

# Kevin G Becker

## List of Publications by Year in descending order

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180  
papers

20,919  
citations

17405

63  
h-index

10424

139  
g-index

182  
all docs

182  
docs citations

182  
times ranked

29968  
citing authors

#	ARTICLE	IF	CITATIONS
1	Resveratrol improves health and survival of mice on a high-calorie diet. <i>Nature</i> , 2006, 444, 337-342.	13.7	3,882
2	Metformin improves healthspan and lifespan in mice. <i>Nature Communications</i> , 2013, 4, 2192.	5.8	1,118
3	Resveratrol Delays Age-Related Deterioration and Mimics Transcriptional Aspects of Dietary Restriction without Extending Life Span. <i>Cell Metabolism</i> , 2008, 8, 157-168.	7.2	1,060
4	The Genetic Association Database. <i>Nature Genetics</i> , 2004, 36, 431-432.	9.4	888
5	LincRNA-p21 Suppresses Target mRNA Translation. <i>Molecular Cell</i> , 2012, 47, 648-655.	4.5	876
6	Analysis of Microarray Data Using Z Score Transformation. <i>Journal of Molecular Diagnostics</i> , 2003, 5, 73-81.	1.2	860
7	miR-24 Inhibits Cell Proliferation by Targeting E2F2, MYC, and Other Cell-Cycle Genes via Binding to 3'UTR MicroRNA Recognition Elements. <i>Molecular Cell</i> , 2009, 35, 610-625.	4.5	544
8	Effects of Sex, Strain, and Energy Intake on Hallmarks of Aging in Mice. <i>Cell Metabolism</i> , 2016, 23, 1093-1112.	7.2	360
9	AGEMAP: A Gene Expression Database for Aging in Mice. <i>PLoS Genetics</i> , 2007, 3, e201.	1.5	355
10	The SIRT1 Activator SRT1720 Extends Lifespan and Improves Health of Mice Fed a Standard Diet. <i>Cell Reports</i> , 2014, 6, 836-843.	2.9	342
11	Transcriptional Profiling of Aging in Human Muscle Reveals a Common Aging Signature. <i>PLoS Genetics</i> , 2006, 2, e115.	1.5	331
12	MicroRNA Expression and Identification of Putative miRNA Targets in Ovarian Cancer. <i>PLoS ONE</i> , 2008, 3, e2436.	1.1	303
13	Microarray analysis of gene expression in the prefrontal cortex in schizophrenia: a preliminary study. <i>Schizophrenia Research</i> , 2002, 58, 11-20.	1.1	261
14	Genome-wide Analysis of Histone Methylation Reveals Chromatin State-Based Regulation of Gene Transcription and Function of Memory CD8+ T Cells. <i>Immunity</i> , 2009, 30, 912-925.	6.6	256
15	SRT1720 improves survival and healthspan of obese mice. <i>Scientific Reports</i> , 2011, 1, 70.	1.6	249
16	Analysis of gene expression in multiple sclerosis lesions using cDNA microarrays. <i>Annals of Neurology</i> , 1999, 46, 425-428.	2.8	237
17	Application of cDNA microarrays to examine gene expression differences in schizophrenia. <i>Brain Research Bulletin</i> , 2001, 55, 641-650.	1.4	236
18	Global analysis of stress-regulated mRNA turnover by using cDNA arrays. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 10611-10616.	3.3	212

#	ARTICLE	IF	CITATIONS
19	Calorie restriction in humans inhibits the PI3K/AKT pathway and induces a younger transcription profile. <i>Aging Cell</i> , 2013, 12, 645-651.	3.0	208
20	SIRT1 extends survival of male mice on a standard diet and preserves bone and muscle mass. <i>Aging Cell</i> , 2014, 13, 787-796.	3.0	208
21	Transcriptional profiling of aging in human muscle reveals a common aging signature. <i>PLoS Genetics</i> , 2005, preprint, e115.	1.5	208
22	Senescence-associated lncRNAs: senescence-associated long noncoding RNAs. <i>Aging Cell</i> , 2013, 12, 890-900.	3.0	184
23	PubMatrix: a tool for multiplex literature mining. <i>BMC Bioinformatics</i> , 2003, 4, 61.	1.2	180
24	Altered Extracellular Vesicle Concentration, Cargo, and Function in Diabetes. <i>Diabetes</i> , 2018, 67, 2377-2388.	0.3	176
25	Sex-Dependent Metabolic, Neuroendocrine, and Cognitive Responses to Dietary Energy Restriction and Excess. <i>Endocrinology</i> , 2007, 148, 4318-4333.	1.4	167
26	Control of gene expression during T cell activation: alternate regulation of mRNA transcription and mRNA stability. <i>BMC Genomics</i> , 2005, 6, 75.	1.2	163
27	PAR-CLIP analysis uncovers AUF1 impact on target RNA fate and genome integrity. <i>Nature Communications</i> , 2014, 5, 5248.	5.8	156
28	Metformin-mediated increase in DICER1 regulates microRNA expression and cellular senescence. <i>Aging Cell</i> , 2016, 15, 572-581.	3.0	153
29	The common variants/multiple disease hypothesis of common complex genetic disorders. <i>Medical Hypotheses</i> , 2004, 62, 309-317.	0.8	144
30	Hippocampal gene expression patterns underlying the enhancement of memory by running in aged mice. <i>Neurobiology of Aging</i> , 2010, 31, 1937-1949.	1.5	135
31	Mice Fed Rapamycin Have an Increase in Lifespan Associated with Major Changes in the Liver Transcriptome. <i>PLoS ONE</i> , 2014, 9, e83988.	1.1	132
32	Commensal bacteria contribute to insulin resistance in aging by activating innate B1a cells. <i>Science Translational Medicine</i> , 2018, 10, .	5.8	121
33	FACS purification of immunolabeled cell types from adult rat brain. <i>Journal of Neuroscience Methods</i> , 2012, 203, 10-18.	1.3	119
34	Tomatidine enhances lifespan and healthspan in <i>C. elegans</i> through mitophagy induction via the SKN-1/Nrf2 pathway. <i>Scientific Reports</i> , 2017, 7, 46208.	1.6	116
35	CREB phosphorylation regulates striatal transcriptional responses in the self-administration model of methamphetamine addiction in the rat. <i>Neurobiology of Disease</i> , 2013, 58, 132-143.	2.1	115
36	Methamphetamine Causes Differential Alterations in Gene Expression and Patterns of Histone Acetylation/Hypoacetylation in the Rat Nucleus Accumbens. <i>PLoS ONE</i> , 2012, 7, e34236.	1.1	111

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37	iTRAQ Analysis of Complex Proteome Alterations in 3xTgAD Alzheimer's Mice: Understanding the Interface between Physiology and Disease. <i>PLoS ONE</i> , 2008, 3, e2750.	1.1	110
38	DNA polymerase $\beta$ deficiency leads to neurodegeneration and exacerbates Alzheimer disease phenotypes. <i>Nucleic Acids Research</i> , 2015, 43, 943-959.	6.5	110
39	Systematic analysis, comparison, and integration of disease based human genetic association data and mouse genetic phenotypic information. <i>BMC Medical Genomics</i> , 2010, 3, 1.	0.7	106
40	Gene expression atlas of the mouse central nervous system: impact and interactions of age, energy intake and gender. <i>Genome Biology</i> , 2007, 8, R234.	13.9	103
41	AMPK agonist AICAR improves cognition and motor coordination in young and aged mice. <i>Learning and Memory</i> , 2014, 21, 119-126.	0.5	102
42	Skeletal muscle <i>ex vivo</i> mitochondrial respiration parallels decline in <i>in vivo</i> oxidative capacity, cardiorespiratory fitness, and muscle strength: The Baltimore Longitudinal Study of Aging. <i>Aging Cell</i> , 2018, 17, e12725.	3.0	101
43	Region-specific transcriptional response to chronic nicotine in rat brain. <i>Brain Research</i> , 2001, 909, 194-203.	1.1	96
44	Mannose receptor modulates macrophage polarization and allergic inflammation through miR-511-3p. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 350-364.e8.	1.5	91
45	Amitriptyline-Mediated Cognitive Enhancement in Aged 3xTg Alzheimer's Disease Mice Is Associated with Neurogenesis and Neurotrophic Activity. <i>PLoS ONE</i> , 2011, 6, e21660.	1.1	82
46	Autism, asthma, inflammation, and the hygiene hypothesis. <i>Medical Hypotheses</i> , 2007, 69, 731-740.	0.8	81
47	Development of a highly specialized cDNA array for the study and diagnosis of epithelial ovarian cancer. <i>Cancer Research</i> , 2002, 62, 2923-8.	0.4	81
48	Gene expression characteristics of CD28null memory phenotype CD8+ T cells and its implication in T-cell aging. <i>Immunological Reviews</i> , 2005, 205, 190-206.	2.8	80
49	A Novel Combination of Factors, Termed SPIE, which Promotes Dopaminergic Neuron Differentiation from Human Embryonic Stem Cells. <i>PLoS ONE</i> , 2009, 4, e6606.	1.1	79
50	Gene Expression Profile of Herpesvirus-Infected T Cells Obtained Using Immunomicroarrays: Induction of Proinflammatory Mechanisms. <i>Journal of Virology</i> , 2001, 75, 11641-11650.	1.5	78
51	CHD5, a Brain-Specific Paralog of Mi2 Chromatin Remodeling Enzymes, Regulates Expression of Neuronal Genes. <i>PLoS ONE</i> , 2011, 6, e24515.	1.1	76
52	Extension of Lifespan in <i>C. elegans</i> by Naphthoquinones That Act through Stress Hormesis Mechanisms. <i>PLoS ONE</i> , 2011, 6, e21922.	1.1	76
53	Methamphetamine Induces Dopamine D1 Receptor-Dependent Endoplasmic Reticulum Stress-Related Molecular Events in the Rat Striatum. <i>PLoS ONE</i> , 2009, 4, e6092.	1.1	76
54	Extremely Long-Range Chromatin Loops Link Topological Domains to Facilitate a Diverse Antibody Repertoire. <i>Cell Reports</i> , 2016, 14, 896-906.	2.9	75

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55	MNKs act as a regulatory switch for eIF4E1 and eIF4E3 driven mRNA translation in DLBCL. <i>Nature Communications</i> , 2014, 5, 5413.	5.8	73
56	Role of heparin binding growth factors in nigrostriatal dopamine system development and Parkinson's disease. <i>Brain Research</i> , 2007, 1147, 77-88.	1.1	71
57	Nontelomeric TRF2-REST Interaction Modulates Neuronal Gene Silencing and Fate of Tumor and Stem Cells. <i>Current Biology</i> , 2008, 18, 1489-1494.	1.8	71
58	VENNTUREâ€“A Novel Venn Diagram Investigational Tool for Multiple Pharmacological Dataset Analysis. <i>PLoS ONE</i> , 2012, 7, e36911.	1.1	71
59	Hippocampal Transcriptomic and Proteomic Alterations in the BTBR Mouse Model of Autism Spectrum Disorder. <i>Frontiers in Physiology</i> , 2015, 6, 324.	1.3	70
60	Absence of cannabinoid 1 receptor in beta cells protects against high-fat/high-sugar diet-induced beta cell dysfunction and inflammation in murine islets. <i>Diabetologia</i> , 2018, 61, 1470-1483.	2.9	69
61	Protease Activated Receptor Signaling Is Required for African Trypanosome Traversal of Human Brain Microvascular Endothelial Cells. <i>PLoS Neglected Tropical Diseases</i> , 2009, 3, e479.	1.3	68
62	An overview of the BioCreative 2012 Workshop Track III: interactive text mining task. <i>Database: the Journal of Biological Databases and Curation</i> , 2013, 2013, bas056-bas056.	1.4	68
63	Claudin-7 Is Frequently Overexpressed in Ovarian Cancer and Promotes Invasion. <i>PLoS ONE</i> , 2011, 6, e22119.	1.1	66
64	Genetic and environmental pathways to complex diseases. <i>BMC Systems Biology</i> , 2009, 3, 46.	3.0	65
65	Application of z-score transformation to Affymetrix data. <i>Applied Bioinformatics</i> , 2003, 2, 209-17.	1.7	64
66	Minimal Peroxide Exposure of Neuronal Cells Induces Multifaceted Adaptive Responses. <i>PLoS ONE</i> , 2010, 5, e14352.	1.1	61
67	Gene expression and pathway analysis of ovarian cancer cells selected for resistance to cisplatin, paclitaxel, or doxorubicin. <i>Journal of Ovarian Research</i> , 2011, 4, 21.	1.3	61
68	MIR100 host gene-encoded lncRNAs regulate cell cycle by modulating the interaction between HuR and its target mRNAs. <i>Nucleic Acids Research</i> , 2018, 46, 10405-10416.	6.5	61
69	Histone acetylation is associated with differential gene expression in the rapid and robust memory CD8+ T-cell response. <i>Blood</i> , 2006, 108, 3363-3370.	0.6	60
70	Effects of aging and calorie restriction on the global gene expression profiles of mouse testis and ovary. <i>BMC Biology</i> , 2008, 6, 24.	1.7	59
71	Microarray Analysis Reveals Interleukin-6 as a Novel Secretory Product of the Hypothalamo-neurohypophyseal System. <i>Journal of Biological Chemistry</i> , 2003, 278, 19280-19285.	1.6	58
72	Elongator and codon bias regulate protein levels in mammalian peripheral neurons. <i>Nature Communications</i> , 2018, 9, 889.	5.8	58

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73	A Mechanism for the Inhibition of Neural Progenitor Cell Proliferation by Cocaine. <i>PLoS Medicine</i> , 2008, 5, e117.	3.9	58
74	Dynamic BRG1 Recruitment during T Helper Differentiation and Activation Reveals Distal Regulatory Elements. <i>Molecular and Cellular Biology</i> , 2011, 31, 1512-1527.	1.1	56
75	Single Round of Antigen Receptor Signaling Programs Naive B Cells to Receive T Cell Help. <i>Immunity</i> , 2010, 32, 355-366.	6.6	54
76	The mitochondrial uncoupler <i>DNP</i> triggers brain cell <i>mTOR</i> signaling network reprogramming and <i>CREB</i> pathway up-regulation. <i>Journal of Neurochemistry</i> , 2015, 134, 677-692.	2.1	53
77	Novel RNA-binding activity of MYF5 enhances <i>Ccnd1</i> / <i>Cyclin D1</i> mRNA translation during myogenesis. <i>Nucleic Acids Research</i> , 2016, 44, 2393-2408.	6.5	52
78	Long-Term Artificial Sweetener Acesulfame Potassium Treatment Alters Neurometabolic Functions in C57BL/6J Mice. <i>PLoS ONE</i> , 2013, 8, e70257.	1.1	50
79	TCR <sup>2</sup> repertoire of CD4+ and CD8+ T cells is distinct in richness, distribution, and CDR3 amino acid composition. <i>Journal of Leukocyte Biology</i> , 2016, 99, 505-513.	1.5	50
80	Transcriptional Changes Common to Human Cocaine, Cannabis and Phencyclidine Abuse. <i>PLoS ONE</i> , 2006, 1, e114.	1.1	50
81	Methamphetamine Preconditioning Alters Midbrain Transcriptional Responses to Methamphetamine-Induced Injury in the Rat Striatum. <i>PLoS ONE</i> , 2009, 4, e7812.	1.1	49
82	Non-Steroidal Anti-inflammatory Drugs Decrease E2F1 Expression and Inhibit Cell Growth in Ovarian Cancer Cells. <i>PLoS ONE</i> , 2013, 8, e61836.	1.1	49
83	Blast traumatic brain injury-induced cognitive deficits are attenuated by preinjury or postinjury treatment with the glucagon-like peptide-1 receptor agonist, exendin-4. <i>Alzheimer's and Dementia</i> , 2016, 12, 34-48.	0.4	48
84	Sporadic Alzheimer disease fibroblasts display an oxidative stress phenotype. <i>Free Radical Biology and Medicine</i> , 2012, 53, 1371-1380.	1.3	47
85	Conserved and Differential Effects of Dietary Energy Intake on the Hippocampal Transcriptomes of Females and Males. <i>PLoS ONE</i> , 2008, 3, e2398.	1.1	46
86	Disulfiram Treatment Normalizes Body Weight in Obese Mice. <i>Cell Metabolism</i> , 2020, 32, 203-214.e4.	7.2	46
87	Age-Related Brain Expression and Regulation of the Chemokine CCL4/MIP-1 <sup>2</sup> in APP/PS1 Double-Transgenic Mice. <i>Journal of Neuropathology and Experimental Neurology</i> , 2014, 73, 362-374.	0.9	45
88	Metabolic and molecular framework for the enhancement of endurance by intermittent food deprivation. <i>FASEB Journal</i> , 2018, 32, 3844-3858.	0.2	45
89	Growth Factor Signals in Neural Cells. <i>Journal of Biological Chemistry</i> , 2009, 284, 2493-2511.	1.6	44
90	Age-associated changes in basal NF- $\kappa$ B function in human CD4+ T lymphocytes via dysregulation of PI3 kinase. <i>Aging</i> , 2014, 6, 957-969.	1.4	44

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91	Identification of Transformation-Related Pathways in a Breast Epithelial Cell Model Using a Ribonomics Approach. <i>Cancer Research</i> , 2008, 68, 7730-7735.	0.4	43
92	Genome-wide profiling identifies a subset of methamphetamine (METH)-induced genes associated with METH-induced increased H4K5Ac binding in the rat striatum. <i>BMC Genomics</i> , 2013, 14, 545.	1.2	43
93	AUF1 promotes let-7b loading on Argonaute 2. <i>Genes and Development</i> , 2015, 29, 1599-1604.	2.7	41
94	Delineation of a Conserved Arrestin-Biased Signaling Repertoire In Vivo. <i>Molecular Pharmacology</i> , 2015, 87, 706-717.	1.0	40
95	Sarcopenia, Aging and Prospective Interventional Strategies. <i>Current Medicinal Chemistry</i> , 2019, 25, 5588-5596.	1.2	40
96	Down-Regulation of eIF4GII by miR-520c-3p Represses Diffuse Large B Cell Lymphoma Development. <i>PLoS Genetics</i> , 2014, 10, e1004105.	1.5	39
97	RNA-Binding Protein AUF1 Promotes Myogenesis by Regulating MEF2C Expression Levels. <i>Molecular and Cellular Biology</i> , 2014, 34, 3106-3119.	1.1	39
98	Platelet MicroRNAs: An Overview. <i>Transfusion Medicine Reviews</i> , 2015, 29, 215-219.	0.9	39
99	Transcriptional outcomes and kinetic patterning of gene expression in response to NF- $\kappa$ B activation. <i>PLoS Biology</i> , 2018, 16, e2006347.	2.6	37
100	Time-Dependent c-Myc Transactomes Mapped by Array-Based Nuclear Run-On Reveal Transcriptional Modules in Human B Cells. <i>PLoS ONE</i> , 2010, 5, e9691.	1.1	37
101	The sharing of cDNA microarray data. <i>Nature Reviews Neuroscience</i> , 2001, 2, 438-440.	4.9	36
102	ATM regulates a DNA damage response posttranscriptional RNA operon in lymphocytes. <i>Blood</i> , 2011, 117, 2441-2450.	0.6	36
103	Cognitive Impairments Induced by Concussive Mild Traumatic Brain Injury in Mouse Are Ameliorated by Treatment with Phenserine via Multiple Non-Cholinergic and Cholinergic Mechanisms. <i>PLoS ONE</i> , 2016, 11, e0156493.	1.1	36
104	A Murine Dopamine Neuron-Specific cDNA Library and Microarray: Increased COXI Expression during Methamphetamine Neurotoxicity. <i>Neurobiology of Disease</i> , 2001, 8, 822-833.	2.1	35
105	Transcriptional profiling in an MPNST-derived cell line and normal human Schwann cells. <i>Neuron Glia Biology</i> , 2004, 1, 135-147.	2.0	35
106	Sequential Enhancer Sequestration Dysregulates Recombination Center Formation at the IgH Locus. <i>Molecular Cell</i> , 2018, 70, 21-33.e6.	4.5	35
107	Enhanced Upregulation of CRH mRNA Expression in the Nucleus Accumbens of Male Rats after a Second Injection of Methamphetamine Given Thirty Days Later. <i>PLoS ONE</i> , 2014, 9, e84665.	1.1	35
108	Novel RNA- and FMRP-binding protein TRF2-S regulates axonal mRNA transport and presynaptic plasticity. <i>Nature Communications</i> , 2015, 6, 8888.	5.8	34

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109	Gonadal Transcriptome Alterations in Response to Dietary Energy Intake: Sensing the Reproductive Environment. <i>PLoS ONE</i> , 2009, 4, e4146.	1.1	33
110	Molecular changes in brain aging and Alzheimer's disease are mirrored in experimentally silenced cortical neuron networks. <i>Neurobiology of Aging</i> , 2012, 33, 205.e1-205.e18.	1.5	33
111	Possible angiogenic roles for claudin-4 in ovarian cancer. <i>Cancer Biology and Therapy</i> , 2009, 8, 1806-1814.	1.5	32
112	IL-10 transcription is negatively regulated by BAF180, a component of the SWI/SNF chromatin remodeling enzyme. <i>BMC Immunology</i> , 2012, 13, 9.	0.9	32
113	DNA methylation signatures reveal that distinct combinations of transcription factors specify human immune cell epigenetic identity. <i>Immunity</i> , 2021, 54, 2465-2480.e5.	6.6	31
114	Differential gene expression patterns in cyclooxygenase-1 and cyclooxygenase-2 deficient mouse brain. <i>Genome Biology</i> , 2007, 8, R14.	13.9	30
115	Euglycemic Agent-mediated Hypothalamic Transcriptomic Manipulation in the N171 $\hat{a}$ €82Q Model of Huntington Disease Is Related to Their Physiological Efficacy*. <i>Journal of Biological Chemistry</i> , 2012, 287, 31766-31782.	1.6	30
116	AF5, a CNS Cell Line Immortalized with an N-Terminal Fragment of SV40 Large T: Growth, Differentiation, Genetic Stability, and Gene Expression. <i>Experimental Neurology</i> , 2002, 175, 318-337.	2.0	29
117	Transcriptome analysis of age-, gender- and diet-associated changes in murine thymus. <i>Cellular Immunology</i> , 2007, 245, 42-61.	1.4	29
118	Divergence of transcriptional landscape occurs early in B cell activation. <i>Epigenetics and Chromatin</i> , 2015, 8, 20.	1.8	28
119	Mild traumatic brain injury-induced hippocampal gene expressions: The identification of target cellular processes for drug development. <i>Journal of Neuroscience Methods</i> , 2016, 272, 4-18.	1.3	28
120	Profile of changes in gene expression in cultured hippocampal neurones evoked by the GABAB receptor agonist baclofen. <i>Physiological Genomics</i> , 2005, 22, 93-98.	1.0	27
121	MyD88 Expression by CNS-Resident Cells is Pivotal for Eliciting Protective Immunity in Brain Abscesses. <i>ASN Neuro</i> , 2009, 1, AN20090004.	1.5	27
122	Topoisomerase 3 $\hat{I}$ 2 interacts with RNAi machinery to promote heterochromatin formation and transcriptional silencing in <i>Drosophila</i> . <i>Nature Communications</i> , 2018, 9, 4946.	5.8	27
123	Extracellular <sc>RNA</sc> profiles with human age. <i>Aging Cell</i> , 2018, 17, e12785.	3.0	27
124	Neonatal dopamine depletion induces changes in morphogenesis and gene expression in the developing cortex. <i>Neurotoxicity Research</i> , 2007, 11, 107-130.	1.3	26
125	Paradoxical microRNAs. <i>Cell Cycle</i> , 2011, 10, 751-759.	1.3	26
126	miR-570 interacts with mitochondrial ATPase subunit g (ATP5L) encoding mRNA in stored platelets. <i>Platelets</i> , 2017, 28, 74-81.	1.1	26



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127	Muscle cannabinoid 1 receptor regulates $\text{IL-6}$ and myostatin expression, governing physical performance and whole-body metabolism. <i>FASEB Journal</i> , 2019, 33, 5850-5863.	0.2	26
128	Age-associated alterations in inducible gene transcription in human CD4+ T lymphocytes. <i>Aging</i> , 2013, 5, 18-36.	1.4	26
129	Methamphetamine-Induced Dopamine-Independent Alterations in Striatal Gene Expression in the 6-Hydroxydopamine Hemiparkinsonian Rats. <i>PLoS ONE</i> , 2010, 5, e15643.	1.1	25
130	Methamphetamine Preconditioning Causes Differential Changes in Striatal Transcriptional Responses to Large Doses of the Drug. <i>Dose-Response</i> , 2011, 9, dose-response.1.	0.7	25
131	GIT2 Acts as a Systems-Level Coordinator of Neurometabolic Activity and Pathophysiological Aging. <i>Frontiers in Endocrinology</i> , 2015, 6, 191.	1.5	25
132	A rapid method for microarray cross platform comparisons using gene expression signatures. <i>Molecular and Cellular Probes</i> , 2007, 21, 35-46.	0.9	24
133	Gene Expression Profiling Reveals Distinct Cocaine-Responsive Genes in Human Fetal CNS Cell Types. <i>Journal of Addiction Medicine</i> , 2009, 3, 218-226.	1.4	24
134	Multiple Oxygen Tension Environments Reveal Diverse Patterns of Transcriptional Regulation in Primary Astrocytes. <i>PLoS ONE</i> , 2011, 6, e21638.	1.1	24
135	Human Umbilical Cord Matrix Mesenchymal Stem Cells Suppress the Growth of Breast Cancer by Expression of Tumor Suppressor Genes. <i>PLoS ONE</i> , 2015, 10, e0123756.	1.1	22
136	Topoisomerase $3\beta$ knockout mice show transcriptional and behavioural impairments associated with neurogenesis and synaptic plasticity. <i>Nature Communications</i> , 2020, 11, 3143.	5.8	22
137	Co-Regulation of the DAF-16 Target Gene, <i>cyp-35B1/dod-13</i> , by HSF-1 in <i>C. elegans</i> Dauer Larvae and <i>daf-2</i> Insulin Pathway Mutants. <i>PLoS ONE</i> , 2011, 6, e17369.	1.1	21
138	Cortical gene transcription response patterns to water maze training in aged mice. <i>BMC Neuroscience</i> , 2011, 12, 63.	0.8	21
139	Toll-like receptor 2 (TLR2)-TLR9 crosstalk dictates $\text{IL-12}$ family cytokine production in microglia. <i>Glia</i> , 2012, 60, 29-42.	2.5	21
140	RPTOR, a novel target of miR-155, elicits a fibrotic phenotype of cystic fibrosis lung epithelium by upregulating CTGF. <i>RNA Biology</i> , 2016, 13, 837-847.	1.5	21
141	Conserved and species-specific molecular denominators in mammalian skeletal muscle aging. <i>Npj Aging and Mechanisms of Disease</i> , 2017, 3, 8.	4.5	21
142	MicroRNAs Modulate Oxidative Stress in Hypertension through PARP-1 Regulation. <i>Oxidative Medicine and Cellular Longevity</i> , 2017, 2017, 1-12.	1.9	21
143	Tumor-Derived Thymic Stromal Lymphopoietin Expands Bone Marrow B-cell Precursors in Circulation to Support Metastasis. <i>Cancer Research</i> , 2019, 79, 5826-5838.	0.4	21
144	Microarray screening of suppression subtractive hybridization-PCR cDNA libraries identifies novel RNAs regulated by dehydration in the rat supraoptic nucleus. <i>Physiological Genomics</i> , 2006, 24, 163-172.	1.0	20

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145	Molecular analysis of bovine spongiform encephalopathy infection by cDNA arrays. <i>Journal of General Virology</i> , 2007, 88, 1356-1362.	1.3	20
146	Significant modulation of mitochondrial electron transport system by nicotine in various rat brain regions. <i>Mitochondrion</i> , 2009, 9, 186-195.	1.6	20
147	Distinct inhibitory effects on mTOR signaling by ethanol and INK128 in diffuse large B-cell lymphoma. <i>Cell Communication and Signaling</i> , 2015, 13, 15.	2.7	20
148	Experience Modulates the Effects of Histone Deacetylase Inhibitors on Gene and Protein Expression in the Hippocampus: Impaired Plasticity in Aging. <i>Journal of Neuroscience</i> , 2015, 35, 11729-11742.	1.7	20
149	Altered learning, memory, and social behavior in type 1 taste receptor subunit 3 knock-out mice are associated with neuronal dysfunction. <i>Journal of Biological Chemistry</i> , 2017, 292, 11508-11530.	1.6	20
150	Na <sup>+</sup> ve rat umbilical cord matrix stem cells significantly attenuate mammary tumor growth through modulation of endogenous immune responses. <i>Cytotherapy</i> , 2013, 15, 586-597.	0.3	18
151	Amitriptyline Improves Motor Function via Enhanced Neurotrophin Signaling and Mitochondrial Functions in the Murine N171-82Q Huntington Disease Model. <i>Journal of Biological Chemistry</i> , 2015, 290, 2728-2743.	1.6	18
152	A double blind placebo controlled randomized trial of the effect of acute uric acid changes on inflammatory markers in humans: A pilot study. <i>PLoS ONE</i> , 2017, 12, e0181100.	1.1	18
153	Disease and phenotype gene set analysis of disease-based gene expression in mouse and human. <i>Physiological Genomics</i> , 2010, 42A, 162-167.	1.0	17
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