome Editing. Experimental Animals, 2014, 63, 349-355.	0.7	12
106	All-in-One CRISPR-Cas9/Fokl-dCas9 Vector-Mediated Multiplex Genome Engineering in Cultured Cells. Methods in Molecular Biology, 2017, 1498, 41-56.	0.4	12
107	Establishment of Functional Genomics Pipeline in Mouse Epiblast-Like Tissue by Combining Transcriptomic Analysis and Gene Knockdown/Knockin/Knockout, Using RNA Interference and CRISPR/Cas9. Human Gene Therapy, 2016, 27, 436-450.	1.4	11
108	Generation of and characterization of anti-IL-11 antibodies using newly established Il11-deficient mice. Biochemical and Biophysical Research Communications, 2018, 505, 453-459.	1.0	11

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109	Reinvestigation of Disulfide-bonded Oligomeric Forms of the Unfolded Protein Response Transducer ATF6. Cell Structure and Function, 2020, 45, 9-21.	0.5	11
110	Regenerating islet-derived protein (Reg)3β plays a crucial role in attenuation of ileitis and colitis in mice. Biochemistry and Biophysics Reports, 2020, 21, 100738.	0.7	11
111	Enhancer activity sensitive to the orientation of the gene it regulates in the chordategenome. Developmental Biology, 2013, 375, 79-91.	0.9	10
112	Cancer induction and suppression with transcriptional control and epigenome editing technologies. Journal of Human Genetics, 2018, 63, 187-194.	1.1	10
113	Differential micronucleus frequency in isogenic human cells deficient in DNA repair pathways is a valuable indicator for evaluating genotoxic agents and their genotoxic mechanisms. Environmental and Molecular Mutagenesis, 2018, 59, 529-538.	0.9	10
114	Anephrogenic phenotype induced by SALL1 gene knockout in pigs. Scientific Reports, 2019, 9, 8016.	1.6	10
115	Current Overview of TALEN Construction Systems. Methods in Molecular Biology, 2017, 1630, 25-36.	0.4	10
116	Non-RVD mutations that enhance the dynamics of the TAL repeat array along the superhelical axis improve TALEN genome editing efficacy. Scientific Reports, 2016, 6, 37887.	1.6	9
117	TAp63 represses transcription of MYCN/NCYM gene and its high levels of expression are associated with favorable outcome in neuroblastoma. Biochemical and Biophysical Research Communications, 2019, 518, 311-318.	1.0	9
118	Pathological characteristics of <i>Ccdc85c</i> knockout rats: a rat model of genetic hydrocephalus. Experimental Animals, 2020, 69, 26-33.	0.7	9
119	TALEN-Mediated Gene Editing of slc24a5 (Solute Carrier Family 24, Member 5) in Kawakawa, Euthynnus affinis. Journal of Marine Science and Engineering, 2021, 9, 1378.	1.2	8
120	Single-Cell-State Culture of Human Pluripotent Stem Cells Increases Transfection Efficiency. BioResearch Open Access, 2016, 5, 127-136.	2.6	7
121	Unexpected heterogeneity derived from Cas9 ribonucleoproteinâ€introduced clonal cells at the <i><scp>HPRT</scp>1</i> locus. Genes To Cells, 2018, 23, 255-263.	0.5	7
122	MET Activation by a Macrocyclic Peptide Agonist that Couples to Biological Responses Differently from HGF in a Context-Dependent Manner. International Journal of Molecular Sciences, 2018, 19, 3141.	1.8	6
123	Development of a protein-based system for transient epigenetic repression of immune checkpoint molecule and enhancement of antitumour activity of natural killer cells. British Journal of Cancer, 2020, 122, 823-834.	2.9	6
124	Epithelial DLD-1 Cells with Disrupted E-cadherin Gene Retain the Ability to Form Cell Junctions and Apico-basal Polarity. Cell Structure and Function, 2015, 40, 79-94.	0.5	5
125	Homeolog-specific targeted mutagenesis in Xenopus laevis using TALENs. In Vitro Cellular and Developmental Biology - Animal, 2015, 51, 879-884.	0.7	5
126	Hox13 is essential for formation of a sensory organ at the terminal end of the sperm duct in Ciona. Developmental Biology, 2020, 458, 120-131.	0.9	5

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127	Various strategies of effector accumulation to improve the efficiency of genome editing and derivative methodologies. In Vitro Cellular and Developmental Biology - Animal, 2020, 56, 359-366.	0.7	5
128	FASTâ€id system for enrichment of cells with <scp>TALEN</scp> â€induced mutations and large deletions. Genes To Cells, 2014, 19, 419-431.	0.5	4
129	Nucleotide receptor P2RY4 is required for head formation via induction and maintenance of head organizer in <i>Xenopus laevis</i> . Development Growth and Differentiation, 2019, 61, 186-197.	0.6	4
130	Temporal effects of Notch signaling and potential cooperation with multiple downstream effectors on adenohypophysis cell specification in zebrafish. Genes To Cells, 2016, 21, 492-504.	0.5	3
131	Establishment of pten knockout medaka with transcription activator–like effector nucleases (TALENs) as a model of PTEN deficiency disease. PLoS ONE, 2017, 12, e0186878.	1.1	3
132	HpSumf1 is involved in the activation of sulfatases responsible for regulation of skeletogenesis during sea urchin development. Development Genes and Evolution, 2011, 221, 157-166.	0.4	2
133	TALEN-mediated targeted editing of the GDE5 gene suppresses fibroblastic cell proliferation. Bioscience, Biotechnology and Biochemistry, 2017, 81, 2164-2167.	0.6	1
134	Genetic Tools for Self-Organizing Culture of Mouse Embryonic Stem Cells via Small Regulatory RNA-Mediated Technologies, CRISPR/Cas9, and Inducible RNAi. Methods in Molecular Biology, 2017, 1622, 269-292.	0.4	1
135	Six1 is required for signaling center formation and labialâ€lingual asymmetry in developing lower incisors. Developmental Dynamics, 2020, 249, 1098-1116.	0.8	1
136	Genome editing for dissecting and curing human genetic diseases. Journal of Human Genetics, 2018, 63, 105-105.	1.1	0