Abraham A Palmer

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264 11,343 6.4 5.93 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
218	The Collaborative Cross, a community resource for the genetic analysis of complex traits. <i>Nature Genetics</i> , 2004 , 36, 1133-7	36.3	822
217	The nature and identification of quantitative trait loci: a community's view. <i>Nature Reviews Genetics</i> , 2003 , 4, 911-6	30.1	330
216	High-resolution genetic mapping using the Mouse Diversity outbred population. <i>Genetics</i> , 2012 , 190, 437-47	4	306
215	GWAS of lifetime cannabis use reveals new risk loci, genetic overlap with psychiatric traits, and a causal influence of schizophrenia. <i>Nature Neuroscience</i> , 2018 , 21, 1161-1170	25.5	270
214	Genome-wide association analyses of risk tolerance and risky behaviors in over 1 million individuals identify hundreds of loci and shared genetic influences. <i>Nature Genetics</i> , 2019 , 51, 245-257	36.3	259
213	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. <i>Nature Neuroscience</i> , 2018 , 21, 1656-1669	25.5	257
212	A genomewide screen of 345 families for autism-susceptibility loci. <i>American Journal of Human Genetics</i> , 2003 , 73, 886-97	11	224
211	Meta-analysis of Genome-wide Association Studies for Neuroticism, and the Polygenic Association With Major Depressive Disorder. <i>JAMA Psychiatry</i> , 2015 , 72, 642-50	14.5	222
21 0	The latent structure of impulsivity: impulsive choice, impulsive action, and impulsive personality traits. <i>Psychopharmacology</i> , 2016 , 233, 3361-70	4.7	214
209	Genetics of caffeine consumption and responses to caffeine. <i>Psychopharmacology</i> , 2010 , 211, 245-57	4.7	169
208	Genome-Wide Association Study Meta-Analysis of the Alcohol Use Disorders Identification Test (AUDIT) in Two Population-Based Cohorts. <i>American Journal of Psychiatry</i> , 2019 , 176, 107-118	11.9	156
207	Genetic Background Limits Generalizability of Genotype-Phenotype Relationships. <i>Neuron</i> , 2016 , 91, 1253-1259	13.9	143
206	Behavioral differences among C57BL/6 substrains: implications for transgenic and knockout studies. <i>Journal of Neurogenetics</i> , 2008 , 22, 315-31	1.6	142
205	Genetic analysis in the Collaborative Cross breeding population. <i>Genome Research</i> , 2011 , 21, 1223-38	9.7	132
204	Latent TGF-beta-binding protein 4 modifies muscular dystrophy in mice. <i>Journal of Clinical Investigation</i> , 2009 , 119, 3703-12	15.9	123
203	Meta-analysis of Genome-Wide Association Studies for Extraversion: Findings from the Genetics of Personality Consortium. <i>Behavior Genetics</i> , 2016 , 46, 170-82	3.2	122
202	Quantitative trait locus mapping methods for diversity outbred mice. <i>G3: Genes, Genomes, Genetics</i> , 2014 , 4, 1623-33	3.2	116

201	Glyoxalase 1 increases anxiety by reducing GABAA receptor agonist methylglyoxal. <i>Journal of Clinical Investigation</i> , 2012 , 122, 2306-15	15.9	95
200	Genetic variation and population substructure in outbred CD-1 mice: implications for genome-wide association studies. <i>PLoS ONE</i> , 2009 , 4, e4729	3.7	94
199	Voluntary ethanol drinking in C57BL/6J and DBA/2J mice before and after sensitization to the locomotor stimulant effects of ethanol. <i>Psychopharmacology</i> , 2001 , 155, 91-9	4.7	92
198	Evaluation of genetic variability in the dopamine receptor D2 in relation to behavioral inhibition and impulsivity/sensation seeking: an exploratory study with d-amphetamine in healthy participants. <i>Experimental and Clinical Psychopharmacology</i> , 2009 , 17, 374-83	3.2	91
197	Annexin A6 modifies muscular dystrophy by mediating sarcolemmal repair. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 6004-9	11.5	90
196	Pavlovian fear memory circuits and phenotype models of PTSD. <i>Neuropharmacology</i> , 2012 , 62, 638-46	5.5	88
195	The genetics of human personality. <i>Genes, Brain and Behavior</i> , 2018 , 17, e12439	3.6	88
194	A common and unstable copy number variant is associated with differences in Glo1 expression and anxiety-like behavior. <i>PLoS ONE</i> , 2009 , 4, e4649	3.7	87
193	Variation in the form of Pavlovian conditioned approach behavior among outbred male Sprague-Dawley rats from different vendors and colonies: sign-tracking vs. goal-tracking. <i>PLoS ONE</i> , 2013 , 8, e75042	3.7	87
192	Genome-wide association studies and the problem of relatedness among advanced intercross lines and other highly recombinant populations. <i>Genetics</i> , 2010 , 185, 1033-44	4	86
191	Gene expression differences in mice divergently selected for methamphetamine sensitivity. <i>Mammalian Genome</i> , 2005 , 16, 291-305	3.2	78
190	Genome-Wide Association Studies of Impulsive Personality Traits (BIS-11 and UPPS-P) and Drug Experimentation in up to 22,861 Adult Research Participants Identify Loci in the and genes. <i>Journal of Neuroscience</i> , 2019 , 39, 2562-2572	6.6	77
189	Selection for contextual fear conditioning affects anxiety-like behaviors and gene expression. <i>Genes, Brain and Behavior</i> , 2007 , 6, 736-49	3.6	75
188	Prenatal protein deprivation in rats induces changes in prepulse inhibition and NMDA receptor binding. <i>Brain Research</i> , 2004 , 996, 193-201	3.7	74
187	Genome-wide association study of behavioral, physiological and gene expression traits in outbred CFW mice. <i>Nature Genetics</i> , 2016 , 48, 919-26	36.3	72
186	Syntax for calculation of discounting indices from the monetary choice questionnaire and probability discounting questionnaire. <i>Journal of the Experimental Analysis of Behavior</i> , 2016 , 106, 156-6	3 ^{.1}	70
185	Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits. <i>Nature Neuroscience</i> , 2020 , 23, 809-818	25.5	69
184	Impulsivity as a mechanism linking child abuse and neglect with substance use in adolescence and adulthood. <i>Development and Psychopathology</i> , 2018 , 30, 417-435	4.3	66

183	Genome-wide association study of d-amphetamine response in healthy volunteers identifies putative associations, including cadherin 13 (CDH13). <i>PLoS ONE</i> , 2012 , 7, e42646	3.7	65
182	Hardy-Weinberg disequilibrium identified genotyping error of the serotonin transporter (SLC6A4) promoter polymorphism. <i>Psychiatric Genetics</i> , 2006 , 16, 31-4	2.9	64
181	Prepulse startle deficit in the Brown Norway rat: A potential genetic model <i>Behavioral Neuroscience</i> , 2000 , 114, 374-388	2.1	64
180	Role of Glyoxalase 1 (Glo1) and methylglyoxal (MG) in behavior: recent advances and mechanistic insights. <i>Frontiers in Genetics</i> , 2012 , 3, 250	4.5	63
179	QTLRel: an R package for genome-wide association studies in which relatedness is a concern. <i>BMC Genetics</i> , 2011 , 12, 66	2.6	58
178	Candidate gene studies of a promising intermediate phenotype: failure to replicate. Neuropsychopharmacology, 2013, 38, 802-16	8.7	57
177	Genome-wide association study of delay discounting in 23,217 adult research participants of European ancestry. <i>Nature Neuroscience</i> , 2018 , 21, 16-18	25.5	56
176	Psychopharmacology of theobromine in healthy volunteers. <i>Psychopharmacology</i> , 2013 , 228, 109-18	4.7	54
175	Genome-Wide Association Study of Loneliness Demonstrates a Role for Common Variation. <i>Neuropsychopharmacology</i> , 2017 , 42, 811-821	8.7	53
174	Rats are the smart choice: Rationale for a renewed focus on rats in behavioral genetics. <i>Neuropharmacology</i> , 2014 , 76 Pt B, 250-8	5.5	52
173	Association between the casein kinase 1 epsilon gene region and subjective response to D-amphetamine. <i>Neuropsychopharmacology</i> , 2006 , 31, 1056-63	8.7	51
172	Csnk1e is a genetic regulator of sensitivity to psychostimulants and opioids. <i>Neuropsychopharmacology</i> , 2012 , 37, 1026-35	8.7	49
171	Genome-wide association study of alcohol use disorder identification test (AUDIT) scores in 20B28 research participants of European ancestry. <i>Addiction Biology</i> , 2019 , 24, 121-131	4.6	49
170	Genetic variation for life history sensitivity to seasonal warming in Arabidopsis thaliana. <i>Genetics</i> , 2014 , 196, 569-77	4	47
169	Prenatal protein deprivation alters dopamine-mediated behaviors and dopaminergic and glutamatergic receptor binding. <i>Brain Research</i> , 2008 , 1237, 62-74	3.7	47
168	Bioinformatic analysis of autism positional candidate genes using biological databases and computational gene network prediction. <i>Genes, Brain and Behavior</i> , 2003 , 2, 303-20	3.6	47
167	Catechol-O-methyltransferase val158met genotype modulates sustained attention in both the drug-free state and in response to amphetamine. <i>Psychiatric Genetics</i> , 2010 , 20, 85-92	2.9	47
166	Mapping of Craniofacial Traits in Outbred Mice Identifies Major Developmental Genes Involved in Shape Determination. <i>PLoS Genetics</i> , 2015 , 11, e1005607	6	45

165	Social neuroscience and its potential contribution to psychiatry. World Psychiatry, 2014, 13, 131-9	14.4	45	
164	Effects of a Drd2 deletion mutation on ethanol-induced locomotor stimulation and sensitization suggest a role for epistasis. <i>Behavior Genetics</i> , 2003 , 33, 311-24	3.2	45	
163	A large-scale genome-wide association study meta-analysis of cannabis use disorder. <i>Lancet Psychiatry,the</i> , 2020 , 7, 1032-1045	23.3	43	
162	OPRM1 gene variants modulate amphetamine-induced euphoria in humans. <i>Genes, Brain and Behavior</i> , 2011 , 10, 199-209	3.6	41	
161	Interrelationships among parental family history of substance misuse, delay discounting, and personal substance use. <i>Psychopharmacology</i> , 2016 , 233, 39-48	4.7	39	
160	Genome-wide association for fear conditioning in an advanced intercross mouse line. <i>Behavior Genetics</i> , 2012 , 42, 437-48	3.2	38	
159	A role for casein kinase 1 epsilon in the locomotor stimulant response to methamphetamine. <i>Psychopharmacology</i> , 2009 , 203, 703-11	4.7	38	
158	Differences in aggressive behavior and DNA copy number variants between BALB/cJ and BALB/cByJ substrains. <i>Behavior Genetics</i> , 2010 , 40, 201-10	3.2	38	
157	Dark matter: are mice the solution to missing heritability?. Frontiers in Genetics, 2011, 2, 32	4.5	36	
156	Hnrnph1 Is A Quantitative Trait Gene for Methamphetamine Sensitivity. <i>PLoS Genetics</i> , 2015 , 11, e100	5763	36	
155	Negative emotionality: monoamine oxidase B gene variants modulate personality traits in healthy humans. <i>Journal of Neural Transmission</i> , 2009 , 116, 1323-34	4.3	35	
154	Fine mapping of QTL for prepulse inhibition in LG/J and SM/J mice using F(2) and advanced intercross lines. <i>Genes, Brain and Behavior</i> , 2010 , 9, 759-67	3.6	34	
153	Corticotropin-releasing factor overexpression decreases ethanol drinking and increases sensitivity to the sedative effects of ethanol. <i>Psychopharmacology</i> , 2004 , 176, 386-97	4.7	34	
152	Strain differences in Fos expression following airpuff startle in Spontaneously Hypertensive and Wistar Kyoto rats. <i>Neuroscience</i> , 1999 , 89, 965-78	3.9	34	
151	A simulation study of permutation, bootstrap, and gene dropping for assessing statistical significance in the case of unequal relatedness. <i>Genetics</i> , 2013 , 193, 1015-8	4	33	
150	Identification of quantitative trait loci for prepulse inhibition in rats. <i>Psychopharmacology</i> , 2003 , 165, 270-9	4.7	33	
149	Genome-wide association for methamphetamine sensitivity in an advanced intercross mouse line. <i>Genes, Brain and Behavior</i> , 2012 , 11, 52-61	3.6	32	
148	Practical considerations regarding the use of genotype and pedigree data to model relatedness in the context of genome-wide association studies. <i>G3: Genes, Genomes, Genetics</i> , 2013 , 3, 1861-7	3.2	31	

147	Casein kinase 1 enables nucleus accumbens amphetamine-induced locomotion by regulating AMPA receptor phosphorylation. <i>Journal of Neurochemistry</i> , 2011 , 118, 237-47	6	31
146	Acute and chronic responses to the convulsant pilocarpine in DBA/2J and A/J mice. <i>Neuroscience</i> , 2007 , 149, 465-75	3.9	31
145	Reverse selection for differential response to the locomotor stimulant effects of ethanol provides evidence for pleiotropic genetic influence on locomotor response to other drugs of abuse. <i>Alcoholism: Clinical and Experimental Research</i> , 2003 , 27, 1535-47	3.7	31
144	Prepulse startle deficit in the Brown Norway rat: a potential genetic model. <i>Behavioral Neuroscience</i> , 2000 , 114, 374-88	2.1	30
143	Phenome-wide investigation of health outcomes associated with genetic predisposition to loneliness. <i>Human Molecular Genetics</i> , 2019 , 28, 3853-3865	5.6	29
142	High-resolution genetic mapping of complex traits from a combined analysis of F2 and advanced intercross mice. <i>Genetics</i> , 2014 , 198, 103-16	4	29
141	Strong genetic influences on measures of behavioral-regulation among inbred rat strains. <i>Genes, Brain and Behavior</i> , 2013 , 12, 490-502	3.6	29
140	Fine-mapping of muscle weight QTL in LG/J and SM/J intercrosses. <i>Physiological Genomics</i> , 2010 , 42A, 33-8	3.6	29
139	Initial sensitivity, tolerance and cross-tolerance to allopregnanolone- and ethanol-induced hypothermia in selected mouse lines. <i>Psychopharmacology</i> , 2002 , 162, 313-22	4.7	28
138	Recent Efforts to Dissect the Genetic Basis of Alcohol Use and Abuse. <i>Biological Psychiatry</i> , 2020 , 87, 609-618	7.9	28
137	Propensity for social interaction predicts nicotine-reinforced behaviors in outbred rats. <i>Genes, Brain and Behavior</i> , 2014 , 13, 202-12	3.6	27
136	Glyoxalase 1 and its substrate methylglyoxal are novel regulators of seizure susceptibility. <i>Epilepsia</i> , 2013 , 54, 649-57	6.4	27
135	A major QTL on chromosome 11 influences psychostimulant and opioid sensitivity in mice. <i>Genes, Brain and Behavior</i> , 2009 , 8, 795-805	3.6	27
134	Genetic factors modulating the response to stimulant drugs in humans. <i>Current Topics in Behavioral Neurosciences</i> , 2012 , 12, 537-77	3.4	26
133	Distinct genetic regions modify specific muscle groups in muscular dystrophy. <i>Physiological Genomics</i> , 2011 , 43, 24-31	3.6	25
132	Sensitivity to the locomotor stimulant effects of ethanol and allopregnanolone is influenced by common genes <i>Behavioral Neuroscience</i> , 2002 , 116, 126-137	2.1	25
131	More aroused, less fatigued: fatty acid amide hydrolase gene polymorphisms influence acute response to amphetamine. <i>Neuropsychopharmacology</i> , 2010 , 35, 613-22	8.7	24
130	Identification of a novel, fast-acting GABAergic antidepressant. <i>Molecular Psychiatry</i> , 2018 , 23, 384-391	15.1	23

129	Genetic determinants for intramuscular fat content and water-holding capacity in mice selected for high muscle mass. <i>Mammalian Genome</i> , 2011 , 22, 530-43	3.2	23	
128	Fine-mapping alleles for body weight in LG/J ISM/J Fland F(34) advanced intercross lines. <i>Mammalian Genome</i> , 2011 , 22, 563-71	3.2	23	
127	Genetic analysis of impulsive personality traits: Examination of a priori candidates and genome-wide variation. <i>Psychiatry Research</i> , 2018 , 259, 398-404	9.9	23	
126	Assessment of behaviors modeling aspects of schizophrenia in Csmd1 mutant mice. <i>PLoS ONE</i> , 2012 , 7, e51235	3.7	22	
125	Use of chromosome substitution strains to identify seizure susceptibility loci in mice. <i>Mammalian Genome</i> , 2007 , 18, 23-31	3.2	22	
124	Rapid selection response for contextual fear conditioning in a cross between C57BL/6J and A/J: behavioral, QTL and gene expression analysis. <i>Behavior Genetics</i> , 2008 , 38, 277-91	3.2	22	
123	Significance thresholds for quantitative trait locus mapping under selective genotyping. <i>Genetics</i> , 2007 , 177, 1963-6	4	22	
122	Behavioral sensitization to ethanol is modulated by environmental conditions, but is not associated with cross-sensitization to allopregnanolone or pentobarbital in DBA/2J mice. <i>Neuroscience</i> , 2005 , 131, 263-73	3.9	22	
121	Locomotor activity responses to ethanol, other alcohols and GABA-A acting compounds in forward-and reverse-selected FAST and SLOW mouse lines <i>Behavioral Neuroscience</i> , 2002 , 116, 958-967	2.1	22	
120	Using Heterogeneous Stocks for Fine-Mapping Genetically Complex Traits. <i>Methods in Molecular Biology</i> , 2019 , 2018, 233-247	1.4	22	
119	Further evidence of association between amphetamine response and SLC6A2 gene variants. <i>Psychopharmacology</i> , 2009 , 206, 501-11	4.7	21	
118	Metal-Binding Pharmacophore Library Yields the Discovery of a Glyoxalase 1 Inhibitor. <i>Journal of Medicinal Chemistry</i> , 2019 , 62, 1609-1625	8.3	20	
117	Fine-mapping QTLs in advanced intercross lines and other outbred populations. <i>Mammalian Genome</i> , 2014 , 25, 271-92	3.2	20	
116	Congenic dissection of a major QTL for methamphetamine sensitivity implicates epistasis. <i>Genes, Brain and Behavior</i> , 2012 , 11, 623-32	3.6	20	
115	Genetic architecture of fear conditioning in chromosome substitution strains: relationship to measures of innate (unlearned) anxiety-like behavior. <i>Mammalian Genome</i> , 2007 , 18, 221-8	3.2	20	
114	Genome wide association analysis in a mouse advanced intercross line. <i>Nature Communications</i> , 2018 , 9, 5162	17.4	20	
113	Incentive salience attribution, "sensation-seeking" and "novelty-seeking" are independent traits in a large sample of male and female heterogeneous stock rats. <i>Scientific Reports</i> , 2019 , 9, 2351	4.9	19	
112	Heterogeneous stock rats: a model to study the genetics of despair-like behavior in adolescence. <i>Genes, Brain and Behavior</i> , 2018 , 17, 139-148	3.6	19	

111	Anxiety and fear in a cross of C57BL/6J and DBA/2J mice: mapping overlapping and independent QTL for related traits. <i>Genes, Brain and Behavior</i> , 2011 , 10, 604-14	3.6	19
110	Polymorphisms in dopamine transporter (SLC6A3) are associated with stimulant effects of D-amphetamine: an exploratory pharmacogenetic study using healthy volunteers. <i>Behavior Genetics</i> , 2010 , 40, 255-61	3.2	19
109	Forward, Relaxed, and Reverse Selection for Reduced and Enhanced Sensitivity to Ethanol's Locomotor Stimulant Effects in Mice. <i>Alcoholism: Clinical and Experimental Research</i> , 2002 , 26, 593-602	3.7	19
108	QTL Analysis of Type I and Type IIA Fibers in Soleus Muscle in a Cross between LG/J and SM/J Mouse Strains. <i>Frontiers in Genetics</i> , 2011 , 2, 99	4.5	18
107	Latent inhibition and conditioning in rat strains which show differential prepulse inhibition. <i>Behavior Genetics</i> , 2001 , 31, 325-33	3.2	18
106	Genome-wide Associations Reveal Human-Mouse Genetic Convergence and Modifiers of Myogenesis, CPNE1 and STC2. <i>American Journal of Human Genetics</i> , 2019 , 105, 1222-1236	11	18
105	Premature responding is associated with approach to a food cue in male and female heterogeneous stock rats. <i>Psychopharmacology</i> , 2016 , 233, 2593-605	4.7	17
104	Neuronal overexpression of Glo1 or amygdalar microinjection of methylglyoxal is sufficient to regulate anxiety-like behavior in mice. <i>Behavioural Brain Research</i> , 2016 , 301, 119-23	3.4	17
103	Does COMT genotype influence the effects of d-amphetamine on executive functioning?. <i>Genes, Brain and Behavior</i> , 2013 , 12, 13-20	3.6	17
102	Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. <i>Genes, Brain and Behavior</i> , 2006 , 5, 506-17	3.6	17
101	Content and Performance of the MiniMUGA Genotyping Array: A New Tool To Improve Rigor and Reproducibility in Mouse Research. <i>Genetics</i> , 2020 , 216, 905-930	4	17
100	Forward, relaxed, and reverse selection for reduced and enhanced sensitivity to ethanol's locomotor stimulant effects in mice. <i>Alcoholism: Clinical and Experimental Research</i> , 2002 , 26, 593-602	3.7	17
99	Photochemical control of the infectivity of adenoviral vectors using a novel photocleavable biotinylation reagent. <i>Chemistry and Biology</i> , 2002 , 9, 567-73		16
98	Nociceptin attenuates the escalation of oxycodone self-administration by normalizing CeA-GABA transmission in highly addicted rats. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 2140-2148	11.5	15
97	Genetic variation associated with euphorigenic effects of d-amphetamine is associated with diminished risk for schizophrenia and attention deficit hyperactivity disorder. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 5968-73	11.5	15
96	Glo1 inhibitors for neuropsychiatric and anti-epileptic drug development. <i>Biochemical Society Transactions</i> , 2014 , 42, 461-7	5.1	15
95	Methamphetamine-induced conditioned place preference in LG/J and SM/J mouse strains and an F45/F46 advanced intercross line. <i>Frontiers in Genetics</i> , 2012 , 3, 126	4.5	15
94	Traits of fear resistance and susceptibility in an advanced intercross line. <i>European Journal of Neuroscience</i> , 2013 , 38, 3314-24	3.5	15

93	Modulation of Tcf7l2 expression alters behavior in mice. PLoS ONE, 2011, 6, e26897	3.7	15
92	Multivariate GWAS of psychiatric disorders and their cardinal symptoms reveal two dimensions of cross-cutting genetic liabilities		15
91	Locomotor activity responses to ethanol, other alcohols, and GABA-A acting compounds in forward-and reverse-selected FAST and SLOW mouse lines. <i>Behavioral Neuroscience</i> , 2002 , 116, 958-67	2.1	15
90	Genetic influences on ADHD symptom dimensions: Examination of a priori candidates, gene-based tests, genome-wide variation, and SNP heritability. <i>American Journal of Medical Genetics Part B:</i> Neuropsychiatric Genetics, 2017 , 174, 458-466	3.5	14
89	Emerging phenotyping strategies will advance our understanding of psychiatric genetics. <i>Nature Neuroscience</i> , 2020 , 23, 475-480	25.5	14
88	The circadian clock gene Csnk1e regulates rapid eye movement sleep amount, and nonrapid eye movement sleep architecture in mice. <i>Sleep</i> , 2014 , 37, 785-93, 793A-793C	1.1	14
87	Genome-Wide Association Study in 3,173 Outbred Rats Identifies Multiple Loci for Body Weight, Adiposity, and Fasting Glucose. <i>Obesity</i> , 2020 , 28, 1964-1973	8	14
86	Are attention lapses related to d-amphetamine liking?. <i>Psychopharmacology</i> , 2010 , 208, 201-9	4.7	13
85	Mapping a mouse limbic seizure susceptibility locus on chromosome 10. <i>Epilepsia</i> , 2011 , 52, 2076-83	6.4	12
84	Airpuff startle stress elicited fos expression in brain cardiovascular areas of young SHR and WKY rats. <i>Clinical and Experimental Hypertension</i> , 1999 , 21, 1061-81	2.2	11
83	Involvement of central corticotropin-releasing factor (CRF) in suckling-induced inhibition of luteinizing hormone secretion in lactating rats. <i>Journal of Neuroendocrinology</i> , 1993 , 5, 451-9	3.8	11
82	Sex-dependent associations between addiction-related behaviors and the microbiome in outbred rats. <i>EBioMedicine</i> , 2020 , 55, 102769	8.8	11
81	Social and anxiety-like behaviors contribute to nicotine self-administration in adolescent outbred rats. <i>Scientific Reports</i> , 2018 , 8, 18069	4.9	11
80	Genetic and pharmacological manipulation of glyoxalase 1 regulates voluntary ethanol consumption in mice. <i>Addiction Biology</i> , 2017 , 22, 381-389	4.6	10
79	Phenotypic instability between the near isogenic substrains BALB/cJ and BALB/cByJ. <i>Mammalian Genome</i> , 2014 , 25, 564-72	3.2	10
78	Discovery and refinement of muscle weight QTLs in B6 ID2 advanced intercross mice. <i>Physiological Genomics</i> , 2014 , 46, 571-82	3.6	10
77	A large QTL for fear and anxiety mapped using an F2 cross can be dissected into multiple smaller QTLs. <i>Genes, Brain and Behavior</i> , 2013 , 12, 714-22	3.6	10
76	Multivariate analysis of 1.5 million people identifies genetic associations with traits related to self-regulation and addiction. <i>Nature Neuroscience</i> , 2021 , 24, 1367-1376	25.5	10

75	Sensitivity to the locomotor stimulant effects of ethanol and allopregnanolone is influenced by common genes. <i>Behavioral Neuroscience</i> , 2002 , 116, 126-37	2.1	10
74	Fine-mapping of genes determining extrafusal fiber properties in murine soleus muscle. <i>Physiological Genomics</i> , 2017 , 49, 141-150	3.6	9
73	Genome-wide association study meta-analysis of the Alcohol Use Disorder Identification Test (AUDIT) in two population-based cohorts (N=141,932)		9
72	Genetic risk for major depressive disorder and loneliness in sex-specific associations with coronary artery disease. <i>Molecular Psychiatry</i> , 2021 , 26, 4254-4264	15.1	9
71	Individual differences in timing of peak positive subjective responses to d-amphetamine: Relationship to pharmacokinetics and physiology. <i>Journal of Psychopharmacology</i> , 2016 , 30, 330-43	4.6	8
70	Sex-specific linkage scans in opioid dependence. <i>American Journal of Medical Genetics Part B:</i> Neuropsychiatric Genetics, 2017 , 174, 261-268	3.5	8
69	Effect of Forward and Reverse Selection for Ethanol-Induced Locomotor Response on Other Measures of Ethanol Sensitivity. <i>Alcoholism: Clinical and Experimental Research</i> , 2002 , 26, 1322-1329	3.7	8
68	Steep Discounting of Future Rewards as an Impulsivity Phenotype: A Concise Review. <i>Current Topics in Behavioral Neurosciences</i> , 2020 , 47, 113-138	3.4	7
67	A dendritic organization of lateral amygdala neurons in fear susceptible and resistant mice. <i>Neurobiology of Learning and Memory</i> , 2016 , 127, 64-71	3.1	7
66	Mice selectively bred for High and Low fear behavior show differences in the number of pMAPK (p44/42 ERK) expressing neurons in lateral amygdala following Pavlovian fear conditioning. Neurobiology of Learning and Memory, 2014, 112, 195-203	3.1	7
65	QTLs for murine red blood cell parameters in LG/J and SM/J F(2) and advanced intercross lines. <i>Mammalian Genome</i> , 2012 , 23, 356-66	3.2	7
64	Murine Warriors or Worriers: The Saga of Comt1, B2 SINE Elements, and the Future of Translational Genetics. <i>Frontiers in Neuroscience</i> , 2010 , 4, 177	5.1	7
63	Attenuation of Fos expression to airpuff startle stimuli following tympanic membrane rupture. <i>Brain Research</i> , 1998 , 787, 91-8	3.7	7
62	Angiotensin II receptor binding sites in the ventral portion of the bed nucleus of the stria terminalis are reduced by interruption of the medial forebrain bundle. <i>Brain Research</i> , 1998 , 809, 5-11	3.7	7
61	Inappropriate choice of the experimental unit leads to a dramatic overestimation of the significance of quantitative trait loci for prepulse inhibition and startle response in recombinant congenic mice. <i>Neuropsychopharmacology</i> , 2003 , 28, 818; author reply 819	8.7	7
60	Hierarchical investigation of genetic influences on response inhibition in healthy young adults. <i>Experimental and Clinical Psychopharmacology</i> , 2017 , 25, 512-520	3.2	7
59	Trans-ancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders		7
58	The Latent Genetic Structure of Impulsivity and Its Relation to Internalizing Psychopathology. <i>Psychological Science</i> , 2020 , 31, 1025-1035	7.9	7

57	Item-Level Genome-Wide Association Study of the Alcohol Use Disorders Identification Test in Three Population-Based Cohorts. <i>American Journal of Psychiatry</i> , 2021 , applajp202020091390	11.9	7
56	Systems genetic and pharmacological analysis identifies candidate genes underlying mechanosensation in the von Frey test. <i>Genes, Brain and Behavior</i> , 2016 , 15, 604-15	3.6	7
55	Cdh13 and AdipoQ gene knockout alter instrumental and Pavlovian drug conditioning. <i>Genes, Brain and Behavior</i> , 2017 , 16, 686-698	3.6	6
54	Genomic basis of delayed reward discounting. <i>Behavioural Processes</i> , 2019 , 162, 157-161	1.6	6
53	Genetic influences on delayed reward discounting: A genome-wide prioritized subset approach. <i>Experimental and Clinical Psychopharmacology</i> , 2019 , 27, 29-37	3.2	6
52	Multivariate genomic analysis of 1.5 million people identifies genes related to addiction, antisocial behavior, and health		6
51	Meta-analysis of problematic alcohol use in 435,563 individuals identifies 29 risk variants and yields insights into biology, pleiotropy and causality		6
50	Electronic Health Records Are the Next Frontier for the Genetics of Substance Use Disorders. <i>Trends in Genetics</i> , 2019 , 35, 317-318	8.5	5
49	Adapting Genotyping-by-Sequencing and Variant Calling for Heterogeneous Stock Rats. <i>G3: Genes, Genomes, Genetics</i> , 2020 , 10, 2195-2205	3.2	5
48	Inhibition of Glyoxalase 1 reduces alcohol self-administration in dependent and nondependent rats. <i>Pharmacology Biochemistry and Behavior</i> , 2018 , 167, 36-41	3.9	5
47	Translational genetic approaches to substance use disorders: bridging the gap between mice and humans. <i>Human Genetics</i> , 2012 , 131, 931-9	6.3	5
46	Differences between SHR and WKY following the airpuff startle stimulus in the number of Fos expressing, RVLM projecting neurons. <i>Clinical and Experimental Hypertension</i> , 2002 , 24, 125-39	2.2	5
45	Genetic characterization of outbred Sprague Dawley rats and utility for genome-wide association stud	ies	5
44	Integration of genome-wide association and extant brain expression QTL identifies candidate genes influencing prepulse inhibition in inbred F1 mice. <i>Genes, Brain and Behavior</i> , 2016 , 15, 260-70	3.6	5
43	Polygenic contributions to alcohol use and alcohol use disorders across population-based and clinically ascertained samples. <i>Psychological Medicine</i> , 2021 , 51, 1147-1156	6.9	5
42	A locus on mouse Ch10 influences susceptibility to limbic seizure severity: fine mapping and in silico candidate gene analysis. <i>Genes, Brain and Behavior</i> , 2014 , 13, 341-9	3.6	4
41	Differences in cultured cardiac fibroblast populations isolated from SHR and WKY rats. <i>Clinical and Experimental Pharmacology and Physiology</i> , 1995 , 22, S265-7	3	4
40	Genome-wide association study of Alcohol Use Disorder Identification Test (AUDIT) scores in 20,328 research participants of European ancestry		4

39	Polygenic transcriptome risk scores improve portability of polygenic risk scores across ancestries		4
38	Phenome-wide Investigation of Health Outcomes Associated with Genetic Predisposition to Loneliness		4
37	Sensitivity to food and cocaine cues are independent traits in a large sample of heterogeneous stock rats. <i>Scientific Reports</i> , 2021 , 11, 2223	4.9	4
36	Glyoxalase 1 (GLO1) Inhibition or Genetic Overexpression Does Not Alter Ethanol's Locomotor Effects: Implications for GLO1 as a Therapeutic Target in Alcohol Use Disorders. <i>Alcoholism: Clinical and Experimental Research</i> , 2018 , 42, 869-878	3.7	3
35	Polygenic transcriptome risk scores (PTRS) can improve portability of polygenic risk scores across ancestries <i>Genome Biology</i> , 2022 , 23, 23	18.3	3
34	Genome wide association analysis in a mouse advanced intercross line		3
33	Genome-wide association study in two cohorts from a multi-generational mouse advanced intercross line highlights the difficulty of replication		3
32	Dissecting indirect genetic effects from peers in laboratory mice		3
31	Genome wide association study in 3,173 outbred rats identifies multiple loci for body weight, adiposity, and fasting glucose		3
30	Genome-Wide Association Study in Two Cohorts from a Multi-generational Mouse Advanced Intercross Line Highlights the Difficulty of Replication Due to Study-Specific Heterogeneity. <i>G3: Genes, Genomes, Genetics</i> , 2020 , 10, 951-965	3.2	3
29	The Cocaine and Oxycodone Biobanks, Two Repositories from Genetically Diverse and Behaviorally Characterized Rats for the Study of Addiction. <i>ENeuro</i> , 2021 , 8,	3.9	3
28	Effect of forward and reverse selection for ethanol-induced locomotor response on other measures of ethanol sensitivity. <i>Alcoholism: Clinical and Experimental Research</i> , 2002 , 26, 1322-9	3.7	3
27	More on ADORA. <i>Psychopharmacology</i> , 2010 , 212, 699-700	4.7	2
26	Evaluation of delay discounting as a transdiagnostic research domain criteria indicator in 1388 general community adults <i>Psychological Medicine</i> , 2022 , 1-9	6.9	2
25	Modeling epistasis in mice and yeast using the proportion of two or more distinct genetic backgrounds: Evidence for "polygenic epistasis". <i>PLoS Genetics</i> , 2020 , 16, e1009165	6	2
24	Adapting genotyping-by-sequencing and variant calling for heterogeneous stock rats		2
23	Content and performance of the MiniMUGA genotyping array, a new tool to improve rigor and reproducibility in mouse research		2
22	Multidimensional latent structure of risk-related phenotypes in healthy young adults. <i>Experimental and Clinical Psychopharmacology</i> , 2020 , 28, 55-64	3.2	2

21	Forward, Relaxed, and Reverse Selection for Reduced and Enhanced Sensitivity to Ethanol???s Locomotor Stimulant Effects in Mice. <i>Alcoholism: Clinical and Experimental Research</i> , 2002 , 26, 593-602	3.7	2
20	Genome-wide associations reveal human-mouse genetic convergence and modifiers of myogenesis,CP	NE1an	dSTC2
19	Modeling epistasis in mice and yeast using the proportion of two or more distinct genetic backgrounds: evidence for polygenic epistasis□		2
18	Behavioral Genetic Studies in Rats. <i>Methods in Molecular Biology</i> , 2019 , 2018, 319-326	1.4	1
17	Polymorphic SNPs, short tandem repeats and structural variants are responsible for differential gene expression across C57BL/6 and C57BL/10 substrains		1
16	Multivariate GWAS elucidates the genetic architecture of alcohol consumption and misuse, corrects biases, and reveals novel associations with disease		1
15	Integration of evidence across human and model organism studies: A meeting report. <i>Genes, Brain and Behavior</i> , 2021 , 20, e12738	3.6	1
14	Mapping Pathways by Which Genetic Risk Influences Adolescent Externalizing Behavior: The Interplay Between Externalizing Polygenic Risk Scores, Parental Knowledge, and Peer Substance Use. <i>Behavior Genetics</i> , 2021 , 51, 543-558	3.2	1
13	Characterization of cocaine addiction-like behavior in heterogeneous stock rats		1
12	Functional validation of a finding from a mouse genome-wide association study shows that Azi2 influences the acute locomotor simulant response to methamphetamine. <i>Genes, Brain and Behavior</i> , 2021 , 20, e12760	3.6	1
11	Genetic and Pharmacological Manipulations of Glyoxalase 1 Mediate Ethanol Withdrawal Seizure Susceptibility in Mice. <i>Brain Sciences</i> , 2021 , 11,	3.4	1
10	Analysis of independent cohorts of outbred CFW mice reveals novel loci for behavioral and physiological traits and identifies factors determining reproducibility		1
9	A natural mutator allele shapes mutation spectrum variation in mice <i>Nature</i> , 2022 , 605, 497-502	50.4	1
8	Assessing the motivational effects of ethanol in mice using a discrete-trial current-intensity intracranial self-stimulation procedure. <i>Drug and Alcohol Dependence</i> , 2020 , 207, 107806	4.9	O
7	Dissecting indirect genetic effects from peers in laboratory mice. <i>Genome Biology</i> , 2021 , 22, 216	18.3	O
6	Genome-Wide Association Study on Three Behaviors Tested in an Open Field in Heterogeneous Stock Rats Identifies Multiple Loci Implicated in Psychiatric Disorders <i>Frontiers in Psychiatry</i> , 2022 , 13, 790566	5	O
5	SNPs, short tandem repeats, and structural variants are responsible for differential gene expression across C57BL/6 and C57BL/10 substrains. <i>Cell Genomics</i> , 2022 , 2, 100102		O
4	Leptin Protects Against the Development and Expression of Cocaine Addiction-Like Behavior in Heterogeneous Stock Rats <i>Frontiers in Behavioral Neuroscience</i> , 2022 , 16, 832899	3.5	O

3	Rufy1 or Hnrnph1 is a likely quantitative trait gene for methamphetamine sensitivity. <i>FASEB Journal</i> , 2013 , 27, lb472	0.9
2	ACNP efforts toward reducing climate change. <i>Neuropsychopharmacology</i> , 2020 , 45, 2137-2138	8.7
1	A mutant allele of glycoprotein M6-B (Gpm6b) facilitates behavioral flexibility but increases delay discounting <i>Genes, Brain and Behavior</i> , 2022 , e12800	3.6