

A conditional knockout resource for the genome-wide s

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Transcriptomic analysis of pluripotent stem cells: insights into health and disease. <i>Genome Medicine</i> , 2011, 3, 68.	3.6	7
2	Targeted gene correction of α 1-antitrypsin deficiency in induced pluripotent stem cells. <i>Nature</i> , 2011, 478, 391-394.	13.7	635
3	A scalable pipeline for highly effective genetic modification of a malaria parasite. <i>Nature Methods</i> , 2011, 8, 1078-1082.	9.0	93
4	A resource of vectors and ES cells for targeted deletion of microRNAs in mice. <i>Nature Biotechnology</i> , 2011, 29, 840-845.	9.4	92
5	pTransgenesis: a cross-species, modular transgenesis resource. <i>Development (Cambridge)</i> , 2011, 138, 5451-5458.	1.2	52
6	Mouse genomic variation and its effect on phenotypes and gene regulation. <i>Nature</i> , 2011, 477, 289-294.	13.7	1,461
7	A knockout mouse resource for the biomedical research community. <i>Annals of the New York Academy of Sciences</i> , 2011, 1245, 24-26.	1.8	58
8	Following the genes: a framework for animal modeling of psychiatric disorders. <i>BMC Biology</i> , 2011, 9, 76.	1.7	27
9	Research Highlights. <i>Pharmacogenomics</i> , 2011, 12, 1637-1639.	0.6	1
10	Targeted isolation of cloned genomic regions by recombineering for haplotype phasing and isogenic targeting. <i>Nucleic Acids Research</i> , 2011, 39, e137-e137.	6.5	14
11	Swift, flexible knockouts. <i>Nature Methods</i> , 2011, 8, 617-617.	9.0	0
12	Requirement of argininosuccinate lyase for systemic nitric oxide production. <i>Nature Medicine</i> , 2011, 17, 1619-1626.	15.2	189
13	Rapid-Throughput Skeletal Phenotyping of 100 Knockout Mice Identifies 9 New Genes That Determine Bone Strength. <i>PLoS Genetics</i> , 2012, 8, e1002858.	1.5	73
14	Structural and Functional Characterization of Two Alternative Splicing Variants of Mouse Endothelial Cell-Specific Chemotaxis Regulator (ECSCR). <i>International Journal of Molecular Sciences</i> , 2012, 13, 4920-4936.	1.8	7
15	Culture parameters for stable expansion, genetic modification and germline transmission of rat pluripotent stem cells. <i>Biology Open</i> , 2012, 1, 58-65.	0.6	32
16	Community gene annotation in practice. <i>Database: the Journal of Biological Databases and Curation</i> , 2012, 2012, bas009-bas009.	1.4	26
17	Bromodomain-dependent stage-specific male genome programming by Brdt. <i>EMBO Journal</i> , 2012, 31, 3809-3820.	3.5	216
18	Disruption of Mouse Cenpj, a Regulator of Centriole Biogenesis, Phenocopies Seckel Syndrome. <i>PLoS Genetics</i> , 2012, 8, e1003022.	1.5	84

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19	Modelling Human Regulatory Variation in Mouse: Finding the Function in Genome-Wide Association Studies and Whole-Genome Sequencing. <i>PLoS Genetics</i> , 2012, 8, e1002544.	1.5	16
20	Experimental Validation of <i>Ankrd17</i> and <i>Anapc10</i> , Two Novel Meiotic Genes Predicted by Computational Models in Mice. <i>Biology of Reproduction</i> , 2012, 86, 102.	1.2	5
21	Exploring the elephant: histopathology in high-throughput phenotyping of mutant mice. <i>DMM Disease Models and Mechanisms</i> , 2012, 5, 19-25.	1.2	32
22	Disruption of IFT Complex A Causes Cystic Kidneys without Mitotic Spindle Misorientation. <i>Journal of the American Society of Nephrology: JASN</i> , 2012, 23, 641-651.	3.0	103
23	New insights into behaviour using mouse ENU mutagenesis. <i>Human Molecular Genetics</i> , 2012, 21, R72-R81.	1.4	27
24	Germline ablation of SMUG1 DNA glycosylase causes loss of 5-hydroxymethyluracil- and UNG-backup uracil-excision activities and increases cancer predisposition of <i>Ung^Δ/Δ⁺Msh2^Δ/Δ⁺</i> mice. <i>Nucleic Acids Research</i> , 2012, 40, 6016-6025.	6.5	89
25	Mixture of differentially tagged Tol2 transposons accelerates conditional disruption of a broad spectrum of genes in mouse embryonic stem cells. <i>Nucleic Acids Research</i> , 2012, 40, e97-e97.	6.5	10
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27	Lost in Transgenesis. <i>Circulation Research</i> , 2012, 111, 761-777.	2.0	92
28	Generation of mouse mutants as tools in dissecting the molecular clock. <i>Progress in Brain Research</i> , 2012, 199, 247-265.	0.9	1
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30	Functional genomic resources. <i>Nature Methods</i> , 2012, 9, 35-35.	9.0	1
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32	Cardiac-Specific Inducible and Conditional Gene Targeting in Mice. <i>Circulation Research</i> , 2012, 110, 1498-1512.	2.0	61
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40	Conditional control of gene function by an invertible gene trap in zebrafish. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 15389-15394.	3.3	66
41	Isolation of homozygous mutant mouse embryonic stem cells using a dual selection system. <i>Nucleic Acids Research</i> , 2012, 40, e21-e21.	6.5	21
42	Transposon-mediated BAC transgenesis in human ES cells. <i>Nucleic Acids Research</i> , 2012, 40, e150-e150.	6.5	109
43	Clone DB: an integrated NCBI resource for clone-associated data. <i>Nucleic Acids Research</i> , 2012, 41, D1070-D1078.	6.5	14
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48	The critical role of histone H2A-deubiquitinase Mym1 in hematopoiesis and lymphocyte differentiation. <i>Blood</i> , 2012, 119, 1370-1379.	0.6	87
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50	Sphingosine-1-phosphate lyase expression in embryonic and adult murine tissues. <i>Journal of Lipid Research</i> , 2012, 53, 1920-1931.	2.0	40
51	Accessing and Mining Data from Large-Scale Mouse Phenotyping Projects. <i>International Review of Neurobiology</i> , 2012, 104, 47-70.	0.9	9
52	Recombination-Mediated Genetic Engineering of <i>Plasmodium berghei</i> DNA. <i>Methods in Molecular Biology</i> , 2012, 923, 127-138.	0.4	27
53	Comparison of male chimeric mice generated from microinjection of JM8.N4 embryonic stem cells into C57BL/6J and C57BL/6NTac blastocysts. <i>Transgenic Research</i> , 2012, 21, 1149-1158.	1.3	8
54	Computational tools for comparative phenomics: the role and promise of ontologies. <i>Mammalian Genome</i> , 2012, 23, 669-679.	1.0	19

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87	De Novo Origin of Protein-Coding Genes in Murine Rodents. <i>PLoS ONE</i> , 2012, 7, e48650.	1.1	45
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