## Derek E Wildman

List of Publications by Year in descending order

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141 8,970 papers citations

155

all docs

155 docs citations 50 h-index

> 155 times ranked

48315 88 g-index

12582 citing authors

#	Article	IF	Citations
1	Prolonged myelination in human neocortical evolution. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 16480-16485.	7.1	492
2	Epigenetic and immune function profiles associated with posttraumatic stress disorder. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 9470-9475.	7.1	452
3	Metabolic costs and evolutionary implications of human brain development. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 13010-13015.	7.1	409
4	Largest GWAS of PTSD (N=20 070) yields genetic overlap with schizophrenia and sex differences in heritability. Molecular Psychiatry, 2018, 23, 666-673.	7.9	374
5	Evolution of the mammalian placenta revealed by phylogenetic analysis. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 3203-3208.	7.1	304
6	Evolution of increased glia–neuron ratios in the human frontal cortex. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 13606-13611.	7.1	303
7	Implications of natural selection in shaping 99.4% nonsynonymous DNA identity between humans and chimpanzees: Enlarging genus <i>Homo</i> . Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 7181-7188.	7.1	231
8	The common marmoset genome provides insight into primate biology and evolution. Nature Genetics, 2014, 46, 850-857.	21.4	225
9	A genome-wide association study of post-traumatic stress disorder identifies the retinoid-related orphan receptor alpha (RORA) gene as a significant risk locus. Molecular Psychiatry, 2013, 18, 937-942.	7.9	217
10	Sister grouping of chimpanzees and humans as revealed by genome-wide phylogenetic analysis of brain gene expression profiles. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 2957-2962.	7.1	213
11	A primate subfamily of galectins expressed at the maternal–fetal interface that promote immune cell death. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 9731-9736.	7.1	200
12	Accelerated evolution of the electron transport chain in anthropoid primates. Trends in Genetics, 2004, 20, 578-585.	6.7	181
13	Phylogenetic relationships and divergence times among New World monkeys (Platyrrhini, Primates). Molecular Phylogenetics and Evolution, 2006, 40, 274-280.	2.7	161
14	Genomics, biogeography, and the diversification of placental mammals. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 14395-14400.	7.1	158
15	Epigenetic and inflammatory marker profiles associated with depression in a community-based epidemiologic sample. Psychological Medicine, 2011, 41, 997-1007.	4.5	156
16	Comparative analysis of encephalization in mammals reveals relaxed constraints on anthropoid primate and cetacean brain scaling. Journal of Evolutionary Biology, 2012, 25, 981-994.	1.7	147
17	SLC6A4 methylation modifies the effect of the number of traumatic events on risk for posttraumatic stress disorder. Depression and Anxiety, 2011, 28, 639-647.	4.1	140
18	Sentieon DNASeq Variant Calling Workflow Demonstrates Strong Computational Performance and Accuracy. Frontiers in Genetics, 2019, 10, 736.	2.3	131

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19	Genome-wide adaptive complexes to underground stresses in blind mole rats Spalax. Nature Communications, 2014, 5, 3966.	12.8	124
20	Synaptogenesis and development of pyramidal neuron dendritic morphology in the chimpanzee neocortex resembles humans. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 10395-10401.	7.1	112
21	Genome-wide association study implicates a novel RNA gene, the lincRNA AC068718.1, as a risk factor for post-traumatic stress disorder in women. Psychoneuroendocrinology, 2013, 38, 3029-3038.	2.7	105
22	Molecular evolution tracks macroevolutionary transitions in Cetacea. Trends in Ecology and Evolution, 2014, 29, 336-346.	8.7	105
23	A fully resolved genus level phylogeny of neotropical primates (Platyrrhini). Molecular Phylogenetics and Evolution, 2009, 53, 694-702.	2.7	102
24	Convergent Evolution of Endometrial Prolactin Expression in Primates, Mice, and Elephants Through the Independent Recruitment of Transposable Elements. Molecular Biology and Evolution, 2012, 29, 239-247.	8.9	100
25	Moving primate genomics beyond the chimpanzee genome. Trends in Genetics, 2005, 21, 511-517.	6.7	93
26	Emergence of hormonal and redox regulation of galectin-1 in placental mammals: Implication in maternal–fetal immune tolerance. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 15819-15824.	7.1	86
27	Dolphin genome provides evidence for adaptive evolution of nervous system genes and a molecular rate slowdown. Proceedings of the Royal Society B: Biological Sciences, 2012, 279, 3643-3651.	2.6	86
28	Cytochrome c oxidase: Evolution of control via nuclear subunit addition. Biochimica Et Biophysica Acta - Bioenergetics, 2012, 1817, 590-597.	1.0	86
29	Glucocorticoid receptor DNA methylation, childhood maltreatment and major depression. Journal of Affective Disorders, 2016, 206, 181-188.	4.1	83
30	Galectins: guardians of eutherian pregnancy at the maternal–fetal interface. Trends in Endocrinology and Metabolism, 2012, 23, 23-31.	7.1	82
31	Molecular Evolution of Aerobic Energy Metabolism in Primates. Molecular Phylogenetics and Evolution, 2001, 18, 26-36.	2.7	81
32	DICER1 and microRNA regulation in post-traumatic stress disorder with comorbid depression. Nature Communications, 2015, 6, 10106.	12.8	81
33	Rapid electrostatic evolution at the binding site for cytochrome c on cytochrome c oxidase in anthropoid primates. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 6379-6384.	7.1	79
34	<i>ADCYAP1R1</i> GENOTYPE, POSTTRAUMATIC STRESS DISORDER, AND DEPRESSION AMONG WOMEN EXPOSED TO CHILDHOOD MALTREATMENT. Depression and Anxiety, 2013, 30, 251-258.	4.1	77
35	An analysis of gene expression in PTSD implicates genes involved in the glucocorticoid receptor pathway and neural responses to stress. Psychoneuroendocrinology, 2015, 57, 1-13.	2.7	77
36	Adaptive evolution of cytochrome c oxidase subunit VIII in anthropoid primates. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 5873-5878.	7.1	76

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37	Phylogenomic analyses reveal convergent patterns of adaptive evolution in elephant and human ancestries. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 20824-20829.	7.1	75
38	Functionally important glycosyltransferase gain and loss during catarrhine primate emergence. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 559-564.	7.1	74
39	Gene Expression and Methylation Signatures of <i>MAN2C1</i> are Associated with PTSD. Disease Markers, 2011, 30, 111-121.	1.3	69
40	Extracellular Vesicles and the Promise of Continuous Liquid Biopsies. Journal of Pathology and Translational Medicine, 2018, 52, 1-8.	1.1	68
41	Evolutionary genetics and implications of small size and twinning in callitrichine primates.  Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 1467-1472.	7.1	66
42	The tempo and mode of New World monkey evolution and biogeography in the context of phylogenomic analysis. Molecular Phylogenetics and Evolution, 2015, 82, 386-399.	2.7	66
43	Severe preeclampsia is characterized by increased placental expression of galectin-1. Journal of Maternal-Fetal and Neonatal Medicine, 2008, 21, 429-442.	1.5	65
44	Elevated systemic expression of ER stress related genes is associated with stress-related mental disorders in the Detroit Neighborhood Health Study. Psychoneuroendocrinology, 2014, 43, 62-70.	2.7	65
45	Mitochondrial evidence for the origin of hamadryas baboons. Molecular Phylogenetics and Evolution, 2004, 32, 287-296.	2.7	64
46	Molecular Variation at the SLC6A3 Locus Predicts Lifetime Risk of PTSD in the Detroit Neighborhood Health Study. PLoS ONE, 2012, 7, e39184.	2.5	64
47	Evaluating the impact of trauma and PTSD on epigenetic prediction of lifespan and neural integrity. Neuropsychopharmacology, 2020, 45, 1609-1616.	5.4	63
48	The core transcriptome of mammalian placentas and the divergence of expression with placental shape. Placenta, 2017, 57, 71-78.	1.5	62
49	Distinct genomic signatures of adaptation in pre- and postnatal environments during human evolution. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 3215-3220.	7.1	61
50	Developmental Changes in the Transcriptome of Human Cerebral Cortex Tissue: Long Noncoding RNA Transcripts. Cerebral Cortex, 2014, 24, 1451-1459.	2.9	58
51	Interaction between polygenic risk for cigarette use and environmental exposures in the Detroit neighborhood health study. Translational Psychiatry, 2013, 3, e290-e290.	4.8	52
52	Gene expression and methylation signatures of MAN2C1 are associated with PTSD. Disease Markers, 2011, 30, 111-21.	1.3	51
53	Geobiology reveals how human kidney stones dissolve in vivo. Scientific Reports, 2018, 8, 13731.	3.3	50
54	Genetic Association Analysis of 300 Genes Identifies a Risk Haplotype in SLC18A2 for Post-traumatic Stress Disorder in Two Independent Samples. Neuropsychopharmacology, 2014, 39, 1872-1879.	5.4	49

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55	The evolution of embryo implantation. International Journal of Developmental Biology, 2014, 58, 155-161.	0.6	48
56	Molecular evolution of the cytochrome c oxidase subunit 5A gene in primates. BMC Evolutionary Biology, 2008, 8, 8.	3.2	46
57	Genomic data reject the hypothesis of a prosimian primate clade. Journal of Human Evolution, 2011, 61, 295-305.	2.6	45
58	Longitudinal epigenetic variation of DNA methyltransferase genes is associated with vulnerability to post-traumatic stress disorder. Psychological Medicine, 2014, 44, 3165-3179.	4.5	45
59	Adrenal androgen production in catarrhine primates and the evolution of adrenarche. American Journal of Physical Anthropology, 2012, 147, 389-400.	2.1	44
60	A review of inter- and intraspecific variation in the eutherian placenta. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20140072.	4.0	44
61	FKBP5 DNA methylation does not mediate the association between childhood maltreatment and depression symptom severity in the Detroit Neighborhood Health Study. Journal of Psychiatric Research, 2018, 96, 39-48.	3.1	44
62	ORIGINAL ARTICLE: Chorioamnionitis and Increased Galectinâ€1 Expression in PPROM – An Antiâ€Inflammatory Response in the Fetal Membranes?. American Journal of Reproductive Immunology, 2008, 60, 298-311.	1.2	43
63	Over-expression of the thrombin receptor (PAR-1) in the placenta in preeclampsia: A mechanism for the intersection of coagulation and inflammation. Journal of Maternal-Fetal and Neonatal Medicine, 2008, 21, 345-355.	1.5	42
64	Inhibitory interneurons of the human prefrontal cortex display conserved evolution of the phenotype and related genes. Proceedings of the Royal Society B: Biological Sciences, 2010, 277, 1011-1020.	2.6	42
65	Frequency of alcohol consumption in humans; the role of metabotropic glutamate receptors and downstream signaling pathways. Translational Psychiatry, 2015, 5, e586-e586.	4.8	39
66	PTSD is associated with an increase in aged T cell phenotypes in adults living in Detroit. Psychoneuroendocrinology, 2016, 67, 133-141.	2.7	39
67	Epigenetic meta-analysis across three civilian cohorts identifies <i>NRG1</i> and <i>HGS</i> as blood-based biomarkers for post-traumatic stress disorder. Epigenomics, 2018, 10, 1585-1601.	2.1	39
68	Coadaptive evolution in cytochrome c oxidase: 9 of 13 subunits show accelerated rates of nonsynonymous substitution in anthropoid primates. Molecular Phylogenetics and Evolution, 2004, 33, 944-950.	2.7	33
69	The human progesterone receptor shows evidence of adaptive evolution associated with its ability to act as a transcription factor. Molecular Phylogenetics and Evolution, 2008, 47, 637-649.	2.7	33
70	Functional Divergence of the Nuclear Receptor <i>NR2C1</i> as a Modulator of Pluripotentiality During Hominid Evolution. Genetics, 2016, 203, 905-922.	2.9	33
71	Dynamic Gene Expression in the Human Cerebral Cortex Distinguishes Children from Adults. PLoS ONE, 2012, 7, e37714.	2.5	32
72	Income and Markers of Immunological Cellular Aging. Psychosomatic Medicine, 2016, 78, 657-666.	2.0	32

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73	Ancient origin of placental expression in the growth hormone genes of anthropoid primates. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 17083-17088.	7.1	31
74	Epigenetic Signatures May Explain the Relationship between Socioeconomic Position and Risk of Mental Illness: Preliminary Findings from an Urban Community-Based Sample. Biodemography and Social Biology, 2013, 59, 68-84.	1.0	31
75	Evidence of a Conserved Molecular Response to Selection for Increased Brain Size in Primates. Genome Biology and Evolution, 2017, 9, 700-713.	2.5	31
76	Asthma-Related Immune Responses in Youth With Asthma. Psychosomatic Medicine, 2015, 77, 892-902.	2.0	30
77	Noninvasive methods for collecting fresh hair tissue. Molecular Ecology, 1999, 8, 1749-1750.	3.9	29
78	Transcriptional Regulation of the Novel Toll-like Receptor Tlr13. Journal of Biological Chemistry, 2009, 284, 20540-20547.	3.4	29
79	PTSD and obesity in the Detroit neighborhood health study. General Hospital Psychiatry, 2013, 35, 671-673.	2.4	29
80	RORA and posttraumatic stress trajectories: main effects and interactions with childhood physical abuse history. Brain and Behavior, 2015, 5, e00323.	2.2	29
81	The Dopamine D <sub>3</sub> Receptor Gene and Posttraumatic Stress Disorder. Journal of Traumatic Stress, 2014, 27, 379-387.	1.8	28
82	Associations between the SS variant of 5-HTTLPR and PTSD among adults with histories of childhood emotional abuse: Results from two African American independent samples. Journal of Affective Disorders, 2014, 161, 91-96.	4.1	28
83	Epigenetic predictors of all-cause mortality are associated with objective measures of neighborhood disadvantage in an urban population. Clinical Epigenetics, 2020, 12, 44.	4.1	28
84	Primate phylogenomics: developing numerous nuclear non-coding, non-repetitive markers for ecological and phylogenetic applications and analysis of evolutionary rate variation. BMC Genomics, 2009, 10, 247.	2.8	27
85	Adaptive history of single copy genes highly expressed in the term human placenta. Genomics, 2009, 93, 33-41.	2.9	27
86	Elephant Transcriptome Provides Insights into the Evolution of Eutherian Placentation. Genome Biology and Evolution, 2012, 4, 713-725.	2.5	27
87	Out of Africa, but how and when? The case of hamadryas baboons (Papio hamadryas). Journal of Human Evolution, 2014, 76, 154-164.	2.6	25
88	Recommendations for performance optimizations when using GATK3.8 and GATK4. BMC Bioinformatics, 2019, 20, 557.	2.6	25
89	The phylogenetic history of New World monkey $\hat{l}^2$ globin reveals a platyrrhine $\hat{l}^2$ to $\hat{l}'$ gene conversion in the atelid ancestry. Molecular Phylogenetics and Evolution, 2005, 35, 225-234.	2.7	23
90	Socioeconomic status, family negative emotional climate, and anti-inflammatory gene expression among youth with asthma. Psychoneuroendocrinology, 2018, 91, 62-67.	2.7	23

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91	Episodic Positive Selection in Ape Cytochrome c Oxidase Subunit IV. Molecular Biology and Evolution, 2002, 19, 1812-1815.	8.9	22
92	Evolution of the Couple Cytochrome c and Cytochrome c Oxidase in Primates. Advances in Experimental Medicine and Biology, 2012, 748, 185-213.	1.6	22
93	The rs1049353 Polymorphism in the CNR1 Gene Interacts With Childhood Abuse to Predict Posttraumatic Threat Symptoms. Journal of Clinical Psychiatry, 2015, 76, e1622-e1623.	2.2	22
94	Mothers' Attachment is Linked to Their Children's Anti-Inflammatory Gene Expression via Maternal Warmth. Social Psychological and Personality Science, 2017, 8, 796-805.	3.9	19
95	Neighborhood environment, social cohesion, and epigenetic aging. Aging, 2021, 13, 7883-7899.	3.1	19
96	A map of the common chimpanzee genome. BioEssays, 2002, 24, 490-493.	2.5	17
97	New Onto-Tools: Promoter-Express, nsSNPCounter and Onto-Translate. Nucleic Acids Research, 2006, 34, W626-W631.	14.5	17
98	Phylogeny of the Ferungulata (Mammalia: Laurasiatheria) as determined from phylogenomic data. Molecular Phylogenetics and Evolution, 2009, 52, 660-664.	2.7	17
99	Rapid Fractionation and Isolation of Whole Blood Components in Samples Obtained from a Community-based Setting. Journal of Visualized Experiments, 2015, , .	0.3	17
100	Phylogenetic comparisons suggest that distance from the locus control region guides developmental expression of primate beta-type globin genes. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 3186-3191.	7.1	16
101	Spontaneous Abortion and Preterm Labor and Delivery in Nonhuman Primates: Evidence from a Captive Colony of Chimpanzees (Pan troglodytes). PLoS ONE, 2011, 6, e24509.	2.5	16
102	Synaptosomal Lactate Dehydrogenase Isoenzyme Composition Is Shifted toward Aerobic Forms in Primate Brain Evolution. Brain, Behavior and Evolution, 2014, 83, 216-230.	1.7	16
103	<i>FKBP5</i> : A Key Mediator of How Vertebrates Flexibly Cope with Adversity. BioScience, 2020, 70, 1127-1138.	4.9	16
104	The impact of psychopathology, social adversity and stress-relevant DNA methylation on prospective risk for post-traumatic stress: A machine learning approach. Journal of Affective Disorders, 2021, 282, 894-905.	4.1	16
105	Psychosocial experiences modulate asthma-associated genes through gene-environment interactions. ELife, 2021, 10, .	6.0	15
106	Silencing, Positive Selection and Parallel Evolution: Busy History of Primate Cytochromes c. PLoS ONE, 2011, 6, e26269.	2.5	14
107	Methylomic profiles reveal sex-specific differences in leukocyte composition associated with post-traumatic stress disorder. Brain, Behavior, and Immunity, 2019, 81, 280-291.	4.1	14
108	Ancient evolutionary origins of epigenetic regulation associated with posttraumatic stress disorder. Frontiers in Human Neuroscience, 2014, 8, 284.	2.0	13

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109	Burden of post-traumatic stress disorder in postgenocide Rwandan population following exposure to 1994 genocide against the Tutsi: A meta-analysis. Journal of Affective Disorders, 2020, 275, 7-13.	4.1	13
110	Advancing human health in the decade ahead: pregnancy as a key window for discovery. American Journal of Obstetrics and Gynecology, 2020, 223, 312-321.	1.3	13
111	OCPAT: an online codon-preserved alignment tool for evolutionary genomic analysis of protein coding sequences. Source Code for Biology and Medicine, 2007, 2, 5.	1.7	11
112	Maternal weight affects placental DNA methylation of genes involved in metabolic pathways in the common marmoset monkey ( <i>Callithrix jacchus</i> ). American Journal of Primatology, 2020, 82, e23101.	1.7	10
113	Rapid Nonsynonymous Evolution of the Iron-Sulfur Protein in Anthropoid Primates. Journal of Bioenergetics and Biomembranes, 2005, 37, 35-41.	2.3	9
114	Development and evaluation of new mask protocols for gene expression profiling in humans and chimpanzees. BMC Bioinformatics, 2009, 10, 77.	2.6	9
115	IFPA award in placentology lecture: Phylogenomic origins and evolution of the mammalian placenta. Placenta, 2016, 48, S31-S39.	1.5	9
116	The association between residential proximity to brownfield sites and high-traffic areas and measures of immunity. Journal of Exposure Science and Environmental Epidemiology, 2020, 30, 824-834.	3.9	9
117	Humankind's Place in a Phylogenetic Classification of Living Primates. , 2004, , 293-311.		9
118	Divergent lactate dehydrogenase isoenzyme profile in cellular compartments of primate forebrain structures. Molecular and Cellular Neurosciences, 2017, 82, 137-142.	2.2	7
119	Leukocyte methylomic imprints of exposure to the genocide against the Tutsi in Rwanda: a pilot epigenome-wide analysis. Epigenomics, 2022, 14, 11-25.	2.1	7
120	Characterization of human cortical gene expression in relation to glucose utilization. American Journal of Human Biology, 2013, 25, 418-430.	1.6	6
121	Population Distributions of Thymic Function in Adults: Variation by Sociodemographic Characteristics and Health Status. Biodemography and Social Biology, 2016, 62, 208-221.	1.0	6
122	Socioeconomic status, financial stress, and glucocorticoid resistance among youth with asthma: Testing the moderation effects of maternal involvement and warmth. Brain, Behavior, and Immunity, 2021, 96, 92-99.	4.1	6
123	Relationship between Maternal and Fetal Plasma Glucose and Insulin Concentrations during Graded Maternal Hyperglycemic States in Primates. American Journal of Perinatology, 2006, 23, 369-376.	1.4	5
124	Development and annotation of shotgun sequence libraries from New World monkeys. Molecular Ecology Resources, 2012, 12, 950-955.	4.8	4
125	Heterogeneous Mediation Analysis on Epigenomic PTSD and Traumatic Stress in a Predominantly African American Cohort. Journal of the American Statistical Association, 2022, 117, 1669-1683.	3.1	4
126	A prospective and controlled in vivo in vivo letermine if acute episodes of high glucose concentrations in the extra-embryonic celomic cavity could be related to spontaneous abortion. Journal of Maternal-Fetal and Neonatal Medicine, 2012, 25, 1848-1851.	1.5	3

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127	From PPROM to caul: The evolution of membrane rupture in mammals. Applied & Translational Genomics, 2013, 2, 70-77.	2.1	3
128	Further Support for an Association between the Memory-Related Gene WWC1 and Posttraumatic Stress Disorder: Results from the Detroit Neighborhood Health Study. Biological Psychiatry, 2014, 76, e25-e26.	1.3	3
129	Transcriptomic profiling of fetal membranes of mice deficient in biglycan and decorin as a model of preterm birthâ€. Biology of Reproduction, 2021, 104, 611-623.	2.7	3
130	Trophoblast, Galectin-1 and pre-eclampsia. American Journal of Obstetrics and Gynecology, 2006, 195, S138.	1.3	2
131	Morris Goodman, Ph.D. (1925–2010). A Remembrance. Molecular Phylogenetics and Evolution, 2011, 58, 1-3.	2.7	2
132	Placental Development, Evolution, and Epigenetics of Primate Pregnancies., 2013,, 55-81.		2
133	Highâ€throughput RNA sequencing reveals structural differences of orthologous brainâ€expressed genes between western lowland gorillas and humans. Journal of Comparative Neurology, 2016, 524, 288-308.	1.6	2
134	DNA methylation of Nuclear Factor of Activated T Cells 1 mediates the prospective relation between exposure to different traumatic event types and post-traumatic stress disorder. Psychiatry Research, 2022, 311, 114510.	3.3	2
135	Reply to Skoyles: Decline in growth rate, not muscle mass, predicts the human childhood peak in brain metabolism. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E4910.	7.1	1
136	Ancestral resurrection of anthropoid estrogen receptor $\hat{l}^2$ demonstrates functional consequences of positive selection. Molecular Phylogenetics and Evolution, 2017, 117, 2-9.	2.7	1
137	Prediction of adenocarcinoma development using game theory. , 2017, 2017, 1668-1671.		1
138	IDChase: Mitigating Identifier Migration Trap in Biological Databases. Communications in Computer and Information Science, 2009, , 461-472.	0.5	1
139	Morris Goodman (1925–2010). Journal of Human Evolution, 2011, 60, 673-676.	2.6	0
140	Morris Goodman (1925–2010): Founder of the field of molecular anthropology. Evolutionary Anthropology, 2011, 20, 1-2.	3.4	0
141	Editorial for 25th Anniversary Issue of Molecular Phylogenetics and Evolution. Molecular Phylogenetics and Evolution, 2017, 117, 1.	2.7	0