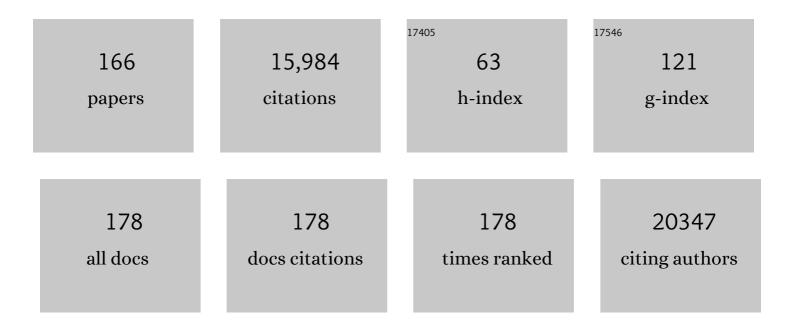
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Immune-responsive gene 1 protein links metabolism to immunity by catalyzing itaconic acid production. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 7820-7825.	3.3	765
2	Waardenburg's syndrome patients have mutations in the human homologue of the Pax-3 paired box gene. Nature, 1992, 355, 635-636.	13.7	688
3	Genome-wide, large-scale production of mutant mice by ENU mutagenesis. Nature Genetics, 2000, 25, 444-447.	9.4	658
4	Antagonistic Interactions between FGF and BMP Signaling Pathways: A Mechanism for Positioning the Sites of Tooth Formation. Cell, 1997, 90, 247-255.	13.5	560
5	Pax: A murine multigene family of paired box-containing genes. Genomics, 1991, 11, 424-434.	1.3	424
6	Degree of methylation of transgenes is dependent on gamete of origin. Nature, 1987, 328, 251-254.	13.7	411
7	De Novo Mutations in Synaptic Transmission Genes Including DNM1 Cause Epileptic Encephalopathies. American Journal of Human Genetics, 2014, 95, 360-370.	2.6	388
8	Variations of cervical vertebrate after expression of a Hox-1.1 transgene in mice. Cell, 1990, 61, 301-308.	13.5	382
9	Pax genes and organogenesis. BioEssays, 1997, 19, 755-765.	1.2	360
10	Impaired insulin secretory capacity in mice lacking a functional vitamin D receptor. FASEB Journal, 2003, 17, 1-14.	0.2	360
11	undulated, a mutation affecting the development of the mouse skeleton, has a point mutation in the paired box of Pax 1. Cell, 1988, 55, 531-535.	13.5	332
12	Teeth: where and how to make them. Trends in Genetics, 1999, 15, 59-65.	2.9	331
13	From hype to reality: data science enabling personalized medicine. BMC Medicine, 2018, 16, 150.	2.3	278
14	Deletion of Deoxyribonucleic Acid Binding Domain of the Vitamin D Receptor Abrogates Genomic and Nongenomic Functions of Vitamin D. Molecular Endocrinology, 2002, 16, 1524-1537.	3.7	267
15	Characterization and Developmental Expression of Pax9, a Paired-Box-Containing Gene Related to Pax1. Developmental Biology, 1995, 170, 701-716.	0.9	266
16	Craniofacial abnormalities induced by ectopic expression of the homeobox gene Hox-1.1 in transgenic mice. Cell, 1989, 58, 337-347.	13.5	264
17	Beethoven, a mouse model for dominant, progressive hearing loss DFNA36. Nature Genetics, 2002, 30, 257-258.	9.4	246
18	Integrating Pathways of Parkinson's Disease in a Molecular Interaction Map. Molecular Neurobiology, 2014, 49, 88-102.	1.9	231

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19	Revolutionizing medicine in the 21 st century through systems approaches. Biotechnology Journal, 2012, 7, 992-1001.	1.8	225
20	1Â,25-dihydroxyvitamin D3 is a potent suppressor of interferon Â-mediated macrophage activation. Blood, 2005, 106, 4351-4358.	0.6	221
21	The hallmarks of <scp>P</scp> arkinson's disease. FEBS Journal, 2013, 280, 5981-5993.	2.2	214
22	The Notch ligand Jagged1 is required for inner ear sensory development. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 3873-3878.	3.3	206
23	Humanized Mice for Modeling Human Infectious Disease: Challenges, Progress, and Outlook. Cell Host and Microbe, 2009, 6, 5-9.	5.1	202
24	Overexpression of activin A in the skin of transgenic mice reveals new activities of activin in epidermal morphogenesis, dermal fibrosis and wound repair. EMBO Journal, 1999, 18, 5205-5215.	3.5	195
25	The European dimension for the mouse genome mutagenesis program. Nature Genetics, 2004, 36, 925-927.	9.4	195
26	Neurodegeneration by Activation of the Microglial Complement-Phagosome Pathway. Journal of Neuroscience, 2014, 34, 8546-8556.	1.7	192
27	Making sense of big data in health research: Towards an EU action plan. Genome Medicine, 2016, 8, 71.	3.6	190
28	Singleâ€cell transcriptomics reveals distinct inflammationâ€induced microglia signatures. EMBO Reports, 2018, 19, .	2.0	186
29	Systems medicine and integrated care to combat chronic noncommunicable diseases. Genome Medicine, 2011, 3, 43.	3.6	181
30	Mutations in STX1B, encoding a presynaptic protein, cause fever-associated epilepsy syndromes. Nature Genetics, 2014, 46, 1327-1332.	9.4	178
31	The P4 Health Spectrum – A Predictive, Preventive, Personalized and Participatory Continuum for Promoting Healthspan. Progress in Cardiovascular Diseases, 2017, 59, 506-521.	1.6	178
32	Diseases as network perturbations. Current Opinion in Biotechnology, 2010, 21, 566-571.	3.3	167
33	The ventralizing effect of the notochord on somite differentiation in chick embryos. Anatomy and Embryology, 1993, 188, 239-45.	1.5	163
34	A gene for autosomal dominant sacral agenesis maps to the holoprosencephaly region at 7q36. Nature Genetics, 1995, 11, 93-95.	9.4	150
35	Global implementation of genomic medicine: We are not alone. Science Translational Medicine, 2015, 7, 290ps13.	5.8	146
36	Cloning and Expression Analysis of a Novel Mesodermally Expressed Cadherin. Developmental Biology, 1995, 169, 337-346.	0.9	132

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37	A novel Fanconi anaemia subtype associated with a dominant-negative mutation in RAD51. Nature Communications, 2015, 6, 8829.	5.8	130
38	The SOX10 / Sox10 gene from human and mouse: sequence, expression, and transactivation by the encoded HMG domain transcription factor. Human Genetics, 1998, 103, 115-123.	1.8	129
39	Pax1 and Pax9 activate Bapx1 to induce chondrogenic differentiation in the sclerotome. Development (Cambridge), 2003, 130, 473-482.	1.2	128
40	Systems genomics evaluation of the SH-SY5Y neuroblastoma cell line as a model for Parkinson's disease. BMC Genomics, 2014, 15, 1154.	1.2	126
41	Functional Annotation of Mouse Genome Sequences. Science, 2001, 291, 1251-1255.	6.0	125
42	ENU MUTAGENESIS: Analyzing Gene Function in Mice. Annual Review of Genomics and Human Genetics, 2001, 2, 463-492.	2.5	122
43	Parkinson's disease-associated alterations of the gut microbiome predict disease-relevant changes in metabolic functions. BMC Biology, 2020, 18, 62.	1.7	122
44	GARP: a key receptor controlling FOXP3 in human regulatory T cells. Journal of Cellular and Molecular Medicine, 2009, 13, 3343-3357.	1.6	113
45	Large scale ENU screens in the mouse: genetics meets genomics. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 1998, 400, 25-32.	0.4	108
46	Pax-1, a regulator of sclerotome development is induced by notochord and floor plate signals in avian embryos. Anatomy and Embryology, 1995, 191, 297-310.	1.5	105
47	A mouse model for valproate teratogenicity: parental effects, homeotic transformations, and altered HOX expression. Human Molecular Genetics, 2000, 9, 227-236.	1.4	103
48	Expression of AvianPax1andPax9Is Intrinsically Regulated in the Pharyngeal Endoderm, but Depends on Environmental Influences in the Paraxial Mesoderm. Developmental Biology, 1996, 178, 403-417.	0.9	102
49	Conditional inactivation of Sox9: A mouse model for campomelic dysplasia. Genesis, 2002, 32, 121-123.	0.8	99
50	COVID-19 Disease Map, building a computational repository of SARS-CoV-2 virus-host interaction mechanisms. Scientific Data, 2020, 7, 136.	2.4	99
51	Targeted Disruption of the Peptide Transporter Pept2 Gene in Mice Defines Its Physiological Role in the Kidney. Molecular and Cellular Biology, 2003, 23, 3247-3252.	1.1	96
52	Recessive mutations in <i>SLC13A5</i> result in a loss of citrate transport and cause neonatal epilepsy, developmental delay and teeth hypoplasia. Brain, 2015, 138, 3238-3250.	3.7	96
53	Notochord-Dependent Expression of MFH1 and PAX1 Cooperates to Maintain the Proliferation of Sclerotome Cells during the Vertebral Column Development. Developmental Biology, 1999, 210, 15-29.	0.9	95
54	Hydrogen Peroxide-Mediated Killing of Caenorhabditis elegans by Streptococcus pyogenes. Infection and Immunity, 2002, 70, 5202-5207.	1.0	87

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55	Critical transitions in chronic disease: transferring concepts from ecology to systems medicine. Current Opinion in Biotechnology, 2015, 34, 48-55.	3.3	86
56	Systems medicine disease maps: community-driven comprehensive representation of disease mechanisms. Npj Systems Biology and Applications, 2018, 4, 21.	1.4	84
57	Evaluating the Use of Circulating MicroRNA Profiles for Lung Cancer Detection in Symptomatic Patients. JAMA Oncology, 2020, 6, 714.	3.4	84
58	Pax genes and organogenesis: <i>Pax9</i> meets tooth development. European Journal of Oral Sciences, 1998, 106, 38-43.	0.7	83
59	A Recurrent Missense Variant in AP2M1 Impairs Clathrin-Mediated Endocytosis and Causes Developmental and Epileptic Encephalopathy. American Journal of Human Genetics, 2019, 104, 1060-1072.	2.6	78
60	Zic1 regulates the patterning of vertebral arches in cooperation with Gli3. Mechanisms of Development, 1999, 89, 141-150.	1.7	76
61	Interaction between undulated and Patch leads to an extreme form of spina bifida in double-mutant mice. Nature Genetics, 1995, 11, 60-63.	9.4	75
62	Mice, microbes and models of infection. Nature Reviews Genetics, 2003, 4, 195-205.	7.7	75
63	Thymopoiesis requiresPax9 function in thymic epithelial cells. European Journal of Immunology, 2002, 32, 1175-1181.	1.6	74
64	Loss of SYNJ1 dual phosphatase activity leads to early onset refractory seizures and progressive neurological decline. Brain, 2016, 139, 2420-2430.	3.7	70
65	MINERVA—a platform for visualization and curation of molecular interaction networks. Npj Systems Biology and Applications, 2016, 2, 16020.	1.4	68
66	Mutation in the βA3/A1-Crystallin Encoding Gene Cryba1 Causes a Dominant Cataract in the Mouse. Genomics, 1999, 62, 67-73.	1.3	67
67	Reverse engineering and verification of gene networks: Principles, assumptions, and limitations of present methods and future perspectives. Journal of Biotechnology, 2009, 144, 190-203.	1.9	67
68	Gene Regulatory Network Inference of Immunoresponsive Gene 1 (IRG1) Identifies Interferon Regulatory Factor 1 (IRF1) as Its Transcriptional Regulator in Mammalian Macrophages. PLoS ONE, 2016, 11, e0149050.	1.1	66
69	Parkinson's disease mouse models in translational research. Mammalian Genome, 2011, 22, 401-419.	1.0	65
70	Systems Medicine Approaches for the Definition of Complex Phenotypes in Chronic Diseases and Ageing. From Concept to Implementation and Policies. Current Pharmaceutical Design, 2014, 20, 5928-5944.	0.9	63
71	Progressive loss of PAX9 expression correlates with increasing malignancy of dysplastic and cancerous epithelium of the human oesophagus. Journal of Pathology, 2002, 197, 293-297.	2.1	60
72	Respiratory mechanics in mice: strain and sex specific differences. Acta Physiologica Scandinavica, 2002, 174, 367-375.	2.3	58

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73	The role of regulatory T cells in neurodegenerative diseases. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2013, 5, 153-180.	6.6	58
74	The Luxembourg Parkinson's Study: A Comprehensive Approach for Stratification and Early Diagnosis. Frontiers in Aging Neuroscience, 2018, 10, 326.	1.7	57
75	PLAU inferred from a correlation network is critical for suppressor function of regulatory T cells. Molecular Systems Biology, 2012, 8, 624.	3.2	54
76	The clinical-chemical screen in the Munich ENU Mouse Mutagenesis Project: screening for clinically relevant phenotypes. Mammalian Genome, 2000, 11, 543-546.	1.0	53
77	COVID19 Disease Map, a computational knowledge repository of virus–host interaction mechanisms. Molecular Systems Biology, 2021, 17, e10387.	3.2	53
78	Characterization of a Mutation in the Lens-specific MP70 Encoding Gene of the Mouse Leading to a Dominant Cataract. Experimental Eye Research, 2001, 73, 867-876.	1.2	52
79	The battle of two genomes: genetics of bacterial host/pathogen interactions in mice. Mammalian Genome, 2001, 12, 261-271.	1.0	52
80	Analysis of limb patterning in BMP-7-deficient mice. Genesis, 1996, 19, 43-50.	3.1	51
81	A new Pax gene, Pax-9, maps to mouse Chromosome 12. Mammalian Genome, 1993, 4, 354-358.	1.0	50
82	ROS networks: designs, aging, Parkinson's disease and precision therapies. Npj Systems Biology and Applications, 2020, 6, 34.	1.4	50
83	Similar αâ€5ynuclein staining in the colon mucosa in patients with Parkinson's disease and controls. Movement Disorders, 2016, 31, 1567-1570.	2.2	48
84	Influenza H3N2 infection of the collaborative cross founder strains reveals highly divergent host responses and identifies a unique phenotype in CAST/EiJ mice. BMC Genomics, 2016, 17, 143.	1.2	48
85	Community-driven roadmap for integrated disease maps. Briefings in Bioinformatics, 2019, 20, 659-670.	3.2	48
86	Common diseases alter the physiological age-related blood microRNA profile. Nature Communications, 2020, 11, 5958.	5.8	46
87	A roadmap towards personalized immunology. Npj Systems Biology and Applications, 2018, 4, 9.	1.4	43
88	A rare loss-of-function variant of ADAM17 is associated with late-onset familial Alzheimer disease. Molecular Psychiatry, 2020, 25, 629-639.	4.1	42
89	Integration and Visualization of Translational Medicine Data for Better Understanding of Human Diseases. Big Data, 2016, 4, 97-108.	2.1	41
90	The large-scale Munich ENU-mouse-mutagenesis screen. Mammalian Genome, 2000, 11, 507-510.	1.0	40

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91	Toward Omics-Based, Systems Biomedicine, and Path and Drug Discovery Methodologies for Depression-Inflammation Research. Molecular Neurobiology, 2016, 53, 2927-2935.	1.9	40
92	Reliable recovery of inbred mouse lines using cryopreserved spermatozoa. Mammalian Genome, 1999, 10, 773-776.	1.0	38
93	Screening for dysmorphological abnormalities—a powerful tool to isolate new mouse mutants. Mammalian Genome, 2000, 11, 528-530.	1.0	38
94	Reduced intragraft mRNA expression of matrix metalloproteinases Mmp3, Mmp12, Mmp13 and Adam8, and diminished transplant arteriosclerosis in Ccr5-deficient mice. European Journal of Immunology, 2004, 34, 2568-2578.	1.6	38
95	Tailchaser (Tlc): a new mouse mutation affecting hair bundle differentiation and hair cell survival. Journal of Neurocytology, 1999, 28, 969-985.	1.6	37
96	MICâ€MAC: An automated pipeline for highâ€throughput characterization and classification of threeâ€dimensional microglia morphologies in mouse and human postmortem brain samples. Glia, 2019, 67, 1496-1509.	2.5	36
97	A locus for radiation-induced gastroschisis on mouse Chromosome 7. Mammalian Genome, 1998, 9, 995-997.	1.0	35
98	Murine genes with homology to Drosophila segmentation genes. Development (Cambridge), 1988, 104, 181-186.	1.2	35
99	In Situ Gene Expression Analysis During BMP2-induced Ectopic Bone Formation in Mice Shows Simultaneous Endochondral and Intramembranous Ossification. Growth Factors, 2003, 20, 197-210.	0.5	33
100	Rare GABRA3 variants are associated with epileptic seizures, encephalopathy and dysmorphic features. Brain, 2017, 140, 2879-2894.	3.7	33
101	Emergence of the silicon human and network targeting drugs. European Journal of Pharmaceutical Sciences, 2012, 46, 190-197.	1.9	32
102	A Systems Medicine Clinical Platform for Understanding and Managing Non- Communicable Diseases. Current Pharmaceutical Design, 2014, 20, 5945-5956.	0.9	32
103	The biochemical metabolite screen in the Munich ENU Mouse Mutagenesis Project: determination of amino acids and acylcarnitines by tandem mass spectrometry. Mammalian Genome, 2000, 11, 547-551.	1.0	31
104	The Mouse Brain Metabolome. American Journal of Pathology, 2015, 185, 1699-1712.	1.9	31
105	Spatio-temporal distribution of chondromodulin-I mRNA in the chicken embryo: Expression during cartilage development and formation of the heart and eye. , 1999, 216, 233-243.		30
106	Neurodegeneration and neuroinflammation are linked, but independent of alphaâ€synuclein inclusions, in a seeding/spreading mouse model of Parkinson's disease. Glia, 2022, 70, 935-960.	2.5	30
107	Structure, expression and chromosomal localization of Zfp-1, a murine zinc finger protein gene. Nucleic Acids Research, 1989, 17, 10427-10438.	6.5	28
108	PAX1, a member of the paired box-containing class of developmental control genes, is mapped to human chromosome 20p11.2 by in Situ hybridization (ISH and FISH). Genomics, 1992, 14, 740-744.	1.3	28

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109	Inhibitory action of BMPs onPax1 expression and on shoulder girdle formation during limb development. Developmental Dynamics, 1998, 213, 199-206.	0.8	27
110	Deep sequencing of sncRNAs reveals hallmarks and regulatory modules of the transcriptome during Parkinson's disease progression. Nature Aging, 2021, 1, 309-322.	5.3	26
111	Pax Genes and Skeletal Development. Annals of the New York Academy of Sciences, 1996, 785, 27-33.	1.8	25
112	Pax genes and sclerotome development. Seminars in Cell and Developmental Biology, 1996, 7, 129-136.	2.3	25
113	From Systems Biology to Systems Biomedicine. Current Opinion in Biotechnology, 2012, 23, 604-608.	3.3	24
114	V76D mutation in a conserved gD-crystallin region leads to dominant cataracts in mice. Mammalian Genome, 2002, 13, 452-455.	1.0	23
115	Identification of immunological relevant phenotypes in ENU mutagenized mice. Mammalian Genome, 2000, 11, 526-527.	1.0	22
116	Exome-wide analysis of mutational burden in patients with typical and atypical Rolandic epilepsy. European Journal of Human Genetics, 2018, 26, 258-264.	1.4	22
117	A comprehensive and comparative phenotypic analysis of the collaborative founder strains identifies new and known phenotypes. Mammalian Genome, 2020, 31, 30-48.	1.0	22
118	Cadherin-11 is highly expressed in rhabdomyosarcomas and during differentiation of myoblastsin vitro. , 1999, 187, 164-172.		19
119	PARK7/DJ-1 promotes pyruvate dehydrogenase activity and maintains Treg homeostasis during ageing. Nature Metabolism, 2022, 4, 589-607.	5.1	18
120	Large-scale N-ethyl-N-nitrosourea mutagenesis of mice - from phenotypes to genes. Experimental Physiology, 2000, 85, 635-643.	0.9	17
121	Comparative analysis of the genomic organization of Pax9 and its conserved physical association with Nkx2-9 in the human, mouse, and pufferfish genomes. Mammalian Genome, 2001, 12, 232-237.	1.0	17
122	Understanding complexity in neurodegenerative diseases: in silico reconstruction of emergence. Frontiers in Physiology, 2012, 3, 291.	1.3	16
123	Recessive mutations in <i>SLC35A3</i> cause early onset epileptic encephalopathy with skeletal defects. American Journal of Medical Genetics, Part A, 2017, 173, 1119-1123.	0.7	16
124	Ethylnitrosourea-Induced Mutation in Mice Leads to the Expression of a Novel Protein in the Eye and to Dominant Cataracts. Genetics, 2001, 157, 1313-1320.	1.2	15
125	Identification of VIMP as a gene inhibiting cytokine production in human CD4+ effector TÂcells. IScience, 2021, 24, 102289.	1.9	14
126	Large-Scale N -Ethyl-N -Nitrosourea Mutagenesis of Mice - from Phenotypes to Genes. Experimental Physiology, 2000, 85, 635-643.	0.9	13

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127	Connecting environmental exposure and neurodegeneration using cheminformatics and high resolution mass spectrometry: potential and challenges. Environmental Sciences: Processes and Impacts, 2019, 21, 1426-1445.	1.7	13
128	Finding and sharing: new approaches to registries of databases and services for the biomedical sciences. Database: the Journal of Biological Databases and Curation, 2010, 2010, baq014-baq014.	1.4	12
129	Platelet mitochondrial membrane potential in Parkinson's disease. Annals of Clinical and Translational Neurology, 2015, 2, 67-73.	1.7	12
130	Rare ABCA7 variants in 2 German families with Alzheimer disease. Neurology: Genetics, 2018, 4, e224.	0.9	12
131	Fine genetic mapping of the proximal part of mouse Chromosome 2 excludes Pax-8 as a candidate gene for Danforth's short tail (Sd). Mammalian Genome, 1993, 4, 324-327.	1.0	11
132	Dynamic cumulative activity of transcription factors as a mechanism of quantitative gene regulation. Genome Biology, 2007, 8, R181.	13.9	11
133	Genetic and molecular control of folate-homocysteine metabolism in mutant mice. Mammalian Genome, 2002, 13, 259-267.	1.0	10
134	The P4 Health Spectrum – A Predictive, Preventive, Personalized and Participatory Continuum for Promoting Healthspan. Progress in Preventive Medicine (New York, N Y), 2017, 2, e0002.	0.7	10
135	Standard Peripheral Blood Mononuclear Cell Cryopreservation Selectively Decreases Detection of Nine Clinically Relevant T Cell Markers. ImmunoHorizons, 2021, 5, 711-720.	0.8	10
136	A nonâ€functioning vitamin D receptor predisposes to leukaemoid reactions in mice. Hematological Oncology, 2010, 28, 185-191.	0.8	9
137	DJâ€1 depletion prevents immunoaging in Tâ€cell compartments. EMBO Reports, 2022, 23, e53302.	2.0	9
138	Molecular basis for skeletal variation: insights from developmental genetic studies in mice. Birth Defects Research Part B: Developmental and Reproductive Toxicology, 2007, 80, 425-450.	1.4	8
139	Mitochondria interaction networks show altered topological patterns in Parkinson's disease. Npj Systems Biology and Applications, 2020, 6, 38.	1.4	7
140	The molecular and genetic analysis of mouse development. FEBS Journal, 1992, 204, 5-11.	0.2	6
141	Accelerating the Development and Validation of New Value-Based Diagnostics by Leveraging Biobanks. Public Health Genomics, 2016, 19, 160-169.	0.6	6
142	Viva Europa, a Land of Excellence in Research and Innovation for Health and Wellbeing. Progress in Preventive Medicine (New York, N Y), 2017, 2, e006.	0.7	6
143	Functional Genomics, Proteomics, Metabolomics and Bioinformatics for Systems Biology. , 2013, , 3-41.		5
144	Large-scale validation of miRNAs by disease association, evolutionary conservation and pathway activity. RNA Biology, 2019, 16, 93-103.	1.5	5

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145	From mouse genetics to systems biology. Mammalian Genome, 2007, 18, 383-388.	1.0	4
146	Maintaining Your Immune SystemOne Method for Enhanced Longevity. Science of Aging Knowledge Environment: SAGE KE, 2004, 2004, 2pe-2.	0.9	4
147	Mapping of the Mod-1 locus on mouse Chromosome 9. Mammalian Genome, 1993, 4, 333-337.	1.0	3
148	From parathyroid to thymus, via glial cells. Nature Medicine, 2000, 6, 860-861.	15.2	3
149	Neurological Diseases from a Systems Medicine Point of View. Methods in Molecular Biology, 2016, 1386, 221-250.	0.4	3
150	Genes associated with Parkinson's disease respond to increasing polychlorinated biphenyl levels in the blood of healthy females. Environmental Pollution, 2019, 250, 107-117.	3.7	3
151	COVID-19 and beyond:Âa call for action andÂaudacious solidarity to all the citizens and nations,Âit is humanity's fight. F1000Research, 0, 9, 1130.	0.8	3
152	On Different Aspects of Network Analysis in Systems Biology. , 2013, , 181-207.		3
153	The genetic map around the tail kinks (tk) locus on mouse Chromosome 9. Mammalian Genome, 1993, 4, 560-564.	1.0	2
154	"4D Biology for health and disease―workshop report. New Biotechnology, 2011, 28, 291-293.	2.4	2
155	Quantitative trait locus mapping identifies a locus linked toÂstriatal dopamine and points to collagen <scp>IV</scp> alphaâ€6 chain asÂa novel regulator of striatal axonal branching in mice. Genes, Brain and Behavior, 2021, 20, e12769.	1.1	2
156	Computational Infrastructures for Data and Knowledge Management in Systems Biology. , 2013, , 377-397.		2
157	Moratorium call. Nature, 1988, 334, 560-560.	13.7	1
158	Development(s) in mouse genetics. Biochemistry and Cell Biology, 1990, 68, 404-407.	0.9	0
159	Whose law for sharing research tools?. Nature, 1998, 396, 509-509.	13.7	Ο
160	From Developmental Biology to Developmental Toxicology. Annals of the New York Academy of Sciences, 2000, 919, 239-245.	1.8	0
161	Der neue VBIO: Weichenstellungen für die Zukunft der Biowissenschaften. Biologie in Unserer Zeit, 2007, 37, 275-275.	0.3	0
162	Sudden and unexpected. Nature Genetics, 2007, 39, 1422-1423.	9.4	0

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163	From Diagnosing Diseases to Predicting Diseases. , 2019, , 95-103.		о
164	Workshop Report: Systems Genetics of Neurodegenerative Disease, a Summer School in Systems Medicine, 25th Augustâ~'1st September 2017. Frontiers in Genetics, 2019, 10, 29.	1.1	0
165	Presenting and sharing clinical data using the eTRIKS Standards Master Tree for tranSMART. Bioinformatics, 2019, 35, 1562-1565.	1.8	0
166	The molecular and genetic analysis of mouse development. , 1993, , 35-41.		0