Derek E Wildman

List of Publications by Year
in descending order

141 papers	8,970 citations	38660 50 h-index	48187 88 g-index
155 all docs	155 docs citations	155 times ranked	12582 citing authors

#	Article	IF	Citations
1	Leukocyte methylomic imprints of exposure to the genocide against the Tutsi in Rwanda: a pilot epigenome-wide analysis. Epigenomics, 2022, 14, 11-25.	1.0	7
2	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.	1.7	2
3	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.	1.8	4
4	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.	1.2	3
5	Neighborhood environment, social cohesion, and epigenetic aging. Aging, 2021, 13, 7883-7899.	1.4	19
6	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.	2.0	16
7	Psychosocial experiences modulate asthma-associated genes through gene-environment interactions. ELife, 2021, 10, .	2.8	15
8	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.	2.0	6
9	FKBP5: A Key Mediator of How Vertebrates Flexibly Cope with Adversity. BioScience, 2020, 70, 1127-1138.	2.2	16
10	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.	2.0	13
11	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.	1.8	9
12	Evaluating the impact of trauma and PTSD on epigenetic prediction of lifespan and neural integrity. Neuropsychopharmacology, 2020, 45, 1609-1616.	2.8	63
13	Advancing human health in the decade ahead: pregnancy as a key window for discovery. American Journal of Obstetrics and Gynecology, 2020, 223, 312-321.	0.7	13
14	Epigenetic predictors of all-cause mortality are associated with objective measures of neighborhood disadvantage in an urban population. Clinical Epigenetics, 2020, 12, 44.	1.8	28
15	Maternal weight affects placental DNA methylation of genes involved in metabolic pathways in the common marmoset monkey (Callithrix jacchus). American Journal of Primatology, 2020, 82, e23101.	0.8	10
16	Methylomic profiles reveal sex-specific differences in leukocyte composition associated with post-traumatic stress disorder. Brain, Behavior, and Immunity, 2019, 81, 280-291.	2.0	14
17	Sentieon DNASeq Variant Calling Workflow Demonstrates Strong Computational Performance and Accuracy. Frontiers in Genetics, 2019, 10, 736.	1.1	131
18	Recommendations for performance optimizations when using GATK3.8 and GATK4. BMC Bioinformatics, 2019, 20, 557.	1.2	25

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19	Socioeconomic status, family negative emotional climate, and anti-inflammatory gene expression among youth with asthma. Psychoneuroendocrinology, 2018, 91, 62-67.	1.3	23
20	Largest GWAS of PTSD (N=20â€‰070) yields genetic overlap with schizophrenia and sex differences in heritability. Molecular Psychiatry, 2018, 23, 666-673.	4.1	374
21	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.	1.5	44
22	Epigenetic meta-analysis across three civilian cohorts identifies NRG1 and HGS as blood-based biomarkers for post-traumatic stress disorder. Epigenomics, 2018, 10, 1585-1601.	1.0	39
23	Extracellular Vesicles and the Promise of Continuous Liquid Biopsies. Journal of Pathology and Translational Medicine, 2018, 52, 1-8.	0.4	68
24	Geobiology reveals how human kidney stones dissolve in vivo. Scientific Reports, 2018, 8, 13731.	1.6	50
25	Mothersâ€™ Attachment is Linked to Their Childrenâ€™s Anti-Inflammatory Gene Expression via Maternal Warmth. Social Psychological and Personality Science, 2017, 8, 796-805.	2.4	19
26	Divergent lactate dehydrogenase isoenzyme profile in cellular compartments of primate forebrain structures. Molecular and Cellular Neurosciences, 2017, 82, 137-142.	1.0	7
27	Ancestral resurrection of anthropoid estrogen receptor Î² demonstrates functional consequences of positive selection. Molecular Phylogenetics and Evolution, 2017, 117, 2-9.	1.2	1
28	The core transcriptome of mammalian placentas and the divergence of expression with placental shape. Placenta, 2017, 57, 71-78.	0.7	62
29	Prediction of adenocarcinoma development using game theory. , 2017, 2017, 1668-1671.		1
30	Editorial for 25th Anniversary Issue of Molecular Phylogenetics and Evolution. Molecular Phylogenetics and Evolution, 2017, 117, 1.	1.2	0
31	Evidence of a Conserved Molecular Response to Selection for Increased Brain Size in Primates. Genome Biology and Evolution, 2017, 9, 700-713.	1.1	31
32	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.	0.9	2
33	Income and Markers of Immunological Cellular Aging. Psychosomatic Medicine, 2016, 78, 657-666.	1.3	32
34	IFPA award in placentology lecture: Phylogenomic origins and evolution of the mammalian placenta. Placenta, 2016, 48, S31-S39.	0.7	9
35	Population Distributions of Thymic Function in Adults: Variation by Sociodemographic Characteristics and Health Status. Biodemography and Social Biology, 2016, 62, 208-221.	0.4	6
36	Glucocorticoid receptor DNA methylation, childhood maltreatment and major depression. Journal of Affective Disorders, 2016, 206, 181-188.	2.0	83

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37	Functional Divergence of the Nuclear ReceptorNR2C1as a Modulator of Pluripotentiality During Hominid Evolution. Genetics, 2016, 203, 905-922.	1.2	33
38	PTSD is associated with an increase in aged T cell phenotypes in adults living in Detroit. Psychoneuroendocrinology, 2016, 67, 133-141.	1.3	39
39	Rapid Fractionation and Isolation of Whole Blood Components in Samples Obtained from a Community-based Setting. Journal of Visualized Experiments, 2015, , .	0.2	17
40	RORA and posttraumatic stress trajectories: main effects and interactions with childhood physical abuse history. Brain and Behavior, 2015, 5, e00323.	1.0	29
41	Asthma-Related Immune Responses in Youth With Asthma. Psychosomatic Medicine, 2015, 77, 892-902.	1.3	30
42	DICER1 and microRNA regulation in post-traumatic stress disorder with comorbid depression. Nature Communications, 2015, 6, 10106.	5.8	81
43	A review of inter- and intraspecific variation in the eutherian placenta. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20140072.	1.8	44
44	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.	1.3	77
45	Frequency of alcohol consumption in humans; the role of metabotropic glutamate receptors and downstream signaling pathways. Translational Psychiatry, 2015, 5, e586-e586.	2.4	39
46	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.	1.2	66
47	The rs1049353 Polymorphism in the CNR1 Gene Interacts With Childhood Abuse to Predict Posttraumatic Threat Symptoms. Journal of Clinical Psychiatry, 2015, 76, e1622-e1623.	1.1	22
48	The evolution of embryo implantation. International Journal of Developmental Biology, 2014, 58, 155-161.	0.3	48
49	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.	2.8	49
50	Developmental Changes in the Transcriptome of Human Cerebral Cortex Tissue: Long Noncoding RNA Transcripts. Cerebral Cortex, 2014, 24, 1451-1459.	1.6	58
51	Out of Africa, but how and when? The case of hamadryas baboons (Papio hamadryas). Journal of Human Evolution, 2014, 76, 154-164.	1.3	25
52	The Dopamine D₃ Receptor Gene and Posttraumatic Stress Disorder. Journal of Traumatic Stress, 2014, 27, 379-387.	1.0	28
53	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.	3.3	66
54	Synaptosomal Lactate Dehydrogenase Isoenzyme Composition Is Shifted toward Aerobic Forms in Primate Brain Evolution. Brain, Behavior and Evolution, 2014, 83, 216-230.	0.9	16

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55	Ancient evolutionary origins of epigenetic regulation associated with posttraumatic stress disorder. Frontiers in Human Neuroscience, 2014, 8, 284.	1.0	13
56	Longitudinal epigenetic variation of DNA methyltransferase genes is associated with vulnerability to post-traumatic stress disorder. Psychological Medicine, 2014, 44, 3165-3179.	2.7	45
57	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.	1.3	65
58	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.	2.0	28
59	Molecular evolution tracks macroevolutionary transitions in Cetacea. Trends in Ecology and Evolution, 2014, 29, 336-346.	4.2	105
60	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.	3.3	1
61	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.	3.3	409
62	The common marmoset genome provides insight into primate biology and evolution. Nature Genetics, 2014, 46, 850-857.	9.4	225
63	Genome-wide adaptive complexes to underground stresses in blind mole rats Spalax. Nature Communications, 2014, 5, 3966.	5.8	124
64	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.	0.7	3
65	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.	0.4	31
66	PTSD and obesity in the Detroit neighborhood health study. General Hospital Psychiatry, 2013, 35, 671-673.	1.2	29
67	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.	1.3	105
68	From PPROM to caul: The evolution of membrane rupture in mammals. Applied & Translational Genomics, 2013, 2, 70-77.	2.1	3
69	Placental Development, Evolution, and Epigenetics of Primate Pregnancies. , 2013, , 55-81.		2
70	Characterization of human cortical gene expression in relation to glucose utilization. American Journal of Human Biology, 2013, 25, 418-430.	0.8	6
71	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.	3.3	112
72	ADCYAP1R1GENOTYPE, POSTTRAUMATIC STRESS DISORDER, AND DEPRESSION AMONG WOMEN EXPOSED TO CHILDHOOD MALTREATMENT. Depression and Anxiety, 2013, 30, 251-258.	2.0	77

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73	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.	4.1	217
74	Interaction between polygenic risk for cigarette use and environmental exposures in the Detroit neighborhood health study. Translational Psychiatry, 2013, 3, e290-e290.	2.4	52
75	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.	3.5	100
76	A prospective and controlledin vivostudy to determine 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.	0.7	3
77	Elephant Transcriptome Provides Insights into the Evolution of Eutherian Placentation. Genome Biology and Evolution, 2012, 4, 713-725.	1.1	27
78	Galectins: guardians of eutherian pregnancy at the maternalâ€“fetal interface. Trends in Endocrinology and Metabolism, 2012, 23, 23-31.	3.1	82
79	Prolonged myelination in human neocortical evolution. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 16480-16485.	3.3	492
80	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.	1.2	86
81	Dynamic Gene Expression in the Human Cerebral Cortex Distinguishes Children from Adults. PLoS ONE, 2012, 7, e37714.	1.1	32
82	Molecular Variation at the SLC6A3 Locus Predicts Lifetime Risk of PTSD in the Detroit Neighborhood Health Study. PLoS ONE, 2012, 7, e39184.	1.1	64
83	Evolution of the Couple Cytochrome c and Cytochrome c Oxidase in Primates. Advances in Experimental Medicine and Biology, 2012, 748, 185-213.	0.8	22
84	Cytochrome c oxidase: Evolution of control via nuclear subunit addition. Biochimica Et Biophysica Acta - Bioenergetics, 2012, 1817, 590-597.	0.5	86
85	Development and annotation of shotgun sequence libraries from New World monkeys. Molecular Ecology Resources, 2012, 12, 950-955.	2.2	4
86	Comparative analysis of encephalization in mammals reveals relaxed constraints on anthropoid primate and cetacean brain scaling. Journal of Evolutionary Biology, 2012, 25, 981-994.	0.8	147
87	Adrenal androgen production in catarrhine primates and the evolution of adrenarche. American Journal of Physical Anthropology, 2012, 147, 389-400.	2.1	44
88	Genomic data reject the hypothesis of a prosimian primate clade. Journal of Human Evolution, 2011, 61, 295-305.	1.3	45
89	Gene Expression and Methylation Signatures ofMAN2C1are Associated with PTSD. Disease Markers, 2011, 30, 111-121.	0.6	69
90	Spontaneous Abortion and Preterm Labor and Delivery in Nonhuman Primates: Evidence from a Captive Colony of Chimpanzees (Pan troglodytes). PLoS ONE, 2011, 6, e24509.	1.1	16

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91	Morris Goodman (1925â€“2010). Journal of Human Evolution, 2011, 60, 673-676.	1.3	0
92	Morris Goodman (1925â€“2010): Founder of the field of molecular anthropology. Evolutionary Anthropology, 2011, 20, 1-2.	1.7	0
93	SLC6A4 methylation modifies the effect of the number of traumatic events on risk for posttraumatic stress disorder. Depression and Anxiety, 2011, 28, 639-647.	2.0	140
94	Morris Goodman, Ph.D. (1925â€“2010). A Remembrance. Molecular Phylogenetics and Evolution, 2011, 58, 1-3.	1.2	2
95	Epigenetic and inflammatory marker profiles associated with depression in a community-based epidemiologic sample. Psychological Medicine, 2011, 41, 997-1007.	2.7	156
96	Silencing, Positive Selection and Parallel Evolution: Busy History of Primate Cytochromes c. PLoS ONE, 2011, 6, e26269.	1.1	14
97	Gene expression and methylation signatures of MAN2C1 are associated with PTSD. Disease Markers, 2011, 30, 111-21.	0.6	51
98	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.	1.2	42
99	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.	3.3	452
100	A primate subfamily of galectins expressed at the maternalâ€“fetal interface that promote immune cell death. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 9731-9736.	3.3	200
101	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.	3.3	75
102	Transcriptional Regulation of the Novel Toll-like Receptor Tlr13. Journal of Biological Chemistry, 2009, 284, 20540-20547.	1.6	29
103	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.	3.3	31
104	Development and evaluation of new mask protocols for gene expression profiling in humans and chimpanzees. BMC Bioinformatics, 2009, 10, 77.	1.2	9
105	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.	1.2	27
106	Phylogeny of the Ferungulata (Mammalia: Laurasiatheria) as determined from phylogenomic data. Molecular Phylogenetics and Evolution, 2009, 52, 660-664.	1.2	17
107	A fully resolved genus level phylogeny of neotropical primates (Platyrrhini). Molecular Phylogenetics and Evolution, 2009, 53, 694-702.	1.2	102
108	Adaptive history of single copy genes highly expressed in the term human placenta. Genomics, 2009, 93, 33-41.	1.3	27

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109	IDChase: Mitigating Identifier Migration Trap in Biological Databases. Communications in Computer and Information Science, 2009, , 461-472.	0.4	1
110	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.	1.2	33
111	ORIGINAL ARTICLE: Chorioamnionitis and Increased Galectinâ€1 Expression in PPROM â€“ An Antiâ€inflammatory Response in the Fetal Membranes?. American Journal of Reproductive Immunology, 2008, 60, 298-311.	1.2	43
112	Molecular evolution of the cytochrome c oxidase subunit 5A gene in primates. BMC Evolutionary Biology, 2008, 8, 8.	3.2	46
113	Severe preeclampsia is characterized by increased placental expression of galectin-1. Journal of Maternal-Fetal and Neonatal Medicine, 2008, 21, 429-442.	0.7	65
114	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.	3.3	61
115	Emergence of hormonal and redox regulation of galectin-1 in placental mammals: Implication in maternalâ€“fetal immune tolerance. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 15819-15824.	3.3	86
116	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.	0.7	42
117	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.	3.3	74
118	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.	3.3	158
119	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
120	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.	3.3	304
121	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.	3.3	303
122	Phylogenetic relationships and divergence times among New World monkeys (Platyrrhini, Primates). Molecular Phylogenetics and Evolution, 2006, 40, 274-280.	1.2	161
123	Trophoblast, Galectin-1 and pre-eclampsia. American Journal of Obstetrics and Gynecology, 2006, 195, S138.	0.7	2
124	New Onto-Tools: Promoter-Express, nsSNPCounter and Onto-Translate. Nucleic Acids Research, 2006, 34, W626-W631.	6.5	17
125	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.	3.3	16
126	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.	0.6	5

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127	The phylogenetic history of New World monkey Î² globin reveals a platyrrhine Î² to Î´ gene conversion in the atelid ancestry. Molecular Phylogenetics and Evolution, 2005, 35, 225-234.	1.2	23
128	Moving primate genomics beyond the chimpanzee genome. Trends in Genetics, 2005, 21, 511-517.	2.9	93
129	Rapid Nonsynonymous Evolution of the Iron-Sulfur Protein in Anthropoid Primates. Journal of Bioenergetics and Biomembranes, 2005, 37, 35-41.	1.0	9
130	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.	3.3	79
131	Accelerated evolution of the electron transport chain in anthropoid primates. Trends in Genetics, 2004, 20, 578-585.	2.9	181
132	Mitochondrial evidence for the origin of hamadryas baboons. Molecular Phylogenetics and Evolution, 2004, 32, 287-296.	1.2	64
133	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.	1.2	33
134	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.	3.3	213
135	Humankindâ€™s Place in a Phylogenetic Classification of Living Primates. , 2004, , 293-311.		9
136	Implications of natural selection in shaping 99.4% nonsynonymous DNA identity between humans and chimpanzees: Enlarging genus Homo. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 7181-7188.	3.3	231
137	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.	3.3	76
138	Episodic Positive Selection in Ape Cytochrome c Oxidase Subunit IV. Molecular Biology and Evolution, 2002, 19, 1812-1815.	3.5	22
139	A map of the common chimpanzee genome. BioEssays, 2002, 24, 490-493.	1.2	17
140	Molecular Evolution of Aerobic Energy Metabolism in Primates. Molecular Phylogenetics and Evolution, 2001, 18, 26-36.	1.2	81
141	Noninvasive methods for collecting fresh hair tissue. Molecular Ecology, 1999, 8, 1749-1750.	2.0	29