

Steve D M Brown

List of Publications by Year in descending order

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Version: 2024-02-01

64
papers

4,554
citations

172457

29
h-index

138484

58
g-index

65
all docs

65
docs citations

65
times ranked

8553
citing authors

#	ARTICLE	IF	CITATIONS
1	High-throughput discovery of novel developmental phenotypes. <i>Nature</i> , 2016, 537, 508-514.	27.8	1,001
2	A comparative phenotypic and genomic analysis of C57BL/6J and C57BL/6N mouse strains. <i>Genome Biology</i> , 2013, 14, R82.	9.6	403
3	Genome-Wide Association Study Using Extreme Truncate Selection Identifies Novel Genes Affecting Bone Mineral Density and Fracture Risk. <i>PLoS Genetics</i> , 2011, 7, e1001372.	3.5	233
4	Disease model discovery from 3,328 gene knockouts by The International Mouse Phenotyping Consortium. <i>Nature Genetics</i> , 2017, 49, 1231-1238.	21.4	216
5	Prevalence of sexual dimorphism in mammalian phenotypic traits. <i>Nature Communications</i> , 2017, 8, 15475.	12.8	200
6	A gene-driven ENU-based approach to generating an allelic series in any gene. <i>Mammalian Genome</i> , 2004, 15, 585-591.	2.2	148
7	Analysis of mammalian gene function through broad-based phenotypic screens across a consortium of mouse clinics. <i>Nature Genetics</i> , 2015, 47, 969-978.	21.4	137
8	Novel ENU-induced eye mutations in the mouse: models for human eye disease. <i>Human Molecular Genetics</i> , 2002, 11, 755-767.	2.9	126
9	Genome-wide association of multiple complex traits in outbred mice by ultra-low-coverage sequencing. <i>Nature Genetics</i> , 2016, 48, 912-918.	21.4	124
10	CIB2 interacts with TMC1 and TMC2 and is essential for mechanotransduction in auditory hair cells. <i>Nature Communications</i> , 2017, 8, 43.	12.8	121
11	A large scale hearing loss screen reveals an extensive unexplored genetic landscape for auditory dysfunction. <i>Nature Communications</i> , 2017, 8, 886.	12.8	116
12	Correction of the auditory phenotype in C57BL/6N mice via CRISPR/Cas9-mediated homology directed repair. <i>Genome Medicine</i> , 2016, 8, 16.	8.2	113
13	A mutation in the F-box gene, <i>Fbxo11</i> , causes otitis media in the Jeff mouse. <i>Human Molecular Genetics</i> , 2006, 15, 3273-3279.	2.9	98
14	A hearing and vestibular phenotyping pipeline to identify mouse mutants with hearing impairment. <i>Nature Protocols</i> , 2010, 5, 177-190.	12.0	93
15	Helios is a key transcriptional regulator of outer hair cell maturation. <i>Nature</i> , 2018, 563, 696-700.	27.8	90
16	Mutation at the <i>Evi1</i> Locus in Junbo Mice Causes Susceptibility to Otitis Media. <i>PLoS Genetics</i> , 2006, 2, e149.	3.5	88
17	The Deaf Mouse Mutant Jeff (Jf) is a Single Gene Model of Otitis Media. <i>JARO - Journal of the Association for Research in Otolaryngology</i> , 2003, 4, 130-138.	1.8	79
18	Novel gene function revealed by mouse mutagenesis screens for models of age-related disease. <i>Nature Communications</i> , 2016, 7, 12444.	12.8	79

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19	High-throughput mouse phenomics for characterizing mammalian gene function. <i>Nature Reviews Genetics</i> , 2018, 19, 357-370.	16.3	78
20	Applying the ARRIVE Guidelines to an In Vivo Database. <i>PLoS Biology</i> , 2015, 13, e1002151.	5.6	75
21	Association of the FBXO11 Gene With Chronic Otitis Media With Effusion and Recurrent Otitis Media. <i>JAMA Otolaryngology</i> , 2006, 132, 729.	1.2	65
22	Mouse Models of NMNAT1-Leber Congenital Amaurosis (LCA9) Recapitulate Key Features of the Human Disease. <i>American Journal of Pathology</i> , 2016, 186, 1925-1938.	3.8	61
23	The Functional Annotation of Mammalian Genomes: The Challenge of Phenotyping. <i>Annual Review of Genetics</i> , 2009, 43, 305-333.	7.6	60
24	A resource of targeted mutant mouse lines for 5,061 genes. <i>Nature Genetics</i> , 2021, 53, 416-419.	21.4	60
25	Unraveling the genetics of otitis media: from mouse to human and back again. <i>Mammalian Genome</i> , 2011, 22, 66-82.	2.2	58
26	HIF-1 α -VEGF Pathways Are Critical for Chronic Otitis Media in Junbo and Jeff Mouse Mutants. <i>PLoS Genetics</i> , 2011, 7, e1002336.	3.5	54
27	Otitis media in the Tgfr knockout mouse implicates TGF β 2 signalling in chronic middle ear inflammatory disease. <i>Human Molecular Genetics</i> , 2013, 22, 2553-2565.	2.9	50
28	Gelsolin Plays a Role in the Actin Polymerization Complex of Hair Cell Stereocilia. <i>PLoS ONE</i> , 2010, 5, e11627.	2.5	38
29	A mutation in Nischarin causes otitis media via LIMK1 and NF- κ B pathways. <i>PLoS Genetics</i> , 2017, 13, e1006969.	3.5	36
30	A Wars2 Mutant Mouse Model Displays OXPHOS Deficiencies and Activation of Tissue-Specific Stress Response Pathways. <i>Cell Reports</i> , 2018, 25, 3315-3328.e6.	6.4	35
31	EV11 Acts as an Inducible Negative-Feedback Regulator of NF- κ B by Inhibiting p65 Acetylation. <i>Journal of Immunology</i> , 2012, 188, 6371-6380.	0.8	33
32	Absence of Neuroplastin-65 Affects Synaptogenesis in Mouse Inner Hair Cells and Causes Profound Hearing Loss. <i>Journal of Neuroscience</i> , 2016, 36, 222-234.	3.6	30
33	Age-related changes in the biophysical and morphological characteristics of mouse cochlear outer hair cells. <i>Journal of Physiology</i> , 2020, 598, 3891-3910.	2.9	29
34	Loss of <i>Baiap2l2</i> destabilizes the transducing stereocilia of cochlear hair cells and leads to deafness. <i>Journal of Physiology</i> , 2021, 599, 1173-1198.	2.9	28
35	Advances in mouse genetics for the study of human disease. <i>Human Molecular Genetics</i> , 2021, 30, R274-R284.	2.9	26
36	Biophysical and morphological changes in inner hair cells and their efferent innervation in the ageing mouse cochlea. <i>Journal of Physiology</i> , 2021, 599, 269-287.	2.9	25

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37	Pathophysiological changes in inner hair cell ribbon synapses in the ageing mammalian cochlea. <i>Journal of Physiology</i> , 2020, 598, 4339-4355.	2.9	23
38	Extensive identification of genes involved in congenital and structural heart disorders and cardiomyopathy. , 2022, 1, 157-173.		22
39	Genetic landscape of auditory dysfunction. <i>Human Molecular Genetics</i> , 2018, 27, R130-R135.	2.9	20
40	Interactions between the otitis media gene, <i>Fbxo11</i> , and <i>p53</i> in the mouse embryonic lung. <i>DMM Disease Models and Mechanisms</i> , 2015, 8, 1531-42.	2.4	19
41	Mouse mutant phenotyping at scale reveals novel genes controlling bone mineral density. <i>PLoS Genetics</i> , 2020, 16, e1009190.	3.5	19
42	A genetic modifier suggests that endurance exercise exacerbates Huntington's disease. <i>Human Molecular Genetics</i> , 2018, 27, 1723-1731.	2.9	17
43	N-ethyl-N-Nitrosourea (ENU) Induced Mutations within the <i>Klotho</i> Gene Lead to Ectopic Calcification and Reduced Lifespan in Mouse Models. <i>PLoS ONE</i> , 2015, 10, e0122650.	2.5	16
44	The Actin-Binding Proteins <i>Eps8</i> and <i>Gelsolin</i> Have Complementary Roles in Regulating the Growth and Stability of Mechanosensory Hair Bundles of Mammalian Cochlear Outer Hair Cells. <i>PLoS ONE</i> , 2014, 9, e87331.	2.5	15
45	Cellular content plays a crucial role in Non-typeable <i>Haemophilus influenzae</i> infection of preinflamed <i>Junbo</i> mouse middle ear. <i>Cellular Microbiology</i> , 2019, 21, e12960.	2.1	13
46	Transcript Analysis Reveals a Hypoxic Inflammatory Environment in Human Chronic Otitis Media With Effusion. <i>Frontiers in Genetics</i> , 2019, 10, 1327.	2.3	12
47	Modification of an aggressive model of Alport Syndrome reveals early differences in disease pathogenesis due to genetic background. <i>Scientific Reports</i> , 2019, 9, 20398.	3.3	11
48	Mice with an N-Ethyl-N-Nitrosourea (ENU) Induced Tyr209Asn Mutation in Natriuretic Peptide Receptor 3 (NPR3) Provide a Model for Kyphosis Associated with Activation of the MAPK Signaling Pathway. <i>PLoS ONE</i> , 2016, 11, e0167916.	2.5	11
49	High-throughput discovery of genetic determinants of circadian misalignment. <i>PLoS Genetics</i> , 2020, 16, e1008577.	3.5	10
50	Cellular Immune Response against Non-typeable <i>Haemophilus influenzae</i> Infecting the Preinflamed Middle Ear of the <i>Junbo</i> Mouse. <i>Infection and Immunity</i> , 2019, 87, .	2.2	9
51	A Requirement for <i>Zic2</i> in the Regulation of <i>Nodal</i> Expression Underlies the Establishment of Left-Sided Identity. <i>Scientific Reports</i> , 2018, 8, 10439.	3.3	6
52	Mutation in <i>Fbxo11</i> Leads to Altered Immune Cell Content in Jeff Mouse Model of Otitis Media. <i>Frontiers in Genetics</i> , 2020, 11, 50.	2.3	6
53	Identifying genetic determinants of inflammatory pain in mice using a large-scale gene-targeted screen. <i>Pain</i> , 2022, 163, 1139-1157.	4.2	4
54	Neuroplastin genetically interacts with Cadherin 23 and the encoded isoform Np55 is sufficient for cochlear hair cell function and hearing. <i>PLoS Genetics</i> , 2022, 18, e1009937.	3.5	4

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55	A holistic view of mouse enhancer architectures reveals analogous pleiotropic effects and correlation with human disease. BMC Genomics, 2020, 21, 754.	2.8	3
56	INFRAFRONTIER quality principles in systemic phenotyping. Mammalian Genome, 2021, , 1.	2.2	3
57	Comparative visualization of genotype-phenotype relationships. Nature Methods, 2015, 12, 698-699.	19.0	2
58	The Jeff Mouse Mutant Model for Chronic Otitis Media Manifests Gain-of-Function as Well as Loss-of-Function Effects. Frontiers in Genetics, 2020, 11, 498.	2.3	2
59	Introduction. Mammalian Genome, 2011, 22, 361-361.	2.2	1
60	Introduction to Mammalian Genome Special Issue: Mammalian Genetic Resources. Mammalian Genome, 2022, 33, 1-3.	2.2	0
61	High-throughput discovery of genetic determinants of circadian misalignment. , 2020, 16, e1008577.		0
62	High-throughput discovery of genetic determinants of circadian misalignment. , 2020, 16, e1008577.		0
63	High-throughput discovery of genetic determinants of circadian misalignment. , 2020, 16, e1008577.		0
64	High-throughput discovery of genetic determinants of circadian misalignment. , 2020, 16, e1008577.		0